TSP 5.1
Reference Manual

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and

Clint Cummins

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Introduction

Welcome to the TSP 5.1 Help System

The TSP Help system contains the complete reference manual for TSP, providing a description of every command and command option, organized alphabetically by command. It is not intended as an introduction to the program, nor as a tutorial for the inexperienced TSP user. To learn how to use the program, you should obtain a copy of the TSP 5.1 User's Guide (available at http://www.tspintl.com/support/tsp/ug_online.htm) The User's Guide contains more discussion and examples of how to combine TSP statements and construct TSP programs.

The Help system can be useful in the following ways:

- To read a basic introduction of what TSP and why it is useful, see Introduction to TSP.
- To find out which command to use, see the functional index Commands by Function.
- You know which command or procedure you want to use, but are unsure of the options available. You can look up the details of the command in the Index (click on the Help Topics button above and then click the Index tab) and check the Options.
- To learn more about the methods used in any particular procedure, see the details in References under the command.
- To see how to use a command, look at the Examples.
- To check which results are stored after a procedure and how they are named, consult the Output section of the command entry.
- Basics gives complete definitions of TSP syntax, the interpretation of special characters, and the mathematical and statistical functions available.
- If you want to find out something related to the use of TSP through the Looking Glass, look it up in the TSP through the Looking Glass section.
- Examples of TSP Programs describes where to find sample TSP programs.
- You may also want to look at the TSP read me.
Introduction to TSP

TSP provides regression, forecasting, and other econometric tools on mainframe and personal computers. Areas where TSP can be useful include:

- Applied econometrics, including teaching
- Macro-economic research and forecasting
- Econometric analysis of cross section and panel data
- Sales forecasting
- Financial analysis
- Cost analysis and forecasting

TSP is installed on thousands of computers worldwide. Although TSP was developed by economists (starting with Version 1 in 1967) and most of its uses are in economics, there is nothing in its design that limits its usefulness to economic time series. Any statistical or econometric application involving data sets of up to about 20,000 (or even more) observations is suitable for TSP. For more information, prices, ordering, and upgrades, see our website, http://www.tspintl.com.
Examples of TSP Programs

TSP is a powerful program because it is not limited to its preprogrammed commands. To help you develop your own specialized procedures we have compiled some examples of TSP programs. The example programs can be found in the

\ldots\ldots\ldots\ldots Program Files\TSP 5.1\examples

directory (the exact path may vary depending on where you installed TSP). If you did a custom install of TSP, you may not have installed the examples. In order to get the examples, you will have to reinstall TSP and do either a “Typical” install or a “Custom” with the Examples option check-marked.

The TSP examples are further subdivided into five categories; Miscellaneous, ML PROC, Panel Data, Qualitative Dependent Variables and Time Series. Each of these categories has their own directory in the Examples directory. For a description of each examples, look at the file examples.txt in the Examples directory.

Updated examples may be found on the TSP International website at:

http://www.tspintl.com/examples
Composing Names in TSP

Every name must begin with a letter, _ # % or @ (exceptions: 2SLS and 3SLS commands).

Subsequent characters in a name may be letters, _ # % @, or digits.

# or % may not be used in names that appear in MATRIX commands.

The maximum number of characters permitted in a name is 64 (in versions prior to TSP 4.4 it was 8).
Composing Numbers in TSP

Every number must begin with a ., +, -, or a digit.

No spaces may appear within a number.

One decimal point may appear.

One E or D may appear followed immediately by a one or two digit number with or without a sign. This is interpreted as a power of 10 to multiply the first number.

Example: 1E2 = 100.

With free-format LOAD or READ commands, a . is interpreted as a missing value, and a repeat count with a * may be specified.

The largest value of a series (in absolute value) that may be stored in TSP is 1.E-37, unless OPTIONS DOUBLE ; is used. Values larger than this are set to missing. Scalars and matrices are always stored in double precision.

Examples:

3*0 is treated as 0 0 0
53 . 100 is treated as 53, missing, 100
Introduction

Composing Text Strings in TSP

A text string must be enclosed by matching pair of quotes (" or ").

Quotes are allowed in a string when they are of a different type from the enclosing quotes, i.e. "Can't" or "sometimes" (interpreted as Can’t and “sometimes”).
Composing TSP Commands

- Every statement begins with a command name.

Exceptions:

\[
\begin{align*}
X &= Y; \quad ? \text{ implicit GENR} \\
X(I) &= Y(I); \quad ? \text{ implicit SET} \\
100 <\text{statement}>; \quad ? \text{ statement label for GOTO}
\end{align*}
\]

- The command name may be abbreviated, as long as it is uniquely identified.

- Many statements can have options specified in parentheses after the command name. Option names may be abbreviated, like command names. There are three kinds of options:
  1. Boolean options, either on or off. On is specified by the name of the option, as in PRINT, and off is specified by the option name with NO in front of it, as in NOPRINT.
  2. Options of the form \texttt{option name = option value}. The value may be the name of a variable, a numerical value, or just a keyword, depending on the context.
  3. Options which give lists of variables, and are of the form \texttt{option name = (list of variables)}. Note that the parentheses are required, unless the list contains only one name, or the list is a listname.

- A few commands can be followed by an algebraic formula: GENR, SET, SMPLIF, SELECT, FRML, IDENT, IF, GOTO.

- Most commands are followed by one or more series names, separated by commas or spaces. These series names may include lags. An implicit list (such as \texttt{X1-X5}) can be used directly in a statement without making an intermediate listname. See the LIST command for a complete description of implicit list syntax.

- The end of a statement is marked by a semicolon (;) or dollar sign ($).
Introduction

Composing Algebraic Expressions in TSP

In general, TSP rules for formulas are similar to Fortran or other scientific programming languages.

A lag is indicated by putting an integer or a name in parentheses after a series name. The integer is negative for lags and positive for leads. A + sign is not necessary for leads. If the lag or lead is a name, it must have no more than four characters.

A series may have a single numeric or variable subscript (or lag/lead). A matrix may have a single or double subscript (numeric or variable). See the SET command for detailed rules and examples.

Arithmetic operators are:

<table>
<thead>
<tr>
<th>Operator</th>
<th>Description</th>
</tr>
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<tbody>
<tr>
<td>+</td>
<td>add</td>
</tr>
<tr>
<td>-</td>
<td>subtract</td>
</tr>
<tr>
<td>*</td>
<td>multiply</td>
</tr>
<tr>
<td>/</td>
<td>divide</td>
</tr>
<tr>
<td>** or ^</td>
<td>raise to the power</td>
</tr>
</tbody>
</table>

See TSP Functions for a detailed list of functions and the MATRIX command for matrix functions.

Relational and logical operators are the following:

<table>
<thead>
<tr>
<th>Operator</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>=</td>
<td>.EQ. gives the value 1 when the variables on the left and on the right are equal; otherwise it is zero</td>
</tr>
<tr>
<td>~= or ^=</td>
<td>.NE. gives the value 1 when the variables on the left and on the right are not equal; otherwise it is zero</td>
</tr>
<tr>
<td>&lt;</td>
<td>.LT. gives the value 1 when the variable on the left is less than the variable on the right; otherwise it is zero</td>
</tr>
<tr>
<td>&gt;</td>
<td>.GT. gives the value 1 when the variable on the left is greater than the variable on the right; otherwise it is zero</td>
</tr>
<tr>
<td>&lt;=</td>
<td>.LE. gives the value 1 when the variable on the left is less than or equal to the variable on the right; otherwise it is zero</td>
</tr>
<tr>
<td>&gt;=</td>
<td>.GE. gives the value 1 when the variable on the left is greater than or equal to the variable on the right</td>
</tr>
</tbody>
</table>
right; otherwise it is zero

<table>
<thead>
<tr>
<th>Operator</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td><code>&amp;</code></td>
<td>.AND. gives the value 1 when both the variable on the left and on the right are positive</td>
</tr>
<tr>
<td>`</td>
<td>`</td>
</tr>
<tr>
<td><code>~</code> or <code>^</code></td>
<td>.NOT. gives the value 1 when the variable on the right is negative or zero</td>
</tr>
</tbody>
</table>

**Note:** the .OP. form of the relational and logical operators is the alternative to the symbolic notation (but it cannot be used in nested DOT loops).

As many parentheses as necessary may be used to indicate the order of evaluation of a formula. The special parentheses [] and {} are treated as (). In the absence of parentheses, evaluation proceeds from left to right in the following order:

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<td>3</td>
<td>Multiplication and division</td>
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<tr>
<td>6</td>
<td>.NOT. (~)</td>
</tr>
<tr>
<td>7</td>
<td>.AND. (&amp;) and .OR. (</td>
</tr>
</tbody>
</table>
# TSP Functions

These functions can be used in any `GENR, FRML, IF, SET, SELECT`, or `MATRIX` command. Include the argument (value, series name, or algebraic expression) in the parentheses(). For additional matrix functions, see the `MATRIX` command.

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<td>Cosine (argument in radians)</td>
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<td>Tangent (argument in radians)</td>
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<td>Standard normal cumulative distribution function</td>
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<td>Inverse of the standard normal cumulative distribution function</td>
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<tr>
<td>GAMFN()</td>
<td>Gamma function (not Gamma density)</td>
</tr>
<tr>
<td>LGAMFN()</td>
<td>Log of Gamma function</td>
</tr>
<tr>
<td>DLGAMFN()</td>
<td>Derivative of LGAMFN = DIGAMMA()</td>
</tr>
<tr>
<td>TRIGAMMA()</td>
<td>Derivative of DIGAMMA() [non-differentiable]</td>
</tr>
<tr>
<td>FACT()</td>
<td>Factorial: FACT(X) = X! = GAMFN(X+1)</td>
</tr>
<tr>
<td>LFACT()</td>
<td>Log of factorial</td>
</tr>
<tr>
<td>SIGN()</td>
<td>Sign: -1 for X&lt;0, 0 for X=0, 1 for X&gt;0 [deriv=0]</td>
</tr>
<tr>
<td>POS()</td>
<td>Positive: POS(X) = max(0,X)</td>
</tr>
<tr>
<td>MISS()</td>
<td>Missing: 1 for X missing, 0 otherwise [non-differentiable]</td>
</tr>
<tr>
<td>INT()</td>
<td>Integer: truncate (round towards 0) [non-differentiable]</td>
</tr>
<tr>
<td>CEIL()</td>
<td>Ceiling: round away from 0 [non-differentiable]</td>
</tr>
<tr>
<td>ROUND()</td>
<td>Round to nearest integer (.5 rounds to 1) [non-differentiable]</td>
</tr>
</tbody>
</table>
### Character Set for TSP

<table>
<thead>
<tr>
<th>Character</th>
<th>Symbol</th>
<th>Use</th>
</tr>
</thead>
<tbody>
<tr>
<td>letter</td>
<td>A to Z, #,%@</td>
<td>Parts of names. Lowercase letters are allowed on most computers; they are treated like uppercase letters. #, % cannot be used as part of a name in the <strong>MATRIX</strong> command.</td>
</tr>
<tr>
<td>digit</td>
<td>0 to 9</td>
<td>Parts of numbers or names</td>
</tr>
<tr>
<td>decimal point</td>
<td>.</td>
<td>Marks the decimal point in numbers; sets off logical operators; specifies string substitution in the DOT procedure</td>
</tr>
<tr>
<td>comma</td>
<td>,</td>
<td>Separates the words in a list and arguments in multivariate functions like <strong>CNORM2</strong>; spaces may be used, but commas are often preferred for clarity</td>
</tr>
<tr>
<td>colon</td>
<td>:</td>
<td>Part of date</td>
</tr>
<tr>
<td>semicolon</td>
<td>;</td>
<td>Marks the end of a statement</td>
</tr>
<tr>
<td>dollar sign</td>
<td>$</td>
<td>Equivalent to ;. Semicolon is preferred.</td>
</tr>
<tr>
<td>quotation mark</td>
<td>“</td>
<td>Marks the beginning and end of a text string (title or filename); specifies matrix inversion</td>
</tr>
<tr>
<td>apostrophe</td>
<td>’</td>
<td>Marks the beginning and end of a text string; specifies matrix transposition</td>
</tr>
<tr>
<td>parentheses</td>
<td>() [] {}</td>
<td>Encloses a list of options or expressions/lags in algebraic formulas</td>
</tr>
<tr>
<td>question mark</td>
<td>?</td>
<td>Delimits the beginning of comments. (Comments are terminated by the end of the input line or logical record.)</td>
</tr>
<tr>
<td>plus sign</td>
<td>+</td>
<td>Specifies addition</td>
</tr>
<tr>
<td>minus sign</td>
<td>-</td>
<td>Specifies subtraction, a lag, or a list</td>
</tr>
<tr>
<td>star</td>
<td>*</td>
<td>Specifies multiplication or is part of power (**)</td>
</tr>
<tr>
<td>slash</td>
<td>/</td>
<td>Specifies division</td>
</tr>
<tr>
<td>pound sign</td>
<td>#</td>
<td>Matrix Kronecker product (∇)</td>
</tr>
<tr>
<td>percent</td>
<td>%</td>
<td>Matrix Hadamard product (element by element)</td>
</tr>
<tr>
<td>equal sign</td>
<td>=</td>
<td>Specifies equality or definition of data; relational operator (.EQ., .NE., .LE., .GE.).</td>
</tr>
<tr>
<td>ampersand</td>
<td>&amp;</td>
<td>Logical operator (.AND.)</td>
</tr>
<tr>
<td>vertical bar</td>
<td></td>
<td>Logical operator (.OR.); Also used to separate lists of variables in <strong>INST</strong>, <strong>KALMAN</strong>, <strong>LOGIT</strong>, <strong>SAMPSEL</strong>, and <strong>VAR</strong></td>
</tr>
<tr>
<td>caret or hat</td>
<td>^</td>
<td>Logical operator (.NOT., .NE.) or power (**),</td>
</tr>
</tbody>
</table>
depending on context

~ Logical operator (.NOT., .NE.)

< Relational operator (.LT., .LE.)

> Relational operator (.GT., .GE.)

\ Continuation of a line (interactive)

!` Reserved for future use
## Missing Values in TSP Procedures

<table>
<thead>
<tr>
<th>Procedures that drop observations containing missing values</th>
<th>Procedures that cannot execute if the sample contains missing values</th>
</tr>
</thead>
<tbody>
<tr>
<td>AR1, OLSQ, INST, 2SLS, LIML, LAD</td>
<td>ACTFIT</td>
</tr>
<tr>
<td>LSQ, FIML, GMM, ML, SUR, 3SLS</td>
<td>BJEST, BJIDENT, BJFRCST</td>
</tr>
<tr>
<td>PROBIT, TOBIT, SAMPSEL, LOGIT</td>
<td>COINT, UNIT</td>
</tr>
<tr>
<td>CONVERT, FORCST, GENR</td>
<td>ARCH, KALMAN</td>
</tr>
<tr>
<td>MSD, CORR, COVA, MOM</td>
<td>DIVIND, SAMA</td>
</tr>
<tr>
<td>GRAPH, PLOT, HIST</td>
<td>SOLVE, SIML</td>
</tr>
<tr>
<td>PANEL, VAR</td>
<td>PRIN</td>
</tr>
</tbody>
</table>
LOGIN.TSP file

*login.tsp* is a special *INPUT* file; it is read automatically at the start of interactive sessions and batch jobs. This is useful for setting default options for a run.

If you use the same options repeatedly, you may want to place them in a *login.tsp* file. Every time TSP starts, it checks for a *login.tsp* file, and executes it first. Normally, TSP looks for *login.tsp* in your working directory. If it does not find one, it looks in the directory in which you installed TSP for DOS and Windows, in the folder in which you installed TSP for Macs, and in the home directory on Unix.

Commands which can be usefully issued in a *login.tsp* file are the following

**OPTIONS MEMORY=** approximate memory to be used by TSP (in Megabytes). This option only works if OPTIONS is the first command in the run, or the first command in the *login.tsp* file.

**INPUT** some file that transforms the data or selects the data you are using for a number of TSP programs.
Command summary

Display Commands

- **ASMBUG**: prints debug output during parsing of TSP commands
- **DATE**: prints current date on screen or printout
- **DEBUG**: prints debug output during execution of TSP commands
- **DIR**: lists files available in current directory *(interactive)*
- **DOC**: adds descriptions to variables
- **GRAPH**: graph one variable against another *(graphics version)*
- **GRAPH**: graphs one variable against another in a scatter plot
- **HELP**: prints command syntax
- **HIST**: one-way bar chart for variable *(graphics version)*
- **NAME**: specifies name and title for TSP run
- **PAGE**: starts a new page in printout
- **PLOT**: plots several variables versus time *(graphics version)*
- **PLOT**: plots several variables versus time
- **PRINT**: prints variables
- **SHOW**: lists currently defined TSP variables by category (SERIES, etc.)
- **SYMTAB**: debug version of SHOW
- **TITLE**: specifies new title for run or immediate printing
- **TSTATS**: prints table of coefficients and t-statistics
Options Commands

- **FREQ**: set the data frequency (None, Annual, Quarterly, Monthly)
- **NOPLOT**: turns residual plots off (OLSQ, INST, AR1, LSQ)
- **NOREPL**: prevents splicing of series, generates missing values instead
- **OPTIONS**: general option setting -- CRT, HARDCOPY, LIMPRN, etc.
- **PLOTS**: turns residual plots off (OLSQ, INST, AR1, LSQ)
- **REPL**: allows updating of series during **GENR** (the default)
- **SELECT**: restricts the set of observations to those meeting a condition
- **SMPL**: set the sample of observations to be processed
- **SMPLIF**: same as SELECT, but restricts starting from current sample
### Moving Data to/from Files Commands

<table>
<thead>
<tr>
<th>Command</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>CLOSE</td>
<td>closes an external input or output file</td>
</tr>
<tr>
<td>DBCOMP</td>
<td>compresses a databank <em>(Databank)</em></td>
</tr>
<tr>
<td>DBCOPY</td>
<td>copies databank for moving to another computer <em>(Databank)</em></td>
</tr>
<tr>
<td>DBDEL</td>
<td>deletes variables from a databank <em>(Databank)</em></td>
</tr>
<tr>
<td>DBLIST</td>
<td>lists all variable names in a databank <em>(Databank)</em></td>
</tr>
<tr>
<td>DBPRINT</td>
<td>prints all series in a databank <em>(Databank)</em></td>
</tr>
<tr>
<td>FETCH</td>
<td>reads a microTSP-format databank</td>
</tr>
<tr>
<td>FORMAT</td>
<td>(option) -- used in READ and WRITE with numbers</td>
</tr>
<tr>
<td>IN</td>
<td>causes automatic searching of databanks listed <em>(Databank)</em></td>
</tr>
<tr>
<td>KEEP</td>
<td>stores TSP variables on specified OUT files <em>(Databank)</em></td>
</tr>
<tr>
<td>LOAD</td>
<td>reads variables from a file (or from program) - same as READ</td>
</tr>
<tr>
<td>NOPRINT</td>
<td>suppresses echoing of commands in a LOAD section</td>
</tr>
<tr>
<td>OUT</td>
<td>causes automatic databank storage in files listed <em>(Databank)</em></td>
</tr>
<tr>
<td>PRINT</td>
<td>same as WRITE</td>
</tr>
<tr>
<td>READ</td>
<td>reads variables from a file (or from program)</td>
</tr>
<tr>
<td>RECOVER</td>
<td>recovers lost program from INDX.TMP file <em>(Interactive)</em></td>
</tr>
<tr>
<td>RESTORE</td>
<td>reads variables from a SAVE file into the program <em>(Interactive)</em></td>
</tr>
<tr>
<td>SAVE</td>
<td>saves all current variables on disk <em>(Interactive)</em></td>
</tr>
<tr>
<td>STORE</td>
<td>writes a microTSP-format databank</td>
</tr>
<tr>
<td>WRITE</td>
<td>writes variables to a file (or to printout)</td>
</tr>
</tbody>
</table>
Command summary

Data Transformations Commands

**CAPITL** accumulates a capital stock from an investment series

**CONVERT** changes a series from higher to lower frequency (stock or flow)

**COPY** copies any variable

**DELETE** deletes any variables

**DIVIND** computes Divisia price and quantity indices

**DUMMY** makes dummy variables from a series

**GENR** creates a series using an algebraic formula

**LENGTH** computes length of a TSP list (useful in **PROCs**)

**NORMAL** normalizes a series (usually a price index, via division)

**RANDOM** random number generator: normal (univariate or multivariate), uniform, Poisson, or empirical (bootstrap)

**RENAME** renames a variable

**SAMA** Seasonal adjustment using the moving average method

**SET** modify a scalar or series/matrix element with an algebraic formula

**SORT** sorting data

**TREND** create linear trend variable (can be repeating like months)
Command summary

Matrix Operations Commands

- **MATRIX**  matrix algebra and transformations
- **MFORM**  change dimensions or type of matrix
- **MMAKE**  create a matrix from several series or a vector from scalars
- **ORTHON**  orthonormalization
- **UNMAKE**  create several series from a matrix or scalars from a vector
- **YLDFAC**  LDL' decomposition (symmetric indefinite matrix)
## Linear Estimation and Data Analysis Commands

<table>
<thead>
<tr>
<th>Command</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>CORR</td>
<td>correlation matrix of several series</td>
</tr>
<tr>
<td>COVA</td>
<td>covariance matrix of several series</td>
</tr>
<tr>
<td>INST</td>
<td>instrumental variables and two stage least squares regression</td>
</tr>
<tr>
<td>KERNEL</td>
<td>computes a Kernel density estimation or regression</td>
</tr>
<tr>
<td>LAD</td>
<td>Least Absolute Deviations estimation (median regression)</td>
</tr>
<tr>
<td>LIML</td>
<td>Limited Information Maximum Likelihood</td>
</tr>
<tr>
<td>LMS</td>
<td>Least Median Squares estimation</td>
</tr>
<tr>
<td>LP</td>
<td>Linear programming (with constraints)</td>
</tr>
<tr>
<td>MSD</td>
<td>mean, standard deviation, minimum, maximum, sum, variance, skewness, kurtosis for a list of series</td>
</tr>
<tr>
<td>OLSQ</td>
<td>Ordinary Least Squares (linear regression), can use weights</td>
</tr>
<tr>
<td>PANEL</td>
<td>panel data estimation (total, within, between, variance components)</td>
</tr>
<tr>
<td>PDL</td>
<td>describes specification of Polynomial Distributed Lag variables (Almon lags), and Shiller Lags; used in OLSQ, INST, AR1, PROBIT, TOBIT</td>
</tr>
<tr>
<td>PRIN</td>
<td>Principal Components (simple factor analysis)</td>
</tr>
</tbody>
</table>
Nonlinear Estimation and Formula Manipulation Commands

**CONST** defines scalars as fixed (non-estimable) constants

**DIFFER** create equations with analytic derivatives of formulas (FRMLs)

**EQUATE** equation substitution, one formula into another

**FIML** Full Information Maximum Likelihood estimation (system of linear or nonlinear equations, identities, implicit equations, general cross-equation restrictions, Multivariate Normal errors)

**FRML** define a linear or nonlinear equation for estimation

**GMM** Generalized Method of Moments estimation, nonlinear, with heteroskedasticity- and autocorrelation-robust standard errors

**IDENT** same as FRML but for an identity -- no implied disturbance (for FIML)

**LSQ** minimum distance estimation of single or multiple equation linear or nonlinear equations, general cross-equation restrictions, additive error terms (see SUR, 3SLS also)

**ML** Maximum Likelihood estimation, log likelihood specified in FRML

**MLPROC** Maximum Likelihood estimation, log likelihood specified in a PROC

**NONLINEAR** options describes iteration methods and options used by ARCH, BJEST, LSQ, FIML, LOGIT, ML, MLPROC, PROBIT, TOBIT, SIML, SOLVE

**PARAM** defines scalars as estimable parameters, can supply starting values

**SUR** Seeming Unrelated Regressions -- LSQ without instruments

**3SLS** Three Stage Least Squares -- LSQ with instruments
QDV (Qualitative Dependent Variable) Commands

**INTERVAL** estimates the Interval model (ordered Probit with known limits)

**LOGIT** estimates binary, multinomial, conditional, and mixed Logit models

**NEGBIN** estimates the Negative Binomial regression model for count data

**NONLINEAR options** describes iteration methods used by ARCH, BJEST, LSQ, FIML, LOGIT, PROBIT, TOBIT, and ML

**ORDPROB** estimates the Ordered Probit model

**POISSON** estimates the Poisson model for count data

**PROBIT** estimates the Probit model (0/1 with normal error term)

**SAMPSEL** estimates a two-equation Sample Selection model

**TOBIT** estimates the Tobit model (0/positive with normal error term)
Hypothesis Testing Commands

- **ANALYZ** computes standard errors for functions of parameters from a previous estimation
- **CDF** distribution functions and P-values
- **COINT** unit root and cointegration tests
- **REGOPT** controls printing and storage of regression diagnostics
Forecasting and Model Simulation Commands

**ACTFIT**  
compares actual and forecasted series

**BJFRCS T**  
forecasts Box-Jenkins ARIMA models

**FORCST**  
computes forecasts for estimated linear models (OLSQ, INST, AR1)

**FORM**  
constructs an equation (FRML) from an estimated linear model

**MODEL**  
orders large simultaneous equation systems for use by SOLVE

**SIML**  
simulation of general nonlinear systems of equations

**SOLVE**  
simulation of large (usually sparse) systems of equations
Time Series Identification and Estimation Commands

AR1 regression with correction for AR(1) (autocorrelated) error
ARCH estimates GARCH-M models
BJEST estimates Box-Jenkins ARIMA models
BJFRCST forecasts Box-Jenkins ARIMA models
BJIDENT identifies the order of Box-Jenkins ARIMA models
COINT unit root and cointegration tests
KALMAN Kalman filter estimation
UNIT synonym for COINT
VAR Vector autoregressions
## Command summary

### Control Flow Commands

<table>
<thead>
<tr>
<th>Command</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>COLLECT</td>
<td>delays execution of commands such as DO loops</td>
</tr>
<tr>
<td>COMPRESS</td>
<td>clears up space occupied by deleted variables</td>
</tr>
<tr>
<td>DO</td>
<td>defines a (numeric-indexed) DO loop</td>
</tr>
<tr>
<td>DOT</td>
<td>defines a (character-indexed) DOT loop</td>
</tr>
<tr>
<td>ELSE</td>
<td>part of IF-THEN-ELSE conditional structure</td>
</tr>
<tr>
<td>END</td>
<td>end of program</td>
</tr>
<tr>
<td>ENDDO</td>
<td>end of DO loop</td>
</tr>
<tr>
<td>ENDDOT</td>
<td>end of DOT loop</td>
</tr>
<tr>
<td>ENDPROC</td>
<td>end of PROC definition</td>
</tr>
<tr>
<td>EXEC</td>
<td>execute a range of command lines</td>
</tr>
<tr>
<td>EXIT</td>
<td>end of COLLECT loop or program</td>
</tr>
<tr>
<td>GOTO</td>
<td>starts execution at the statement label specified</td>
</tr>
<tr>
<td>IF</td>
<td>part of IF-THEN-ELSE conditional structure</td>
</tr>
<tr>
<td>INPUT</td>
<td>read commands from an external file</td>
</tr>
<tr>
<td>LIST</td>
<td>defines a list of variables</td>
</tr>
<tr>
<td>LOCAL</td>
<td>defines local variables in a PROC</td>
</tr>
<tr>
<td>OUTPUT</td>
<td>directs output to a file instead of screen</td>
</tr>
<tr>
<td>PROC</td>
<td>defines a user procedure</td>
</tr>
<tr>
<td>QUIT</td>
<td>stops TSP, without saving backup</td>
</tr>
<tr>
<td>STOP</td>
<td>stops TSP</td>
</tr>
<tr>
<td>SYSTEM</td>
<td>temporary exit to VMS or DOS without losing TSP session</td>
</tr>
<tr>
<td>TERMINAL</td>
<td>restores output to screen after OUTPUT</td>
</tr>
<tr>
<td>THEN</td>
<td>part of IF-THEN-ELSE conditional structure</td>
</tr>
<tr>
<td>USER</td>
<td>user-programmable command</td>
</tr>
</tbody>
</table>

*(Interactive) means the command is available in Interactive mode.*
Interactive Editing Commands and/or Data Commands

**ADD**  adds arguments to previous command (**Interactive**)
**CLEAR** clears TSP's memory (data storage) (**Interactive**)
**DELETE** deletes lines during execution (**Interactive**)
**DROP** deletes arguments from previous command (**Interactive**)
**EDIT** edits a command (**Interactive**)
**ENTER** enter data for a series (**Interactive**)
**FIND** lists lines containing a specific TSP command (**Interactive**)
**RETRY** edits previous command and re-executes it (**Interactive**)
**REVIEW** lists range of TSP command lines (**Interactive**)
**UPDATE** replaces observations in a series (**Interactive**)
# Command summary

## Obsolete Commands

<table>
<thead>
<tr>
<th>Old command</th>
<th>Replacement command</th>
</tr>
</thead>
<tbody>
<tr>
<td>INPROD x y z;</td>
<td><strong>MAT</strong> $z = x'y$</td>
</tr>
<tr>
<td>INV a a i deta;</td>
<td><strong>MAT</strong> $ai = a''$; <strong>MAT</strong> $deta = det(a)$</td>
</tr>
<tr>
<td>MADD x y z;</td>
<td><strong>MAT</strong> $z = x+y$</td>
</tr>
<tr>
<td>MATRAN x xt</td>
<td><strong>MAT</strong> $xt = x'$</td>
</tr>
<tr>
<td>MDIV x y z;</td>
<td><strong>MAT</strong> $z = x/y$</td>
</tr>
<tr>
<td>MEDIV x y z;</td>
<td><strong>MAT</strong> $z = x/y$</td>
</tr>
<tr>
<td>MEMULT x y z;</td>
<td><strong>MAT</strong> $z = x%y$</td>
</tr>
<tr>
<td>MMULT x y z;</td>
<td><strong>MAT</strong> $z = x^z$</td>
</tr>
<tr>
<td>MSQUARE x y;</td>
<td><strong>MAT</strong> $y = x'x$</td>
</tr>
<tr>
<td>MSUB x y z;</td>
<td><strong>MAT</strong> $z = x-y$</td>
</tr>
<tr>
<td>NOSUPRES x;</td>
<td><strong>REGOPT</strong> x;</td>
</tr>
<tr>
<td>SUPRES x;</td>
<td><strong>REGOPT</strong> (NOPRINT) X;</td>
</tr>
<tr>
<td>VGVMLT x y z;</td>
<td><strong>MAT</strong> $z = y$</td>
</tr>
<tr>
<td>YFACT x y;</td>
<td><strong>MAT</strong> $y = CHOL(x)$</td>
</tr>
<tr>
<td>YINV a a i;</td>
<td><strong>MAT</strong> $ai = YINV(a)$</td>
</tr>
<tr>
<td>YQUAD x y z;</td>
<td><strong>MAT</strong> $z = x'y''x'$</td>
</tr>
</tbody>
</table>
Cross-Reference Pointers

Command synonyms

<table>
<thead>
<tr>
<th>For</th>
<th>See</th>
</tr>
</thead>
<tbody>
<tr>
<td>LOAD</td>
<td>READ</td>
</tr>
<tr>
<td>COVA</td>
<td>MSD</td>
</tr>
<tr>
<td>CORR</td>
<td>MSD</td>
</tr>
<tr>
<td>NOSUPRES</td>
<td>REGOPT (PRINT)</td>
</tr>
<tr>
<td>PRINT</td>
<td>WRITE</td>
</tr>
<tr>
<td>SUPRES</td>
<td>REGOPT (NOPRINT)</td>
</tr>
<tr>
<td>2SLS</td>
<td>INST</td>
</tr>
<tr>
<td>UNIT</td>
<td>COINT</td>
</tr>
</tbody>
</table>

Non-command entries

<table>
<thead>
<tr>
<th>Function</th>
<th>See</th>
</tr>
</thead>
<tbody>
<tr>
<td>Functions</td>
<td>BASIC RULES</td>
</tr>
<tr>
<td>FORMAT</td>
<td>READ and WRITE</td>
</tr>
<tr>
<td>NONLINEAR</td>
<td>convergence options used in: LSQ, ML, FIML, BJEST, etc.</td>
</tr>
<tr>
<td>PDL</td>
<td>Polynomial Distributed Lags used in: OLSQ, AR1, etc.</td>
</tr>
</tbody>
</table>

Examples

For examples of | See                  |
<table>
<thead>
<tr>
<th></th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td>AR1</td>
<td>FORM, PDL</td>
</tr>
<tr>
<td>CDF</td>
<td>RANDOM</td>
</tr>
<tr>
<td>DOT</td>
<td>ELSE, LIST, SORT</td>
</tr>
<tr>
<td>ELSE</td>
<td>GOTO</td>
</tr>
<tr>
<td>EQSUB</td>
<td>ANALYZ, ML</td>
</tr>
<tr>
<td>FRML</td>
<td>FORM, LIST</td>
</tr>
<tr>
<td>GENR</td>
<td>DIFFER</td>
</tr>
<tr>
<td>IF</td>
<td>GOTO</td>
</tr>
<tr>
<td>INST</td>
<td>PDL</td>
</tr>
<tr>
<td>LIST</td>
<td>DUMMY, LENGTH</td>
</tr>
<tr>
<td>LOGIT</td>
<td>MMAKE</td>
</tr>
<tr>
<td>LSQ</td>
<td>ANALYZ, EQSUB</td>
</tr>
<tr>
<td>MAT</td>
<td>MMAKE, ORTHON</td>
</tr>
<tr>
<td>OLSQ</td>
<td>PDL</td>
</tr>
</tbody>
</table>
Command summary

<table>
<thead>
<tr>
<th>Command</th>
<th>Options</th>
</tr>
</thead>
<tbody>
<tr>
<td>PROC</td>
<td>LOCAL</td>
</tr>
<tr>
<td>RETRY</td>
<td>ADD, DROP, EDIT</td>
</tr>
<tr>
<td>SET</td>
<td>DO</td>
</tr>
<tr>
<td>THEN</td>
<td>GOTO</td>
</tr>
</tbody>
</table>
Commands

ACTFIT

ACTFIT computes and prints a variety of goodness-of-fit statistics for the actual and predicted values of a series. Theil (references below) suggests using these statistics for evaluating an estimated time series equation or forecast.

**ACTFIT (SILENT,TERSE) <actual series name> <predicted series name>**

After ACTFIT, give the name of the actual data series followed by the name of the fitted or predicted series.

**Output**

ACTFIT prints a title, the names of the series being compared, the time period (sample) over which they are compared, and then a variety of computed statistics on the comparison. These include the correlation of the two series, the mean square error, the mean absolute error, Theil's inequality coefficient (U), changes and percent changes in U, and a decomposition of the source of the discrepancies between the two series: differences in the mean, or differences in the variance.

The following scalar results are stored by ACTFIT:

<table>
<thead>
<tr>
<th>variable</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@R</td>
<td>1</td>
<td>Correlation coefficient</td>
</tr>
<tr>
<td>@R2</td>
<td>1</td>
<td>Correlation coefficient squared</td>
</tr>
<tr>
<td>@RMSE</td>
<td>1</td>
<td>Root mean square error</td>
</tr>
<tr>
<td>@MSE</td>
<td>1</td>
<td>Mean squared error</td>
</tr>
<tr>
<td>@MAE</td>
<td>1</td>
<td>Mean absolute error</td>
</tr>
<tr>
<td>@ME</td>
<td>1</td>
<td>Mean error</td>
</tr>
<tr>
<td>@RMSPE</td>
<td>1</td>
<td>Root-Mean-Squared Percent Error</td>
</tr>
<tr>
<td>@MSPE</td>
<td>1</td>
<td>Mean-Squared Percent Error</td>
</tr>
<tr>
<td>@MAPE</td>
<td>1</td>
<td>Mean Absolute Percent Error</td>
</tr>
<tr>
<td>@MPE</td>
<td>1</td>
<td>Mean Percent Error</td>
</tr>
<tr>
<td>@BETA</td>
<td>1</td>
<td>Regression coef. of Actual on Predicted</td>
</tr>
<tr>
<td>@U66</td>
<td>1</td>
<td>Theil's U Inequality coef. (Changes) U66</td>
</tr>
<tr>
<td>@U66P</td>
<td>1</td>
<td>Theil's U Ineq. coef. (Percent changes) U66P</td>
</tr>
<tr>
<td>@FBIAS</td>
<td>1</td>
<td>Fraction of MSE due to Bias</td>
</tr>
<tr>
<td>Symbol</td>
<td>Value</td>
<td>Description</td>
</tr>
<tr>
<td>----------</td>
<td>-------</td>
<td>--------------------------------------------------</td>
</tr>
<tr>
<td>@FDVAR</td>
<td>1</td>
<td>Fraction of MSE due to different Variation</td>
</tr>
<tr>
<td>@FDCOV</td>
<td>1</td>
<td>Fraction of MSE due to difference Covariation</td>
</tr>
<tr>
<td>@FDB1</td>
<td>1</td>
<td>(Alt.Decomp.) Frac. due to Diff. of BETA from 1</td>
</tr>
<tr>
<td>@FRES</td>
<td>1</td>
<td>(Alt.Decomp.) Frac. due to Residual variance</td>
</tr>
</tbody>
</table>

**Note**

U is defined differently in the 1961 and 1966 references. The 1966 definition is used in TSP Versions 4.0 and 4.1; under this definition U can be greater than one. In TSP Version 4.2 and above, both versions of U are printed. This output is followed by a time series residual plot of the two series if the PLOTS option is on (see the OPTIONS command). If the RESID option is on, the residual series will be stored under the name @RES, whether or not the PLOTS option is on.

**Options**

SILENT/NOSILENT suppresses all printed output.

TERSE/NOTERSE prints a reduced output.

**Example**

ACTFIT R RS;

**References**


ADD (interactive)

Examples

ADD adds a list of variables to the previous statement and re-executes it. It is the opposite of DROP.

ADD <list of variables> ;

ADD offers a convenient means of adding variables to a regression and performing a second estimation (without having to fully retype the command). It is not, however, restricted to this usage, and may be used in any circumstance where this type of command modification is needed.

The command

ADD var1 var2

and the sequence

RETRY
>> INSERT var1
>> INSERT var2
>> EXIT

are identical in function since both permanently modify the previous command by inserting var1 and var2 at the end of the command. The command is then automatically executed in both cases. The only potential difference between these approaches (besides the amount of typing) is in the definition of "previous". RETRY with no line number argument assumes you want to modify the last line typed. ADD will not accept a line number argument, and always modifies the last line that is not itself an ADD (or DROP) command.

ADD and DROP allow you to execute a series of closely related regressions by entering the first estimation command, followed by a series of ADD and DROP commands. Since each ADD or DROP permanently alters the command, each new modification must take all previous modifications into account.

Notes

It is not possible to combine ADD and DROP into one step to perform a replace function, or to make compound modifications to a command. In these circumstances, RETRY must be used.

Examples
ADD

\textit{OLSQ (WEIGHT=POP) YOUNG,C,RSALE,URBAN,CATHOLIC}  
\textit{ADD MARRIED}

will run two regressions, the second of which is:

\textit{OLSQ (WEIGHT=POP) YOUNG,C,RSALE,URBAN,CATHOLIC,MARRIED}

This is also how the command will now look if you \texttt{REVIEW} it, since it has been modified and replaced both in TSPs internal storage and in the backup file.

Another use for ADD might be in producing plots. The following will produce two plots with the same option settings, but two series are added to the second plot.

\texttt{PLOT (MIN=500,MAX=1500,LINES=(1000)) GNP G GNPS H}  
\textit{ADD CONS C CONSS D}
ANALYZ computes the values and estimated covariance matrix for a set of (nonlinear) functions of the parameters estimated by the most recent OLSQ, LIML, LSQ, FIML, PROBIT, etc. procedure. It also computes the Wald test for the hypothesis that the set of functions are jointly zero. If the functions are linear, after an OLSQ command, the F test of the restrictions, and implied restricted original coefficients will be printed. ANALYZ can also be used to compute values and standard errors for function of parameters and series; in this case the result will be two series, one containing the values corresponding to each observation, and the other the standard errors.

The method used linearizes the nonlinear functions around the estimated parameter values and then uses the standard formulas for the variance and covariance of linear functions of random variables. See the references for further discussion of this "delta method". TSP obtains analytic derivatives internally for the nonlinear functions. ANALYZ can also be used to select/reorder a subset of a VCOV matrix and COEF vector, for use in making a Hausman specification test.

ANALYZ (COEF=<input parameter vector>, HALTON, NAMES=(<list of names>), NDRAW=<number of draws>, PRINT, PRMEAN, PRSERIES, SILENT, VCOV=<matrix name>) <list of equation names> ;

Usage

ANALYZ is followed by a list of equation (FRML) names. After estimation procedures with linear models (OLSQ, INST, LIML, PROBIT, ...), these equations specify functions of the estimated coefficients which are to be computed by referring to the coefficients by the names of the associated variables. After estimation procedures with nonlinear models (LSQ and FIML), the equations specify functions of the estimated parameters. ANALYZ has no provision for combining the variances from more than one estimation, because it cannot obtain the associated covariance of the coefficient estimates. The equations must be previously defined by FRML statements; if the FRML statements have variable names on the left hand side, the computed value of each function will be stored under that variable name.
If series names (other than the names of right hand side variables from the previous OLSQ, INST, LIML, or PROBIT estimation) are included in the FRML(s), a series of values will result. One application for this kind of FRML is an elasticity which depends on estimated parameters, and also on data such as income. ANALYZ will compute the standard errors for such a FRML using the covariance matrix of the estimated parameters, and treating the data as fixed constants. See the example below of computing an elasticity series.

Output

If the PRINT option is on, ANALYZ prints a title, the names of the input parameters, the equations in symbolic form, a table of the derived functions and their standard errors, and the chi-squared value of a test that the functions are jointly zero. This chi-squared has degrees of freedom equal to the number of equations. The P-value (significance level) for the chi-squared test is also printed. If the print option is off (the default), only the derived functions and the chi-squared test are printed.

ANALYZ also stores the calculated parameters and their variances in data storage as though they were estimation results, whether or not the PRINT option is on. The results are stored under the following names.

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@RNMSA</td>
<td>list</td>
<td>#eqs</td>
<td>Names of derived parameters.</td>
</tr>
<tr>
<td>@WALD</td>
<td>scalar</td>
<td>1</td>
<td>Value of Wald test.</td>
</tr>
<tr>
<td>@NCOEFA</td>
<td>scalar</td>
<td>1</td>
<td>Number of derived parameters.</td>
</tr>
<tr>
<td>@NCIDA</td>
<td>scalar</td>
<td>1</td>
<td>Degrees of freedom.</td>
</tr>
<tr>
<td>%WALD</td>
<td>scalar</td>
<td>1</td>
<td>P-value (significance) of Wald test.</td>
</tr>
<tr>
<td>@COEFA</td>
<td>vector</td>
<td>#eqs</td>
<td>Values of derived parameters.</td>
</tr>
<tr>
<td>@SESA</td>
<td>vector</td>
<td>#eqs</td>
<td>Standard errors of derived parameter.</td>
</tr>
<tr>
<td>@TA</td>
<td>vector</td>
<td>#eqs</td>
<td>T-statistics (asymptotically normal).</td>
</tr>
<tr>
<td>%TA</td>
<td>vector</td>
<td>#eqs</td>
<td>p-values corresponding to @TA</td>
</tr>
<tr>
<td>@MSD</td>
<td>matrix</td>
<td>#eqs*8</td>
<td>Matrix of simulation results when NDRAW option is used.</td>
</tr>
<tr>
<td>@VCOVA</td>
<td>matrix</td>
<td>#eqs*#eqs</td>
<td>Estimated variance covariance of derived</td>
</tr>
</tbody>
</table>
parameters.

**Method**

Assume that a previous estimation in TSP has stored a vector of $K$ parameter estimates $b$ stored in @COEF and their variance covariance matrix $\text{Var}(b)$ stored in @VCOV. Values and standard errors for the $M$ functions $f(b)$ are desired. To compute these, ANALYZ obtains the first derivatives of $f$ with respect to $b$ analytically:

$$\begin{align*}
G &= \frac{\partial f}{\partial b} \\
\text{Var}[f(b)] &= G' \text{Var}(b) G
\end{align*}$$

The functions $f(b)$ and the matrix $G$ are evaluated at the current values of $b$ and any constants or data values which may appear in $f(b)$. The variance-covariance matrix for $f(b)$ is then (asymptotically, or exactly if $f(b)$ is linear in $b$) defined as

$$\text{Var}[f(b)] = G' \text{Var}(b) G$$

This is known as the "delta method". For example, if $M=1$ and $f(b) = f_1 = 2b_1$, then $G=2$ (with zeros elsewhere if $K>1$), and $\text{Var}(f_1) = 4 \text{Var}(b_1)$.

If the equations are linear, and an OLSQ command was used for estimation, ANALYZ prints the F-statistic for the set of joint restrictions (@FST = @WALD/@NCIDA). In addition, ANALYZ computes and prints the implied restricted original coefficients and their standard errors. These are stored under @COEFC, @VCOVC, etc.

**Options**

- **COEF=** vector containing the values of the parameters in the equations to be analyzed. This vector should correspond to the parameters listed in the NAMES= option, and also to the supplied VCOV matrix. The default is @COEF.

- **HALTON** specifies that a (shuffled) Halton sequence is used for the random draws when the NDRAW option is given. This provides more uniform coverage of the range of values, so it may yield more accurate integration for a given number of draws.

- **NAMES=** specifies an optional list of parameter names which are the labels for an associated covariance matrix supplied by the VCOV= option. The default is @RNMS.
**ANALYZ**

`NDRAW=n` computes asymmetric confidence intervals for nonlinear functions by drawing `n` simulated parameter vectors. These functions can vary over time as well. This is an alternative to the default "delta method" which uses derivatives and is exact for linear functions. The percentiles 2.5% and 97.5% are computed, to construct a two-tailed confidence interval at the 95% significance level. A matrix named @MSD with columns `SE_T_LB2.5%_UB97.5%_MEAN_MIN_MAX_NUM_GOOD` is stored. `NUM_GOOD` is the number of nonmissing results computed. Numeric errors such as division by zero result in missing values. When ANALYZ is used with series and `NDRAW`, the results are stored in series whose names are the name of the parameter computed followed by `_SE_T_LB_UB_MEAN_MIN_MAX_NUMGOOD`. See the Examples for an illustration.

**PRINT/NOPRINT** tells whether or not the ANALYZ input is to be printed. Under the default, NOPRINT, only the results are printed.

**PRMEAN/NOPRMEAN** tells whether summary statistics (mean, standard deviation, minimum, maximum, and median) are to be computed for the derived series when ANALYZ is used on equations containing series. PRMEAN is true by default.

**PRSERIES/NOPRSERIES** tells whether the computed series are to be printed when ANALYZ is used on equations containing series. The default for PRSERIES is TRUE when the number of observations is less than or equal to 100 and FALSE otherwise.

**SILENT/NOSILENT** specifies that no output is to be produced. The results are stored under the names `@RNMSA`, `@COEFA`, etc. Note that `REGOPT(NOPRINT) COEF;` is also needed, to suppress printing of the table of coefficients.

**VCOV=** specifies the name of a variance-covariance matrix of the input parameters (whose names are given by `NAMES=`). The use of these two options enables one to do an ANALYZ on matrices other than the `@VCOV` matrix from a standard estimation procedure. The default is `@VCOV`.

**Examples**

Obtain "long-run" coefficients for models with lagged dependent variables:

```
FRML LR1 ALPHALR = ALPHA/(1-LAMBDA) ;
FRML LR2 PHILR = PHI/(1-PSI) ;
ANALYZ LR1,LR2 ;
```

See the **EQUATION** command for an example of using ANALYZ (with EQSUB) to evaluate and obtain standard errors for restricted parameters in a translog system.
The next example shows how to calculate an elasticity (and its standard errors) when the elasticity changes over the sample.

```
FRML EQ1 LQ1 = A1 + B1*LP1 + B2*LP2 + B12*LP1*LP2 + 
B13*LP1*LP3 + B23*LP2*LP3;
FRML EL1 ELD1 = B1 + B12*LP2 + B13*LP3; ? d(LQ1)/d(LP1)
SMPL 48,95;
LSQ EQ1;
? Obtain and plot elasticity for each year between 1948 and 1995:
SMPL 48,95;
ANALYZ (NOSPLOT) EL1;
PLOT ELD1;
? Compute the average elasticity and its average s.e. - not needed in 
v5.1 and later, as this is automatic.
MSD ELD1 ELD1_SE;
```

Here is an example of using ANALYZ after OLSQ. It computes a chi-squared test of the hypothesis that the sum of the two coefficients is zero (this test statistic equals the standard F-statistic).

```
OLSQ Y C X1 X2;
FRML SUM X1+X2;
ANALYZ SUM;
```

Suppose that we want to extract a few parameters and their associated VCOV matrix from a system with a large number of parameters in arbitrary order:

```
SUPRES COEF;
LSQ (SILENT) EQ1-EQ50; ? estimation with a large number of 
equations
SUPRES;
DOT B1-B5;
FRML EQ. . = . ; ? construct FRML EQB1 B1 = B1; etc.
ENDDOT;
ANALYZ EQB1-EQB5; ? print results for 5 of the parameters only
RENAME @VCOVA VCVB1_5; ? for use later
RENAME @COEFA CB1_B5;
```

Here is an example using random draws to compute asymmetric confidence intervals:

```
FRML EQ1 Y = A+B*X;
LSQ EQ1;
FRML EQS SUM = A+B;
FRML EQR RATIO = A/B;
ANALYZ(NDRAW=500) EQS EQR;
```
In this example, the scalars SUM and RATIO will be stored, and eight statistics on the 500 computations of the two functions will be printed and stored in the 2 x 8 matrix @MSD.

An example of series output and random draws:

```plaintext
PROBIT D C X;
FRML EQP P = CNORM(A+B*X);
ANALYZ(NDRAW=200,NAMES=(A,B)) EQP;
```

In this example, the series P, P_SE, P_T, P_LB, P_UB, P_MEAN, P_MIN, P_MAX, P_NG will be stored and printed.

References


AR1 obtains estimates of a regression equation whose errors are serially correlated. These estimates are efficient if the disturbances in the equation follow an autoregressive process of order one. The estimates may be obtained using one of two different objective functions: exact maximum likelihood (which imposes stationarity by constraining the serial correlation coefficient to be between -1 and 1 and keeps the first observation for estimation), or by Generalized Least Squares (GLS), which drops the first observation.

\[ \text{AR1} \text{(FAIR, FEI, INST=} \text{(list of instrumental variables)}, \text{METHOD=} \text{CORC or HILU or ML or MLGRID, OBJFN=} \text{EXACTML or GLS, REI, RMIN=} \text{<minimum rho value>, RMAX=} \text{<maximum rho value>, RSTART=} \text{<start value for rho>, RSTEP=} \text{<step value for rho>, TSCS, nonlinear options)} \text{<dependent variable name>} \text{<list of independent variables> ;} \]

To obtain estimates of a regression equation which are corrected for first order serial correlation, use the AR1 command as you would an OLSQ command. PDL (polynomial distributed lag) variables may be included in an AR1 statement. See the PDL section for a further description of how to specify these variables. TSP automatically deletes observations with missing values for one or more variables before estimation.

When the SMPL frequency is type PANEL, AR1 can also obtain estimates for a panel data model with fixed (FEI) or random (REI) effects. AR1 estimates can also handle plain time series which have irregular spacing (gaps in the SMPL).

Output

The AR1 procedure produces output that is similar to OLSQ and LSQ (including the iteration log). The equation title and the chosen objective function (method of estimation) are printed first. If the PRINT option is on, this is followed by the list of option values, the starting values for all the coefficients, iteration output for all coefficients, and any grid values for rho and the objective function.
The usual regression output follows, as described under the OLSQ command. The regression statistics are computed from the fitted values and residuals described below. If the objective function chosen was GLS, a common factor test is included in the regression statistics. This test is a likelihood ratio test of the restrictions implied by AR(1) compared to an unconstrained OLS model that includes the lagged dependent as well as the current and lagged right hand side variables. [The test is not well-defined when the model is estimated by ML due to the special treatment of the first observation].

As in OLSQ and INST, a table of coefficient estimates is printed. RHO is always the last coefficient in the table; its inclusion guarantees that the standard errors are always consistent, even if there are lagged dependent variables on the right hand side. The fitted values (@FIT) and residuals (@RES) are computed as follows:

\[
\begin{align*}
@FIT(t) &= X_t b \\
@FIT(t) &= X_t b + \rho(y_{t-1} - X_{t-1} b) \quad \text{if} \quad t = 1, OBJFN = EXACTML \\
@FIT(t) &= y_t - @FIT(t) \\
@RES(t) &= y_t - @FIT(t) 
\end{align*}
\]

AR1 also stores this regression output in data storage for later use. The table below lists the results available after an AR1 command. Note: the number of coefficients (# vars) always includes RHO.

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@RNMS</td>
<td>list</td>
<td>#vars</td>
<td>Names of right hand side variables</td>
</tr>
<tr>
<td>@LHV</td>
<td>list</td>
<td>1</td>
<td>Name of the dependent variable</td>
</tr>
<tr>
<td>@RHO</td>
<td>scalar</td>
<td>1</td>
<td>Serial correlation parameter at convergence</td>
</tr>
<tr>
<td>@SSR</td>
<td>scalar</td>
<td>1</td>
<td>Sum of squared residuals</td>
</tr>
<tr>
<td>@S</td>
<td>scalar</td>
<td>1</td>
<td>Standard error of regression</td>
</tr>
<tr>
<td>@YMEAN</td>
<td>scalar</td>
<td>1</td>
<td>Mean of the transformed dependent variable</td>
</tr>
<tr>
<td>@SDEV</td>
<td>scalar</td>
<td>1</td>
<td>Standard deviation of the dependent variable</td>
</tr>
<tr>
<td>@NOB</td>
<td>scalar</td>
<td>1</td>
<td>Number of observations</td>
</tr>
<tr>
<td>@DW</td>
<td>scalar</td>
<td>1</td>
<td>Durbin-Watson statistic</td>
</tr>
<tr>
<td>@RSQ</td>
<td>scalar</td>
<td>1</td>
<td>R-squared</td>
</tr>
<tr>
<td>@ARSQ</td>
<td>scalar</td>
<td>1</td>
<td>Adjusted R-squared</td>
</tr>
<tr>
<td>@IFCONV</td>
<td>scalar</td>
<td>1</td>
<td>=1 if convergence achieved, =0 otherwise</td>
</tr>
<tr>
<td>@LOGL</td>
<td>scalar</td>
<td>1</td>
<td>Log of likelihood function.</td>
</tr>
<tr>
<td>@COMFAC</td>
<td>scalar</td>
<td>1</td>
<td>Common factor test (if OBJFN=GLS)</td>
</tr>
</tbody>
</table>
AR1

@COEF vector #vars Coefficient estimates.
@SES vector #vars Standard Errors.
@T vector #vars t-statistics.
%T vector #vars p-values for t-statistics.
@COEFAI vector #vars Fixed effect estimates (FEI)
@SESAI vector #vars Standard Errors on fixed effects (FEI).
@TAI vector #vars t-statistics on fixed effects (FEI).
%TAI vector #vars p-values for t-statistics on FEs (FEI).
@VCOV matrix #vars*#vars Variance-covariance of estimated coefficients.
@AI series #obs Fixed effect for each obs, in series form (FEI)
@RES series #obs Fitted residuals from model.
@FIT series #obs Fitted values of dependent variable.

If the regression includes PDL variables, @SLAG, @MLAG, and @LAGF will also be stored (see OLSQ for details).

Method

AR1 uses an initial grid search to local possible multiple local optima (when OBJFN = GLS), and then iterates efficiently to a global optimum with second derivatives. The likelihood function and treatment of the initial observation are described completely in Davidson and MacKinnon (1993).

When OBJFN=EXACTML (the default), AR1 simply maximizes the likelihood function for disturbances that follow a stationary autoregressive process with respect to the serial correlation rho and the coefficients of the independent variables.

For panel data, AR1 with fixed (FEI) or random (REI) effects is similar to the corresponding PANEL regressions, but with an added AR(1) component. The random effects estimator follows Baltagi and Li (1991). It uses analytic second derivatives to obtain quadratic convergence and accurate t-statistics for all parameters (including RHO and RHO_I, the intraclass correlation coefficient, which can be negative). After the fixed effects AR1 estimator, the estimated fixed effects are stored in the matrix @COEFAI and in the series @AI.

Options
FAIR/NOFAIR specifies whether the lagged dependent and independent variables are to be added to the instrument list automatically when doing instrumental variable estimation combined with a serial correlation correction.

FEI/NOFEI specifies that an AR(1) model with panel fixed effects is to be estimated by means of maximum likelihood (or GLS if OBJFN=GLS is specified).

INST= list of instrumental variables. This list should include any exogenous variables that are in the equation such as the constant or time trend, as well as any other variables you wish to use as instruments. After any instruments are added by the FAIR option, there must be at least as many instruments as the number of estimated coefficients (the number of independent variables in the equation, plus one for rho). OBJFN= GLS is implied; the actual objective function is E'PZ*E, where the Es are rho-transformed residuals. See the Examples for a way to reproduce the AR1 estimates with FORM and LSQ.

Fair once argued that the lagged dependent and independent variables must be in the instrument list to obtain consistent estimates when doing instrumental variable estimation with a serial correlation correction. TSP adds them automatically if you use the FAIR option (the default); if you want to specify a different list of instruments, you must suppress this feature with a NOFAIR option.

Fair retracted his claim in 1984; it has since been disproved by Buse (1989), but the alternative instruments for consistency involve pseudo-differencing with the estimated rho (Theil's G2SLS), which is tedious to perform by hand. Buse also showed that the asymptotically most efficient estimator in this case (S2SLS) includes the lagged excluded exogenous variables as well, but he cautions that in small samples this may quickly exhaust the degrees of freedom.

METHOD=ML or MLGRID or CORC or HILU was formerly used to specify the estimation algorithm. This is now specified by the OBJFN= option. METHOD=ML or MLGRID imply OBJFN=EXACTML, while METHOD=CORC or HILU imply OBJFN=GLS. METHOD=ML formerly used the Beach and McKinnon algorithm, while METHOD=CORC used the Cochrane-Orcutt algorithm. Now iterations are done using the Newton-Raphson algorithm (HITER=N in the nonlinear options) which is quadratically convergent (about the same speed as Beach-MacKinnon, but much faster and more accurate than Cochrane-Orcutt). METHOD=HILU refers to Hildreth-Lu, a simple grid search method.
**OBJFN=EXACTML** or GLS specifies the objective function. EXACTML retains the first observation and includes the Jacobian term \( \log(1-rho^2) \), which guarantees stationarity. GLS drops the first observation and does not impose stationarity. It is the same as nonlinear least squares on a rho-differenced equation, and can also be described as "conditional ML" (conditional on the initial residual).

**EXACTML** is the usual default, but if there is a lagged dependent variable on the right-hand side, GLS becomes the default, because EXACTML has a small-sample bias in this case.

GLS uses an initial grid search to locate starting values and potential multiple local optima. It is well known that multiple local optima can occur for GLS, especially when there are lagged dependent variables. Multiple optima are noted in the output if they are detected. AR1 then iterates efficiently to locate an accurate global optimum. EXACTML normally skips the grid search, because no cases of multiple local optima are known when the Jacobian is included. **METHOD=MLGRID** will turn on this grid search.

**REI/NOREI** specifies that an AR(1) model with panel random effects is to be estimated by means of maximum likelihood (GLS is not available for this model).

**RMIN=** specifies the minimum value of the serial correlation parameter rho for the initial grid search (when OBJFN=GLS or METHOD=MLGRID are used). The default value is -0.9.

**RMAX=** specifies the maximum value of rho for the grid search methods. The default value is 1.05 for OBJFN=GLS, or .95 for METHOD=MLGRID.

**RSTEP=** specifies the increment to be used in the grid search over rho. The default value is 0.1, until rho=.8. Then the values .85, .9, .95 are used, plus .9999, 1.0001, and 1.05 when OBJFN=GLS. These last 3 values help to detect optima with rho > 1, which are usually not reached during iterations when rho starts below 1.

**RSTART=** specifies a starting value of rho for the iterative methods. Ordinarily zero is used for OBJFN=EXACTML, but faster convergence may be achieved if a value closer to the true answer is chosen. RSTART can also be used to override the default grid search for OBJFN=GLS, but multiple local optima would not be detected.

**TSCS/NOTSCS** specifies EXACTML estimation for time series-cross section data when the **FREQ** (PANEL) command is in effect (then TSCS is the default) or when SMPL gaps have been set up to separate the cross section units (see the example below). OBJFN=GLS is not implemented for panel data.
(Obsolete) \texttt{WEIGHT=} is a former AR1 option which is no longer supported. The ML or LSQ commands should be used instead to implement a weight.

**Nonlinear options** are described under \texttt{NONLINEAR} in this manual. \texttt{HITER=N/HCOV=N} (second derivatives, the default) and \texttt{G} (first derivatives) are both available. \texttt{MAXIT=0} can be used to avoid iterations and to perform a simple grid search without the additional accuracy of iterations. Also, \texttt{AR1} uses a special default \texttt{TOL=1E-6 (.000001)}.

**Examples**

This example estimates the consumption function for the illustrative model with a serial correlation correction, first using the maximum likelihood method, and then searching over \( \rho \) to verify that the likelihood is unimodal in the relevant range.

\begin{verbatim}
AR1 (PRINT) CONS C GNP ;
AR1 (METHOD=MLGRID, RSTEP=0.05) CONS C GNP ;
\end{verbatim}

The next three estimations are exactly equivalent and demonstrate the \texttt{FAIR} option with instrumental variables:

\begin{verbatim}
SMPL 11,50;
AR1 (INST=(C,G,TIME,LM)) CONS C GNP ;
AR1 (NOFAIR,INST=(C,G,TIME,LM,GNP(-1),CONS(-1))) CONS C GNP;
FORM(NAR=1,PARAM,VARPREF=B) EQAR1 CONS C GNP;
? Drop first observation, to compare with \texttt{AR1(OBJFN=GLS)} results.
SMPL 12,50;
LSQ(INST=(C,G,TIME,LM,GNP(-1),CONS(-1))) EQAR1;
\end{verbatim}

Lagged dependent variable (default \texttt{OBJFN=GLS}, since \texttt{EXACTML} has a small sample bias):

\begin{verbatim}
AR1 CONS C GNP CONS(-1);
\end{verbatim}

Time series-cross section with 10 years of data and 3 cross section units, and fixed effects:

\begin{verbatim}
SMPL 1,10;
FREQ (PANEL,T=10);
AR1 (FEI) SALES C ADV POP GNP ;
AR1 (REI) SALES C ADV POP GNP ;
\end{verbatim}

**References**


ARCH estimates regression models with AutoRegressive Conditional Heteroskedasticity (originated by Robert Engle). It will estimate any model from linear regression to GARCH-M. ARCH models allow the residuals to have a variable variance (but still have zero conditional mean) over the sample. This contrasts with AR(1) models or general transfer function models where the residuals do not have zero conditional mean. ARCH models are often used to model exchange rate fluctuations and stock market returns.

ARCH (E2INIT=method, GT=<list of weighting series>, HEXP=<value of lambda>, HINIT=method, MEAN, NAR=<number of AR terms for regular ARCH>, NMA=<number of MA terms for GARCH>, RELAX, ZERO, nonlinear options) <dependent variable> [ <list of independent variables> ];

Usage

The ARCH command is just like the OLSQ command, except for the options which model the heteroskedasticity of the residuals. The default model estimated if no options are included is GARCH(1,1).

Method

The generalized form of ARCH (GARCH-M) estimated by TSP is given by the following equations (see McCurdy and Morgan):

\[ y_t = \sum_{i=1}^{N} \gamma_i x_{it} + \theta f(\lambda T_i) + \varepsilon_t \quad \varepsilon_t \sim N(0, h_t) \]

\[ f(h_t) = \frac{h_t^k}{k} \quad \text{if } \lambda \neq 0 \]

\[ f(h_t) = \log h_t \quad \text{if } \lambda = 0 \]

\[ h_t = \alpha_0 + \sum_{i=1}^{NAR} \alpha_i \varepsilon_{t-i}^2 + \sum_{j=1}^{NMA} \beta_j h_{t-j} + \sum_{k=1}^{NGT} \varphi_k g_{kt} \]

An immediate issue is the identification of the order of the ARCH or GARCH process. Bollerslev suggests obtaining squared residuals from OLS and using standard Box-Jenkins techniques (BJIDENT in TSP) on these squared residuals. It is also possible to estimate several ARCH models and use likelihood ratio tests to determine the proper specification.
All models are estimated by maximum likelihood (normally with analytic first and second derivatives). Presample values of \( h \) and the disturbance epsilon are initialized by the methods specified in the HINIT and E2INIT options.

ARCH will not work if there are gaps in the SMPL -- instead, it is possible to use dummy variables (as right hand side variables and in GT= ) to exclude observations from the fit.

Default starting values for \textit{gamma} are obtained from the OLS slope coefficients, while \( a(0) \) and presample \( h(t) \) (if HINIT=ESTALL is used) are started from the OLS ML estimate of the variance of the disturbance. Other parameters are initialized to 0. These defaults apply to all models except when NAR=0 and NMA>0, in which case \( a(0)=0 \) and \( b(1)=1 \) are used (to prevent false convergence to the OLS saddlepoint solution). The defaults may be overridden if an @START vector is provided by the user (with a value for each parameter -- see the final example below).

Several constraints are imposed on the ARCH parameters to protect against numerical problems from negative, zero, or infinite variances:

\[ 0 \leq \theta, \alpha, \beta, \leq 1 \]
\[ \alpha + \beta < 1 \text{ if HINIT=STEADY} \]
\[ \alpha_0 \geq 0 \]
\[ \varphi_k \geq 0 \]
\[ \text{presample } h_t \geq 0 \text{ if } \theta \text{ is not estimated.} \]
\[ \text{presample } h_t > 0 \text{ if } \theta \text{ is estimated.} \]
\[ h_t > 0 \]

Use of the ZERO/NOZERO option controls the technique for bounding (see the description under Options below). If a parameter is bounded on convergence, its standard error is set to zero to make the other estimates conditional on it. In this case, it is wise to try some alternative non-bounded starting values to check for an interior ML solution (see the final example).

\textbf{Output}

A title is printed, based on the options, indicating OLS, OLS-W, OLS-M, ARCH, ARCH-M, GARCH, or GARCH-M estimation. Standard iteration output follows with starting values, etc. Then standard regression output is printed; the only difference is the extra coefficients. The order of the estimated coefficients is:
\( \gamma, \theta, \alpha, \beta, \varphi \) \text{ presample } h(t) \\

\( h(t) \) is stored in @HT.

**Options**

- **E2INIT=HINIT** or INDATA or PREDATA specifies the initialization of the presample values of epsilon. The default HINIT sets them equal to \( h(t) \) (their unconditional expectation, as given by the current HINIT option). INDATA reserves the first NAR observations in the current sample to compute residuals and squares these (this was the default in TSP Version 4.3 and earlier). PREDATA attempts to use NAR observations prior to the current sample to compute such squared residuals (if such observations are missing, some observations from the current sample are used).

- **GT** = a list of weighting series \( g(k,t) \). The estimated coefficients for these series are labeled as PHI_series1, PHI_series2, etc. in the output. Note that the constant C should not be in the GT list because it is already included in \( h(t) \) via \( a(0) \). If GT is used without NAR or NMA, the model is called OLS-W or OLS-M by TSP. Use the RELAX option to relax the constraint on phi.

- **HEXP** = value of the exponent of the conditional variance \( h(t) \) in the regression equation. The default value of HEXP is 0.5, which means that the disturbance depends on the standard deviation of its distribution.

- **HINIT=ESTALL** or OLS or **SSR** or STEADY or value specifies the initialization of the presample values of \( ht \). ESTALL estimates them as nuisance parameters (labelled \( H(0), H(-1) \), etc.); this was the default in TSP Version 4.3 and earlier. OLS holds them fixed at the initial ML estimate of the residual variance from OLS. The default SSR sets them equal to SSR/T, where SSR is the sum of squared residuals from the current parameter values (see Fiorentini et al (1996)). STEADY sets them equal to their steady state value

\[
\begin{align*}
    h_0 &= \frac{\alpha_0}{1 - \sum_{i=NAR}^{NMA} \alpha_i + \sum_{j=1}^{NMA} \beta_j}
\end{align*}
\]

Note that the denominator must be strictly positive, and that the expectation of \( \text{phi}(k)g(k,t) \) is assumed to be zero. The user may also specify an arbitrary fixed value.
MEAN/NOMEAN controls whether the \textit{theta} \( f(h,t) \) term appears in the regression. This is labelled THETA in the output. MEAN indicates a GARCH-M, ARCH-M, or OLS-M model, and the GT, NAR, or NMA options are required. The constraints on \textit{theta} can be relaxed if necessary by using the RELAX option. See the HEXP option.

\textbf{NMA=} the number of terms where \( h(t) \) depends on its past values. These coefficients are labelled BETA1, BETA2, etc. in the output. This indicates a GARCH model.

\textbf{NAR=} the number of terms where \( h(t) \) depends on past squared residuals. These coefficients are labelled ALPHA1, ALPHA2, etc. in the output. This indicates a pure ARCH model if NMA=0.

\textbf{RELAX/NORELAX} relaxes the constraints on \textit{theta} and \textit{phi}.

\textbf{ZERO/NOZERO} specifies the method of imposing constraints on the parameters. If the default ZERO option is used, a parameter is set equal to the bound if the trial value violates the bound. Note that ZERO does not apply to parameters with strict inequality constraints, such as \( h(t) \). With NOZERO, the stepsize is squeezed when the bound is violated (until the constraint is met). NOZERO appears to be much slower than ZERO.

\textbf{Nonlinear options} These options control the iteration methods and printing. They are explained in the NONLINEAR section of this manual. Some of the common options are MAXIT, MAXSQZ, PRINT/NOPRINT, and SILENT/NOSILENT.

The default Hessian choices are HITER=N and HCOV=W. Other choices like B, F, and D are also legal, but Fiorentini et al (1996) show that HITER=N results in fast convergence, and HCOV=W yields standard errors that are robust to non-normal disturbances.

If no options are supplied, a GARCH(1,1) model will be estimated.

\textbf{Examples}

\begin{verbatim}
ARCH(NAR=4) Y C X;       # Pure ARCH
ARCH Y C X;              # GARCH(1,1) (the default)
ARCH(NMA=1,NAR=3,MEAN) Y C X; # GARCH-M
ARCH(GT=(G1,G2,G3),MEAN) Y C X; # OLS-M
\end{verbatim}

GARCH(0,1): Here we try some alternative starting values to override the solution with \textit{beta1}=1 that can occur in these models. The order of parameters in @COEF and @START is described below under Output -- the first 3 parameters are the \textit{gammas} and the last 3 are \textit{alpha0}, \textit{beta1}, and \( h(0) \).
Other types of univariate and multivariate GARCH models can be estimated with ML and ML PROC. Please see the TSP User's Guide and our web page http://www.tspintl.com for examples.

References


ASMBUG

Example

ASMBUG turns the DEBUG switch on during the first phase of TSP execution (when the input program is being processed). When the debug switch is on, TSP produces much more printed output than it usually does. This output is normally not of interest to users, but may be helpful to a TSP programmer or consultant.

ASMBUG

Usage

Include the ASMBUG statement directly before any command(s) you believe are being parsed incorrectly. The DEBUG switch will remain on until a NOABUG statement is encountered. While it is on, the internal representation of each statement and equation will be displayed as it is stored.

For DEBUG output during the execution phase of the program, see the DEBUG statement.

Output

ASMBUG produces a considerable amount of output, although not quite as much as the DEBUG statement. The breakdown of every statement into its constituent parts by INPT is printed; if the statement is a GENR, SET, IF, FRML, or IDENT, intermediate results from FRML and PARSE are also printed. Finally the interpreted form of the command is shown, exactly as it is stored in data storage for later execution.

Example

ASMBUG
FRML EQ Y = A**(EXP(GAMMA*X)) + B*LOG(HSQ*Z);
NOABUG

This example will cause various intermediate results during the parsing of the equation to be printed.
BJEST estimates the parameters of an ARIMA (AutoRegressive Integrated Moving Average) univariate time series model by the method of conditional or exact maximum likelihood. The technical details of the method used are described in Box and Jenkins, Chapter 7. TSP uses the notation of Box and Jenkins to describe the time series model. See BJIDENT for identification of a time series model and BJFRCST for forecasting using the estimated model.

\[ BJEST (CONSTANT, CUMPLOT, EXACTML, NAR=<number of AR parameters>, NBACK=<number of back-forecasted residuals>, NDIFF=<degree of differencing>, NLAG=<number of autocorrelations>, NMA=<number of MA parameters>, NSAR=<number of seasonal AR params>, NSDIFF=<degree of seasonal differencing>, NSMA=<number of seasonal MA parameters>, NSSPAN=<span of seasonal>, PLOT, PREVIEW, ROOTS, START, nonlinear options) [<series name>] [START <parameter name> <parameter value> ......] [FIX <parameter name> <parameter value> ......] [ZERO <parameter names> .....] [ZFIX <parameter names> .....] ; \]

Usage

To estimate an ARIMA model, use the BJEST command followed by the options you want in parentheses and then the name of the series and specification of the starting values, if you want to override the default starting values.

The general form of the model which is estimated is the following:

\[ I^r(B) \Phi(B)w(t) = \Delta(B) \Theta(B)a(t) + \text{constant} \]

\[ w(t) \] is the input series after ordinary and seasonal differencing, and \[ a(t) \] is the underlying "white noise" process. \[ B \] is the "backshift" or lag operator:

\[ Bw(t) = w(t-1) = Lw(t) \text{ in the usual notation} \]

\[ I^r(B), \Phi(B), \Delta(B), \Theta(B) \] are all polynomials in \[ B \].

The order of these polynomials is specified by the options NSAR, NAR, NSMA, and NMA respectively. \[ \text{Gamma}(B) \] and \[ \text{Delta}(B) \] are the seasonal AR and MA polynomials and \[ \text{Phi}(B) \] and \[ \text{Theta}(B) \] are the ordinary AR and MA polynomials. All the polynomials are of the form
BJEST

\[ \Phi(B) = 1 - \phi_1 B - \phi_2 B^2 - \phi_3 B^3 - \ldots \]

Note that the coefficients all have minus signs in front of them.

The BJEST procedure uses conditional sum of squares estimation to find the best values of the coefficients of these polynomials, consistent with the unobserved variable at being independently identically distributed. The options specify the exact model which is to be estimated and can also be used to control the iteration process.

As with the other iterative estimation techniques in TSP, it is helpful to specify reasonable starting values for the parameters. Unlike the other nonlinear TSP procedures, the starting values for BJEST can be specified directly in the command, not with a PARAM statement, since the parameters have certain fixed predetermined names in the Box-Jenkins notation.

There are several ways to specify starting values in BJEST. The easiest way is to let TSP "guess" reasonable starting values for the parameters. TSP will choose starting values, for the non-seasonal parameters only, based on the autocorrelations of the time series. You can suppress this feature with NOSTART. If no other information is supplied (see below), TSP will use zero as the starting value for all the parameters. In practice, this generally leads to slower convergence of the parameter estimates. You can also supply starting values in an @START vector.

You may also specify the starting values for any or all of the parameters yourself. For a model with more than one or two parameters, it may be difficult to choose starting values, since the individual parameters often do not possess a simple interpretation. However, you may wish to specify the starting values of at least some of the parameters based, perhaps, on previous estimates. In this case any parameters for which no starting value is supplied will be given the default value (TSP's guess if the START option is on, or zero for NOSTART, or the @START value).

User-supplied starting values are given after the name of the series on the command. Follow the series name with the keyword START and then a series of pairs of parameter names and starting values. The names available are:

- AR(lag) or PHI(lag) an ordinary autoregressive parameter.
- MA(lag) or THETA(lag) an ordinary moving average parameter.
- SAR(lag) or GAMMA(lag) a seasonal autoregressive parameter.
- SMA(lag) or DELTA(lag) a seasonal moving average parameter.
The lag in parentheses should be an actual number giving the position of the coefficient in the lag polynomial, i.e., AR(1) means the coefficient which multiplies the series lagged once (AR(0) is always unity).

Sometimes it is useful to fix a parameter at a certain value while others are being estimated. This may be done to limit the number of parameters being estimated, or to incorporate prior information about a parameter value, or to isolate an estimation problem which is specific to one or a few parameters. The keyword FIX followed by pairs of parameter names and values can be used to achieve this. FIX and START can be used on the same BJEST command to give starting values to some parameters and hold others fixed. The ZERO keyword is used to override the automatic starting values for some parameter(s) with zero(s). The ZFIX keyword fixes some parameter(s) to zero.

BJEST can also be used to check roots of polynomials for stationarity/invertability, without doing any estimation. Just supply the coefficient values in @START, use NAR=p and/or NMA=q, and do not supply a dependent variable. The PRINT option will print the roots, or the @ARSTAT and @MAINV variables can be used.

Output

The output of BJEST begins with a printout of the starting values, followed by an iteration log with one line per iteration giving the value of the objective function and the convergence criterion. If the PRINT option is on, the convert values of the options and the exact time series process being used are printed. In the iteration log, parameter values and changes are printed for each iteration when PRINT is on.

When convergence of the iterative process has been achieved or the maximum number of iterations reached, a message to that effect is printed, and the final results are displayed. These include the conventional statistics on the model: the standard error of at, the R-squared and F-statistic for the hypothesis that all the parameters are zero. The parameter estimates and their standard errors are shown in the usual regression output.

If the ROOTS option is on and the order of any polynomial is greater than one, its roots and moduli are shown so that you can check that they are outside the unit circle (as is required for stationarity). A table of the autocorrelations and Ljung-Box modified Q-statistics of the residuals is printed after this. If the PRINT option is on, the exact model estimated is again printed in lag notation, along with some summary statistics for the residuals. Then comes a printout of the lagged cross correlations between the differenced series wt and the white noise (residual) series at.
If the PLOT option is on, a time series plot of the residuals (at) is printed, using high resolution graphics in the Oxmetrics version. If the CUMPLOT option is on, a normalized cumulative periodogram is also plotted (see the CUMPLOT option above).

The following variables are stored (statistics based on differenced variable if differencing is specified):

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@LHV</td>
<td>list</td>
<td>1</td>
<td>Name of the dependent variable</td>
</tr>
<tr>
<td>@RNMS</td>
<td>list</td>
<td>#vars</td>
<td>Names of right hand side parameters</td>
</tr>
<tr>
<td>@SSR</td>
<td>scalar</td>
<td>1</td>
<td>Sum of squared residuals</td>
</tr>
<tr>
<td>@S</td>
<td>scalar</td>
<td>1</td>
<td>Standard error of the regression</td>
</tr>
<tr>
<td>@S2</td>
<td>scalar</td>
<td>1</td>
<td>Standard error squared</td>
</tr>
<tr>
<td>@YMEAN</td>
<td>scalar</td>
<td>1</td>
<td>Mean of the dependent variable</td>
</tr>
<tr>
<td>@SDEV</td>
<td>scalar</td>
<td>1</td>
<td>Standard deviation of the dependent variable</td>
</tr>
<tr>
<td>@DW</td>
<td>scalar</td>
<td>1</td>
<td>Durbin-Watson statistic</td>
</tr>
<tr>
<td>@RSQ</td>
<td>scalar</td>
<td>1</td>
<td>R-squared</td>
</tr>
<tr>
<td>@ARSQ</td>
<td>scalar</td>
<td>1</td>
<td>Adjusted R-squared</td>
</tr>
<tr>
<td>@IFCONV</td>
<td>scalar</td>
<td>1</td>
<td>=1 if convergence achieved, =0 otherwise</td>
</tr>
<tr>
<td>@LOGL</td>
<td>scalar</td>
<td>1</td>
<td>Log of likelihood function</td>
</tr>
<tr>
<td>@NCOEF</td>
<td>scalar</td>
<td>1</td>
<td>Number of coefficients</td>
</tr>
<tr>
<td>@NCID</td>
<td>scalar</td>
<td>1</td>
<td>Number of identified coefficients</td>
</tr>
<tr>
<td>@COEF</td>
<td>vector</td>
<td>#coefs</td>
<td>Coefficient estimates</td>
</tr>
<tr>
<td>@SES</td>
<td>vector</td>
<td>#coefs</td>
<td>Standard errors of coefficient estimates</td>
</tr>
<tr>
<td>@T</td>
<td>vector</td>
<td>#coefs</td>
<td>T-statistics on coefficients</td>
</tr>
<tr>
<td>@GRAD</td>
<td>vector</td>
<td>#coefs</td>
<td>Values of the gradient at convergence</td>
</tr>
<tr>
<td>@VCOV</td>
<td>matrix</td>
<td>#coefs*</td>
<td>Variance-covariance of estimated coefficients.</td>
</tr>
<tr>
<td>@QSTAT</td>
<td>vector</td>
<td>NLAG</td>
<td>Ljung-Box modified Q-statistics</td>
</tr>
<tr>
<td>%QSTAT</td>
<td>vector</td>
<td>NLAG</td>
<td>P-values for Q-statistics</td>
</tr>
<tr>
<td>@ARSTAT</td>
<td>scalar</td>
<td>1</td>
<td>1 if AR polynomial is stationary</td>
</tr>
<tr>
<td>@MAINV</td>
<td>scalar</td>
<td>1</td>
<td>1 if MA polynomial is invertible</td>
</tr>
<tr>
<td>@FIT</td>
<td>series</td>
<td>#obs</td>
<td>Fitted values of the dependent variable</td>
</tr>
<tr>
<td>@RES</td>
<td>series</td>
<td>#obs</td>
<td>Residuals=actual - fitted values of the dependent variable</td>
</tr>
<tr>
<td>@ARRTRE</td>
<td>vector</td>
<td>NAR</td>
<td>Real parts of the AR roots</td>
</tr>
<tr>
<td>@ARRTIM</td>
<td>vector</td>
<td>NAR</td>
<td>Imaginary parts of the AR roots</td>
</tr>
</tbody>
</table>
Method

The method used by BJEST (for the default method NOEXACTML) to estimate the parameters is essentially the one described by Box and Jenkins in their book. It uses a conventional nonlinear least squares algorithm with numerical derivatives. The major difference between the estimation of time series models and estimation in the traditional (nonlinear least squares) way relates to the use of "back-forecasted" residuals. The likelihood function for the ARIMA model depends on the infinite past sequence of residuals. If we estimate the time series model by simply setting the values of these past residuals to zero, their unconditional expectation, we might seriously misestimate the parameters if the initial disturbance, a0, happens to be very different from zero.

The solution to this problem, suggested by Box and Jenkins, is to invert the representation of the time series process, i.e., write the same process as if the future outcomes were determining the past. Thus it describes the relationships which the time series will, ex post, exhibit. This representation of the backward process constructs back-forecasts of the disturbance series, the at series and then uses these calculated residuals in the likelihood function. By using a reasonable number of these backcasted residuals, the problems introduced by an unusually high positive or negative value for the first disturbance in the time series can be eliminated.

If the process is a pure moving average process, this backcast becomes zero after a fixed number of time periods; consequently, you can set NBACK to a fairly small number in this case.

When the EXACTML option is used, no backcasting is done; the AS 197 algorithm (Melard 1984) is used.
BJEST

HCOV=U (numeric second derivatives) is the default method of computing the standard errors. For iteration, HITER=F is the default, but HITER=U can be chosen as an option.

Options

Note that for all the Box-Jenkins procedures (BJIDENT, BJEST, and BJFRCST), TSP remembers the options from the previous Box-Jenkins command (except for nonlinear options), so that you only need to specify the ones you want to change.

CONSTANT/NOCONS specifies whether a constant term is to be included in the model.

CUMPLOT/NOCUMPLOT specifies whether a cumulative periodogram of the residuals is to be plotted. The number of computations required for this plot goes up with the square of the number of observations, so that it may be better to forego this option if the number of observations is large.

EXACTML/NOEXACT specifies exact (versus conditional) maximum likelihood estimation. EXACTML is recommended for models with a unit root in the MA polynomial.

NAR= the number of autoregressive parameters in the model. The default is zero.

NBACK= the number of back-forecasted residuals to be calculated. The default is 100.

NDIFF= the degree of differencing to be applied to the series. The default is zero.

NLAG= the number of autocorrelations/Q-statistics to calculate. The default is 20.

NMA= the number of moving average parameters to be estimated. The default is zero.

NSAR= the number of seasonal autoregressive parameters to be estimated. The default is zero.

NSDIFF= the degree of seasonal differencing to be applied to the series. The default is zero (no differencing).

NSMA= the number of seasonal moving average parameters to be estimated. The default is zero.
BJEST

**NSPAN**= the span (number of periods) of the seasonal cycle, i.e., for quarterly data, NSPAN should be 4. The default is the current frequency (that is, 1 for annual, 4 for quarterly, 12 for monthly).

**PLOT/NOPLOT** specifies whether the residuals are to be plotted.

**PREVIEW/NOPREVIEW** *(TSP/Oxmetrics only)* specifies that the residual plots are to be displayed in a high-resolution graphics window.

**PRINT/NOPRINT** specifies the level of output desired. NOPRINT should be adequate for most purposes.

**RELAX/NORELAX** specifies whether the AR stationarity and MA invertibility constraints should be turned off during EXACTML iterations.

**ROOTS/NORoots** specifies whether the roots of the polynomial should be printed.

**START/NOSTART** specifies whether the procedure should supply its own starting values for the parameters.

Nonlinear options control iteration and printing. They are explained in the **NONLINEAR** entry.

One special use of these options in BJESt is the combination EXACTML, NOCONST, MAXIT=0, MAXSQZ=123, which forces the program to simply evaluate the ARMA likelihood at the starting values of the parameters in @START.

**Examples**

This example estimates a simple ARMA(1,1) model with no seasonal component; no plots are produced of the results.

**BJEST (NAR=1,NMA=1,NOPLOT,NOCUMPLO) AR9MA5 ;**

This example uses the Nelson (1973) auto sales data; a logarithmic transformation of the series is made before estimation.

**GENR LOGAUTO = LOG(AUTOSALE) ;**

**BJEST (NDIF=1,NSDIFF=1,NMA=2,NSMA=1,NSPAN=12, NBACK=15) LOGAUTO**

**START THETA(1) 0.12 THETA(2) 0.20 DELTA(1) 0.82 ;**

Note that NBACK is specified as 15 since the backcasted residuals fall to exactly zero after NSPAN+NSMA+NMA periods in a pure moving average model.
The next example estimates a third order autoregressive process with one parameter fixed:

\texttt{BJEST GNP START PHI(2) -0.5 PHI(3) 0.1 FIX PHI(1) 0.9 ;}

The model being estimated is

\[ GNP(t) - 0.9*GNP(t-1) - 2*GNP(t-2) - 3*GNP(t-3) = a(t) \]

Exact ML estimation:

\texttt{MMAKE @START .1 .1 .1;}

\texttt{BJEST(NAR=2,NMA=1,NDIFF=1,EXACTML) Y;}

References


Statlib, \texttt{http://lib.stat.cmu.edu/apstat/}
BJFRCST calculates the forecasts of a time series that are implied by an ARIMA model (see the description of BJEST for more information of ARIMA models). The forecasts and other statistics are calculated using the equations in Chapter 5 and Part V, program 4 of Box and Jenkins.

**Usage**

The entry for the BJEST procedure describes the ARIMA model and its notation in some detail. To obtain forecasts for such a model, use the BJFRCST command followed by the options you want (in parentheses) and then the name of the series.

If the BJFRCST command immediately follows the BJEST command for the desired model, no parameter values need to be specified; BJFRCST will automatically use the final estimates generated by BJEST. Alternatively, any or all of the model parameters can be specified by listing the parameters and their values following the series name (as in BJEST). If no value is supplied for S, the standard error of the disturbance, (and if no estimate from BJEST is available), BJFRCST will calculate an estimate of S using the values of the time series up to and including the ORGBEG-th observation.
BJFRCST calculates forecasts for the time series for the NHORIZ periods following the current "origin". The "origin" is the final period for which (conceptually) the time series is observed, i.e., the final period for which no forecast is required. In practice, it is often useful to generate forecasts for periods that have already been observed. These forecasts provide evidence on the reliability of the estimated time series model. It is also useful to generate sets of forecasts for different origins. BJFRCST calculates a complete set of forecasts and associated confidence bounds for each of the origins from ORGBEG to ORGEND inclusive.

Output

The output of BJFRCST starts with a printout of the option settings. The parameter values are also printed. This is followed by a printout of the expanded \( \Phi^* \) and \( \Theta^* \) polynomials implied by these parameters. The \( \Phi^* \) and \( \Theta^* \) polynomials are the generalized autoregressive and moving average polynomials, respectively, that are obtained by taking the product of the ordinary, seasonal, and pure difference polynomials. Next a table is printed of the forecast standard errors, and the psi weights. Note that the standard error in the N-th row of the table is the standard error of the forecast for the N-th period following the origin (the “N-th step ahead”). The psi weights are the coefficients obtained by expressing the model as a pure moving average. The N-th row entry for \( \Psi, \Phi^* \) or \( \Theta^* \) gives the coefficient for the N-th lag.

The next feature of the output is a table of the forecasts, their upper and lower confidence bounds, the actual values of the time series (when available), and the discrepancies between the forecasts and the realizations of the time series. If PLOT is specified, the values in this table are plotted, in high resolution graphics in the Oxmetrics version. The following variables are stored:

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@FIT</td>
<td>series</td>
<td>NHORIZ</td>
<td>Forecast (for last origin, if there was more than one)</td>
</tr>
<tr>
<td>@FITSE</td>
<td>series</td>
<td>NHORIZ</td>
<td>Forecast standard errors (still in logs for EXP option)</td>
</tr>
</tbody>
</table>

Method
BJFRCST forms the generalized autoregressive and moving average lag polynomials, $\phi^*$ and $\theta^*$. These lag polynomials define a simple difference equation for the time series. The forecasts are generated using this difference equation. As in BJEST, backcasted residuals are calculated to obtain initial conditions for the difference equation. Chapter 5 of Box and Jenkins (1976) contains a detailed explanation and analysis of this procedure.

Options

Note that for all the Box-Jenkins procedures (BJIDENT, BJEST, BJFRCST), TSP remembers the options from the previous Box-Jenkins command, so that you only need to specify the ones you want to change.

CONBOUND= specifies the probability for the confidence bounds about the forecasts. The default is 0.95.

CONSTANT/NOCONST specifies whether there is a constant term in the model.

EXP/NOEXP is used when BJEST was used to model log(y), and you want to forecast the original y. The forecast is increased by using the forecast variance for each observation, to make it unbiased, and the forecast confidence interval will be asymmetric. See the Nelson reference for more details.

NAR= the number of ordinary autoregressive parameters in the model. The default is zero.

NBACK= the number of back-forecasted residuals to be calculated. The default is 100.

NDIFF= the degree of ordinary differencing required to obtain stationarity. Note that the undifferenced series is forecasted. The default is zero.

NHORIZ= the number of periods ahead to forecast the series. The default is 20.

NLAG= the number of periods prior to the forecasting origin to include in the plot of the series and the forecasts. The default is 20.

NMA= the number of ordinary moving average parameters in the model. The default is zero.

NSAR= the number of seasonal autoregressive parameters in the model. The default is zero.
**BJFRCST**

**NSDIFF=** the degree of seasonal differencing required to obtain stationarity. Note that the undifferenced series is forecasted. The default is zero.

**NSMA=** the number of seasonal moving average parameters in the model. The default is zero.

**NSSPAN=** the span (length) of the seasonal cycle. For example, with quarterly data, NSSPAN should be 4. The default is the current frequency (that is, 1 for annual, 4 for quarterly, 12 for monthly). If the current FREQ is 0, the default is the value from the last BJ command.

**ORGBEG=** the first origin to use in forecasting. The default is the final observation in the current sample. This option should be specified as a TSP date, e.g., 82:4. Note that this date must lie within the current SMPL.

**ORGEND=** the final origin to use in forecasting. The default is ORGBEG. This option should also be specified as a TSP date. Like ORGBEG, this date must lie within the current SMPL.

**PLOT/NOPLOT** specifies whether the series and the forecasts are plotted.

**PREVIEW/NOPREVIEW** (TSP/Oxmetrics only) specifies that the forecasts are to be displayed in a high-resolution graphics window.

**PRINT/NOPRINT** specifies whether the results will be printed (versus just stored in @FIT).

**RETRIEVE/NORETR** specifies whether BJFRCST should try to find parameter values from an immediately preceding BJEST procedure.

**SILENT/NOSILENT** suppresses all the output (SILENT is equivalent to NOPLOT, NOPRINT)

**Examples**

The first example produces one set of forecasts for the Nelson auto sales data. The log of the original series is forecasted.

```
BJFRCST (PRINT,NMA=2,NSMA=1,NDIFF=1,NSDIFF=1,NSSPAN=12,
       NHORIZ=24,ORGBEG=264) LOGAUTO S 0.111068 THETA(1)
   0.2114 THETA(2) 0.2612 DELTA(1) 0.8471 ;
```

The second example produces the same forecasts, but the model parameters are provided by the immediately preceding BJEST procedure. Notice that the options describing the model do not need to be repeated for the BJFRCST procedure. Also, the level of the original series, rather than its log, are forecasted.
BJFRCST

BJEST (NMA=2,NSMA=1,NDIFF=1,NSDIFF=1,NSSPAN=12) LOGAUTO
START THETA(1) 0.12 THETA(2) 0.20 DELTA(1) 0.82 ;
BJFRCST (EXP,PRINT,NHORIZ=24,ORGBEG=264) LOGAUTO ;

References


BJIDENT prints and plots descriptive statistics which are useful in identifying the process which generated a time series. The process of time series identification is described in the two references; TSP follows the notation of Box and Jenkins, who developed this technique for analyzing time series. BJEST is used for estimating the model you develop and BJFRCST for forecasting with it.

The decisions to be made in the process of identifying a time series process are 1) whether there is a seasonal component, 2) how much ordinary and seasonal differencing is required to make the series stationary, 3) and what are the minimum orders of the autoregressive and moving average polynomials required to explain adequately the time series. Subject to various option settings, BJIDENT will present plots of autocorrelations and partial autocorrelations for various levels of differencing of the input series.

**Usage**

BJIDENT followed by the name of a series is the simplest form of the command. No differencing of the series will be done in this case, and the output will consist of a plot of the series, and a printout and plot of the autocorrelations of the first 20 lags of the series and the first 10 partial autocorrelations.

Autocorrelations are the correlation of the series with its own values lagged once, lagged twice, and so forth for 20 lags. Partial autocorrelations are the correlations measured with the series residual after all the prior lags have been removed. That is, the second partial autocorrelation is the correlation of the series lagged twice with the part of the series which is orthogonal to the first lag.

The options on the BJIDENT command allow you to specify the differencing you want performed on the series, whether there is a seasonal component, and the output you wish to see plotted or printed. You can also increase and decrease the number of autocorrelations which are computed.
If you want to analyze the log or other transformation of your original series, as suggested by Box and Jenkins or Nelson, transform the series using a GENR statement before submitting it to BJIDENT.

**Output**

The output of BJIDENT begins with the plots of the raw and differenced series (if the PLOT option was requested). These plots are high resolution graphics plots in TSP/Oxmetrics, and low-resolution in other versions. Next a table of the autocorrelations is printed with their standard errors and the Ljung-Box portmanteau test, or modified Q-statistic. These modified Q-statistics are distributed independently as chi-squared random variables with degrees of freedom equal to the number of autocorrelations. The null hypothesis is that all the autocorrelations to that order are zero.

Following this table, the partial autocorrelations are printed for each series. If PLOTAC has been specified, BJIDENT then plots the autocorrelation and partial autocorrelation functions of the series and its differences with standard error bands. The inverse autocorrelations are printed if requested. Finally, the ESACF correlations, their p-values, and a table of Indicators is printed. If the PRINT option is on, a table of AR coefficient estimates is printed. The following matrices are stored:

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@AC</td>
<td>matrix</td>
<td>NLAG*#diff</td>
<td>autocorrelations</td>
</tr>
<tr>
<td>@PAC</td>
<td>matrix</td>
<td>NLAGP*#diff</td>
<td>partial autocorrelations</td>
</tr>
<tr>
<td>@IAC</td>
<td>matrix</td>
<td>NLAGP*NLAGP</td>
<td>inverse autocorrelations if requested</td>
</tr>
<tr>
<td>@ESACF</td>
<td>matrix</td>
<td>NAR*NMA</td>
<td>ESACF correlations</td>
</tr>
<tr>
<td>%ESACF</td>
<td>matrix</td>
<td>NAR*NMA</td>
<td>p-values for ESACF correlations</td>
</tr>
<tr>
<td>@PHI</td>
<td>matrix</td>
<td>(NAR*(NAR+1)/2 x (3+NMA) ESACF</td>
<td></td>
</tr>
<tr>
<td>@ESACFI</td>
<td>matrix</td>
<td>NAR*NMA</td>
<td>ESACF Indicators</td>
</tr>
</tbody>
</table>

**Options**

*Note that for all the Box-Jenkins procedures (BJIDENT, BJEST, and BJFRCST), TSP remembers the options from the previous Box-Jenkins command, so that you only need to specify the ones you want to change.*

**BARTLETT/NOBART** specifies that the Bartlett estimate using lower order autocorrelations is to be used for the variance of the ESACF option. NOBART will simply use 1/(T-p-q).
BJIDENT

ESACF/NOESACF computes the extended sample ACF of Tsay and Tiao (1984). This can be useful for identifying stationary and nonstationary ARMA models. The upper left vertex of a triangle of zeroes in the Indicator matrix identifies the order of the ARMA model. The zeroes correspond to nonsignificant autocorrelations. See the examples.

IAC/NOIAC specifies whether the inverse autocorrelations are to be computed and printed.

NAR= maximum order of AR for ESACF. Default is 20.

NDIFF= the degree of differencing to be applied to the series. The default is zero (no differencing). BJIDENT will calculate statistics for all the differences of the series up to and including the NDIFFth order.

NLAG= the number of autocorrelations to be computed. The default is 20.

NLAGP= the number of partial autocorrelations to be computed. The default is 10.

NMA= maximum order of MA for ESACF. Default is 10.

NSDIFF= the degree of seasonal differencing to be applied to the series. The default is zero (no differencing). As in the case of ordinary differencing, BJIDENT will calculate statistics for all the differences of the series up to and including the NSDIFFth order.

NSPAN= the span (number of periods) of the seasonal cycle, i.e., for quarterly data, NSPAN should be 4. The default is the current frequency (that is, 1 for annual, 4 for quarterly, 12 for monthly).

PLOT/NO PLOT specifies whether all the differenced series are to be plotted.

PLOTAC/NO PLOTAC specifies whether the autocorrelations and partial autocorrelations are to be plotted.

PLTRAW/NO PLTRAW specifies whether the original "raw" series is to be plotted.

PREVIEW/NO PREVIEW (TSP/Oxmetrics only) specifies that the raw and differenced series are to be displayed in a high-resolution graphics window if the PLOT option is on.

PRINT/NO PRINT specifies whether the AR coefficients for the ESACF option are to be printed.

SILENT/NO SILENT Turns off all output. Results are still stored in @AC, @PAC, etc.
Example

This example computes the auto sales example from Nelson's book:

```
BJIDENT (IAC, NDIFF=1, NSDIFF=1, NSSPAN=12, NLAG=48, NLAGP=20) AUTOSALE ;
```

The following example shows the ESACF output for Box-Jenkins Series C:

```
BJIDENT (ESACF, NAR=5, NMA=8) CHEM ;
```

The result of the above command is the following matrix:

<table>
<thead>
<tr>
<th></th>
<th>0</th>
<th>1</th>
<th>2</th>
<th>3</th>
<th>4</th>
<th>5</th>
<th>6</th>
<th>7</th>
<th>8</th>
</tr>
</thead>
<tbody>
<tr>
<td>0</td>
<td>9</td>
<td>9</td>
<td>9</td>
<td>9</td>
<td>9</td>
<td>9</td>
<td>9</td>
<td>9</td>
<td>1</td>
</tr>
<tr>
<td>1</td>
<td>9</td>
<td>9</td>
<td>9</td>
<td>9</td>
<td>9</td>
<td>1</td>
<td>1</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>2</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>3</td>
<td>9</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>4</td>
<td>9</td>
<td>9</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>5</td>
<td>9</td>
<td>9</td>
<td>9</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
</tbody>
</table>

A triangle of zeroes with upper left vertex at (2,0) is seen; this indicates an ARMA(2,0) model.

References


CAPITL

Computes a capital stock series from a gross investment series, using a perpetual inventory and a constant rate of depreciation. If I is gross investment, K is the capital stock, and d is the rate of depreciation then CAPITL computes

\[ K(t) = (1-d)K(t-1) + I(t-1) \quad \text{(for the NOEND option)} \]
\[ K(t) = (1-d)K(t-1) + I(t) \quad \text{(for the END option)} \]

It starts from a capital stock benchmark specified by an observation. If the benchmark is in the middle of the sample, CAPITL also applies the backward version of the formula,

\[ K(t) = (K(t+1) - I(t))/(1-d) \quad \text{(for the NOEND option)} \]
\[ K(t) = (K(t+1) - I(t+1))/(1-d) \quad \text{(for the END option)} \]

to compute values of the capital stock in periods before the benchmark. Note that the depreciation rate, d, is stated as a rate applicable to the frequency of the data. For example, for quarterly data, the depreciation rate must be a quarterly rate, one fourth of the annual rate. It is possible to compute a depreciation series, dK(t), and a net investment series, I(t)-dK(t).

CAPITL may be used in any application where a moving average with geometrically declining weights needs to be calculated. Further, by setting the depreciation rate to zero, it will simply cumulate a series.

\[ \text{CAPITL (BENCHOB=obid, BENCHVAL=scalar, END) <investment series> <depreciation rate> <capital stock series>;} \]

Usage

The only arguments required for the CAPITL statement are an investment series, a depreciation rate, and the name to be given to the derived capital stock series. In this case the value of the capital stock at the beginning of the computation is assumed to be zero. To alter these assumptions, see the options below.

CAPITL requires that there be no gaps in the current SMPL.

Output

CAPITL produces no printed output. The capital stock series is stored in data storage.
Options

**BENCHOBS**= an observation identifier for the benchmark observation. This identifier should be contained in the current SMPL. If the frequency is quarterly and the SMPL is 47:4 80:4, for example, the benchmark observation could be 47:4, 56:1, 80:4, etc.

**BENCHVAL**= the value of the capital stock series at the benchmark observation. CAPITL will compute the capital stock both forwards and backwards from this observation.

**END/NOEND** computes the end-of-period capital stock (see the previous formulas).

Examples

**SMPL 1,74;**
**CAPITL (BENCHVAL=145.4,BENCHOBS=4) INV,.04,KSTOCK ;**

In this example, the gross investment series is INV. CAPITL computes capital, KSTOCK. The benchmark applies to the 4th observation and has the value 145.4. The rate of depreciation is .04.

**CAPITL(BENCHOBS=1,BENCHVAL=X(1),END) X,0.0,XACCUM ;**

This example simply sums the series X and stores the result in XACCUM. Note that since the formula gives the end of period capital stock, the last observation of XACCUM contains the sum of all the observations on X.
CDF calculates and prints tail probabilities (P-values or significance levels) or critical values for several cumulative distribution functions. This is useful for hypothesis testing.

CDF (BIVNORM or CHISQ or DICKEYF or F or NORMAL or T or WTDCHI, CONSTANT, DF=<degrees of freedom for CHISQ or F>, DF1=<numerator degrees of freedom for F>, DF2=<denominator degrees of freedom for F>, EIGVAL=<vector of eigenvalues for WTDCHI>, LOWTAIL or UPTAIL or TWOTAIL, INVERSE, NLAGS=<number of lags for augmented Dickey-Fuller test>, NOB=<number of observations for unit root or cointegration test>, NVAR=<number of variables for cointegration test>, PRINT, RHO=<correlation coefficient for BIVNORM>, TREND, TSQ) <test statistic> [<significance level>];

or

<significance level> [critical value]; (for INVERSE)

or

<x value> <y value> [significance level]; (for BIVNORM)

Usage

CDF followed by the value of a scalar test statistic is the simplest form of the command. In this case, a two-tailed probability for the normal distribution will be calculated and printed. If the INVERSE option is used, the first argument must be a probability level; a critical value will be calculated. Arguments need not be scalars; they can be series or matrices. Distributions other than the normal and/or a choice of tail areas may be selected through the options. For hypothesis testing using a wide variety of regression diagnostics, see the REGOPT (PVPRINT) command.

Output

If the PRINT option is on, the input and output values will be printed, along with the degrees of freedom. If a second argument is supplied, it will be filled with the output values and stored (see examples 4 and 6). Input and output arguments may be any numeric TSP variables.

Method

BIVNORM: ACM Algorithm 462. Inverse is not supplied because it is not unique unless x or y is known, etc.
CHISQ: DCDFLIB method: Abramovitz-Stegun formula 26.4.19 converts it to Incomplete Gamma, and then use DiDinato and Morris (1986). Inverse by iteration (trying values of x to yield p -- faster methods are also known). Non-integer degrees of freedom are allowed, and can be used to compute the incomplete gamma function.

F and T: DCDFLIB method: Abramovitz-Stegun formula 26.6.2 converts it to Incomplete Beta, and then use DiDinato and Morris (1993), i.e. ACM Algorithm 708. Inverse by iteration. Non-integer degrees of freedom for F are allowed, and can be used to compute the incomplete beta function.


DICKEYF: Asymptotic values from Tables 3 and 4 in MacKinnon (1994). Finite sample critical values from Cheung and Lai (1995) [augmented Dickey-Fuller] or MacKinnon (1991) [Engle-Granger]. To convert these to finite sample P-values, a logistic interpolation is used with the .05 size and either the .01 or .10 size (whichever is closer to the observed test statistic). Such interpolated P-values are fine for testing at the .01, .05, or .10 sizes, but would be highly speculative outside this range.

WTDCHI: If \( w(i) \) are the eigenvalues (supplied by the option EIGVAL), \( c(i) \) are chi-squared(1) variables, and \( d \) is the test statistic (supplied as an argument), WTDCHI computes the following probability:

\[
\Pr\left\{ \sum_{i=1}^{N} \frac{w(i)c(i)}{\sum_{i=1}^{N} w(i)} < d \right\} = \Pr\left\{ \sum_{i=1}^{N} (w(i) - d)c(i) < 0 \right\}
\]

This is useful for computing P-values for the Durbin-Watson statistic, other ratios of quadratic forms in normal variables, and certain non-nested tests (for example, Vuong (1989) suggests likelihood ratio tests for nonnested hypotheses). The Pan method is used when the number of eigenvalues is less than 90; otherwise the Imhoff method is used. If the absolute values of the smallest eigenvalues are less than 1D-12, they are not used; otherwise duplicate eigenvalues are not checked for. The inverse of this distribution is not implemented.

Options

BIVNORM/CHISQ/DICKEYF/F/NORMAL/T/WTDCHI specifies the bivariate normal, chi-squared, Dickey-Fuller, F, standard normal, student’s t, and weighted chi-squared distributions, respectively.
**CDF**

**CONSTANT/NOCONST** specifies whether a constant term (C) was included in the regression for Dickey-Fuller. NOCONST is only valid for NVAR=1.

**DF**= the degrees of freedom for the chi-squared or student's t distribution, or the number of observations for Dickey-Fuller (also see the NOB= option for Dickey-Fuller). Non-integers allowed for the chi-squared.

**DF1**= the numerator degrees of freedom for the F distribution (can be non-integer).

**DF2**= the denominator degrees of freedom for the F distribution (can be non-integer).

**EIGVAL**= vector of eigenvalues for the weighted chi-squared distribution.

**INVERSE/NONVERSE** specifies the inverse distribution function (input is significance level, output is critical value). Normally the input is a test statistic and the output is a significance level. INVERSE is not defined for bivariate normal.

**LOWTAIL/TWOTAIL/UPTAIL** specifies the area of integration for the density function. TWOTAIL is the default for most symmetric distributions (normal and t), UPTAIL is the default for chi-squared and F, and LOWTAIL is the default for bivariate normal and Dickey-Fuller. TWOTAIL is not defined for bivariate normal.

**NLAGS**= the number of lagged differences in the augmented Dickey-Fuller test. This number is used to compute the approximate finite sample P-value or critical value. The default is zero (assume an unaugmented test). The NOB= option must also be specified for the finite sample value.

**NOB**= the number of observations for the augmented Dickey-Fuller or Engle-Granger tests. This number is used to compute the approximate finite sample P-value or critical value. The default is zero (to compute asymptotic instead of finite sample value).

**NVAR**= the number of variables for an Engle-Granger/Dickey-Fuller cointegration test. The default is 1 (plain unit root test), and the maximum is 6.

**RHO**= the correlation coefficient for the bivariate normal distribution.

**TREND/NOTREND** specifies whether a trend term (1,2,...,T) was included in the regression for Dickey-Fuller.

**TSQ/NOTSQ** specifies whether a squared trend term (1,4,9,...) was included in the regression for Dickey-Fuller.
CDF

PRINT/NOPRINT turns on printing of results. PRINT is true by default if there is no output specified.

Examples

1. To compute the significance level of a Hausman test statistic with 5 degrees of freedom:

\textit{CDF (CHISQ,DF=5) HAUS;}

produces the output:

CHISQ(5) Test Statistic: 7.235999 , Upper tail area: .20367

2. Significance level of the test for AR(1) with lagged dependent variable(s):

\textit{CDF @DHALT;}

\textit{or}

\textit{REGOPT (PVPRINT) DHALT;}

before the regression is run

3. Two-tailed critical value for the normal distribution:

\textit{CDF (INV) .05;}

produces the output:

NORMAL Critical value: 1.959964 , Two-tailed area: .05000

4. Several critical values for the normal distribution:

\textit{READ PX; .1 .05 .01 ;}

\textit{CDF(INV,NORM) PX CRIT;}

\textit{PRINT PX,CRIT;}

5. F critical values:

\textit{CDF(INV,F,DF1=3,DF2=10) .05;}

produces the output:

F(3,10) Critical value: 3.70827 , Upper tail area: .0500

6. Bivariate normal:
CDF

CDF(BIVNORM,RHO=.5,PRINT) -1 -2 PBIV;

produces the output:

BIVNORM Test Statistic: -1.0000 , -2.0000 , Lower tail area: .01327

7. Dickey-Fuller unit root test:

TREND TIME;
SMPL 2,50;
DY = Y-Y(-1);
OLSQ DY TIME C Y(-1);
CDF(DICKEYF) @T(3);
? The above is equivalent to the following:
UNIT(MAXLAG=0,NOWS) Y;

8. Augmented Dickey-Fuller unit root test, with finite sample P-value:

TREND TIME;
SMPL 2,50;
DY = Y-Y(-1);
SMPL 5,50;
OLSQ DY TIME C Y(-1) DY(-1)-DY(-3);
CDF(DICKEYF,NOB=@NOB,NLAGS=3) @T(3);
? The above is equivalent to the following:
UNIT(MAXLAG=3,NOWS,FINITE) Y;

9. Engle-Granger cointegration test:

TREND TIME;
OLSQ Y1 TIME C Y2 Y3 Y4; EGTEST;
OLSQ Y2 TIME C Y1 Y3 Y4; EGTEST;
OLSQ Y3 TIME C Y1 Y2 Y4; EGTEST;
OLSQ Y4 TIME C Y1 Y2 Y3; EGTEST;
PROC EGTEST;
SMPL 2,50;
DU = @RES-@RES(-1);
OLSQ DU @RES(-1);
CDF(DICKEYF,NVAR=4) @T;
SMPL 1,50;
ENDPROC;
? The above is equivalent to the following:
COINT (NOUNIT,MAXLAG=0,ALLORD) Y1-Y4;

10. Verify critical values for Durbin-Watson statistic, for regression with 10 observations and 2 RHS variables:
SMPL 1,10; OLSQ Y C X1;
SET PI = 4^ATAN(1); SET F = PI/(2*@NOB); TREND I;
EIGB = 4*SIN(I*F)**2;
SELECT I <= (@NOB-@NCOEF); ? use largest eigenvalues for dL
CDF (WTDCHI,EIG=EIGB) .879; ? dL for 5% (n=10, k=1)
? use smallest eigenvalues for dU
SELECT (@NCOEF <= I) & (I <= (@NOB-1));
MMAKE dU 1.320 1.165 1.001; ? dU for 5%, 2.5%, 1% (n=10, k=1)
CDF (WTDCHI,EIG=EIGB,PRINT) dU;

11. Reproduce exact P-value for Durbin-Watson statistic (this can be done automatically using REGOPT):

SMPL 1,10;
READ Y X1; 4 2 7 4.5 6 4 3 2 1 3 2 5 3 4.5 4 7.5 8 5 6 ;
REGOPT (DWPVAL=EXACT);
OLSQ Y C X1;
MMAKE X @RNMS;
MAT XPXI = (X=X)@;
TREND OBS; SELECT OBS > 1;
DC = 0; DX1 = X1 - X1(-1);
MMAKE DX DC DX1;
MMAKE BVEC 2 -1; MFORM(BAND,NROW=@NOB) DDP = BVEC;
MAT DMDP = DDP - DX*XPXI*DX=;
? Eigenvalues of DMDP = D^D = - DX*(X=X)@(DX)=
? (same as nonzero eigenvalues of MA, because A = D=D)
MAT ED = EIGVAL(DMDP);
CDF (WTDCHI,EIG=ED) @DW;

References


CLEAR (interactive)

CLEAR terminates an interactive session, and restarts TSP.

*CLEAR ;*

Usage

CLEAR is an alternative to *QUIT*. Use CLEAR when you want to immediately restart TSP without saving any of your data or commands from the current session. Use QUIT when you want to exit completely from TSP. CLEAR would be preferred if you have just been experimenting with some commands, or if you have created some variables in error, and you want to start with a clean slate.
CLOSE

Example

CLOSE closes a file which has been opened by the READ or WRITE command. Subsequent READ statements will read from the beginning of the file, or subsequent WRITE statements will create a new file. CLOSE can also be used to control access to more than 12 files in a single run, or to insure that a newly updated file will be complete in case the program or computer aborts later.

CLOSE (UNIT=unit number, FILE='filename string');

Usage

When TSP processes a READ or WRITE statement which accesses a file, the file is left open so that further READ or WRITE statements will read data from or write data to the next line after those already processed. This is useful when more observations or variables will be read from or written to the file later in the program.

There are several cases in which it may be useful to close the file:

1. During an interactive session, if an error is made reading or writing data, closing the file will allow correction of the error when the corrected READ or WRITE statements are executed. Without the CLOSE command, a new READ statement would read from the current file position, usually causing an "end of file" error message, while a new WRITE statement would append to the lines already written to the file in error (if any).

2. User-controlled access to more than 12 files in a given run is possible with CLOSE. Since the number of simultaneously open files is limited on most operating systems (often it is less than 12), TSP will close the most recently opened file and issue a warning message when access to a new file would result in too many simultaneously open files. If this arbitrary choice of the file to close causes problems with your program, use the CLOSE statement to reduce the number of simultaneously open files.

3. If results from a repeated iterative estimation process are to be saved repeatedly in a file, the CLOSE command could be used to cause repeated creation of the file instead of appending the new results each time to the file. It would be slightly easier to use the OUT command for this type of problem:

OUT databank; KEEP variables; OUT;
4. If important data has been written to a file, and it is likely that later commands may cause TSP to abort (or power failures may occur with the computer), the file may be closed to guarantee that the data is completely written.

For whatever reason, after the READ or WRITE statement, issue the CLOSE command, specifying the file either with a filename string or with a unit number in the options with the parentheses.

Example

```
READ (FILE='FOO.DAT') X Y Z;
CLOSE (FILE='FOO.DAT');
```
COINT performs unit root and cointegration tests. These may be useful for choosing between trend-stationary and difference-stationary specifications for variables in time series regressions. See Davidson and MacKinnon (1993) for an introduction and comprehensive exposition of these concepts. Most of these tests can be done with OLSQ and CDF commands on a few simple lagged and differenced variables, so the main function of COINT is to summarize the key regression results concisely and to automate the selection of the optimal number of lags.

COINT (ALL, ALLORD, COINT, CONST, DF, EG, FINITE, JOH, MAXLAG=<number of lags>,MINLAG=<number of lags>, PP, PRINT, RULE=AIC2, SEAS, SEAST, SEASTSQ, SILENT, TERSE, TREND, TSQ, UNIT, WS) <list of variables> [ | <list of special exogenous trend variables> ] ;

or

UNIT (ALL, NOCOINT, CONST, DF, FINITE, MAXLAG=<number of lags>, MINLAG=<number of lags>, PP, PRINT, RULE=AIC2, SEAS, SEAST, SEASTSQ, SILENT, TERSE, TREND, TSQ, UNIT, WS) <list of variables> [ | <list of special exogenous trend variables> ] ;

Usage

List the variables to be tested, and specify the types of tests, maximum number of augmenting lags, and standard constant/trend variables in the options list. The default performs augmented Weighted Symmetric, Dickey-Fuller, and Engle-Granger tests with 0 to 10 lags. If there are any special exogenous trend variables, such as split sample dummies or trends, give their names after a | (see the explanation under General Options below). The observations over which the test regressions are computed are determined by the current sample. If any observations have missing values within the current sample, COINT drops the missing observations and prints a warning message (or an error message, if a discontinuous sample would result).

Output

The default output prints a table of results plus coefficients for each test on each variable. Two summary tables are also printed with just the optimal lag lengths (one for all the unit root tests, and one for the Engle-Granger tests if ALLORD is used).
For each variable, all the specified types of unit root tests are performed. A table is printed for each type of test. Usually, the rows of this table are: the estimated root (alpha), test statistic, P-value, coefficients of trend variables, number of observations, the log likelihood, AIC, and the standard error squared. The columns of this table are the number of augmenting lags. A summary table is also printed which includes just the test statistics and P-values for the optimal lag length.

COINT usually stores most of these results in data storage for later use (except when a 3-dimensional matrix would be required). The summary tables are always stored. If more than one variable is being tested, @TABWS, @TABDF, and @TABPP are not stored. If ALLORD is used, @TABEG is not stored (but @EG, %EG, and @EGLAG will be stored). In the table of output results below,

\[
\#\text{regs} = \text{MAXLAG} - \text{MINLAG} + 2 \text{ (if MINLAG < MAXLAG)} \\
\#\text{regs} = 1 \text{ (if MINLAG=MAXLAG)} \\
\#\text{stats} = 3 + 2(\#\text{trend-vars} + \#\text{regs(if PRINT is on)}) + 4 + 1 \text{ (for PP 2)} + \#\text{vars}\times3 + 3 \text{ (for PP 2)} \\
\#\text{types} = \text{number of types of unit root tests performed (2 for default, 3 for ALL, etc.)} \\
\#\text{eg} = \text{number of different cointegrating regressions for Engle-Granger type tests (#vars for ALLORD, or 1 by default)}.
\]

Here are the results generally available after a COINT command:

<table>
<thead>
<tr>
<th>Name</th>
<th>Type</th>
<th>Length</th>
<th>Variable Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@TABWS</td>
<td>matrix</td>
<td>#stats*#regs</td>
<td>table for augmented WS tau tests on a single variable.</td>
</tr>
<tr>
<td>@TABDF</td>
<td>matrix</td>
<td>#stats*#regs</td>
<td>augmented Dickey-Fuller tau tests.</td>
</tr>
<tr>
<td>@TABPP</td>
<td>matrix</td>
<td>#stats*#regs</td>
<td>Phillips-Perron Z tests.</td>
</tr>
<tr>
<td>@UNIT</td>
<td>matrix</td>
<td>#types*#vars</td>
<td>summary table of unit root test statistics for optimal lags.</td>
</tr>
<tr>
<td>%UNIT</td>
<td>matrix</td>
<td>#types*#vars</td>
<td>P-values for optimal lags.</td>
</tr>
<tr>
<td>@UNITLAG</td>
<td>matrix</td>
<td>#types*#vars</td>
<td>Optimal lag lengths chosen by RULE.</td>
</tr>
<tr>
<td>@TABEG</td>
<td>matrix</td>
<td>#types*#vars</td>
<td>table for augmented Engle-Granger tests.</td>
</tr>
<tr>
<td>@CIVEG</td>
<td>matrix</td>
<td>#types*#vars</td>
<td>cointegrating vector (normalized)</td>
</tr>
<tr>
<td>%EG</td>
<td>vector</td>
<td>#eg</td>
<td>P-values for optimal lags.</td>
</tr>
<tr>
<td>@EGLAG</td>
<td>vector</td>
<td>#eg</td>
<td>Optimal lag lengths chosen by RULE.</td>
</tr>
<tr>
<td>@TABJOH</td>
<td>matrix</td>
<td>#stats*#regs</td>
<td>table for Johansen tests.</td>
</tr>
</tbody>
</table>
Method

Unit root tests are based on the following regression equation:

\[ y_t = y_0 + y_1 t + y_2 t^2 + v_t \]
\[ \nu_i = \alpha v_{i-1} + u_i \]
\[ = y_0 + y_1 t + y_2 t^2 + \alpha [ y_{t-1} - y_1(t-1) - y_2(t-1)^2 ] + u_t \]
\[ = \alpha y_{t-1} + \delta_0 + \delta_1 t + \delta_2 t^2 + u_t \]
\[ u_t = \varphi_1 u_{t-1} + \varphi_2 u_{t-2} + \ldots + \varphi_L u_{t-L} + e_t \quad [AR(L)] \]

Let \( L=1 \) for illustration:

\[ y_t = \alpha y_{t-1} + \delta_0 + \delta_1 t + \delta_2 t^2 + \varphi_1 [ y_{t-1} - \alpha y_{t-2} - \delta_0 - \delta_1(t-1) - \delta_2(t-1)^2 ] + e_t \]
\[ = [ \alpha + (1-\alpha) \varphi_1 ] y_{t-1} + \beta_0 + \beta_1 t + \beta_2 t^2 + \alpha \varphi_1 dy_{t-1} + e_t \]

All unit root tests are computed from (possibly weighted) OLS regressions on a few lagged or differenced variables. The coefficient of \( y(t-1) \) is printed in the tables as alpha. Accurate asymptotic P-values for Dickey-Fuller, Phillips-Perron, and Engle-Granger (for up to 6 cointegrating variables) are computed using the coefficients in the MacKinnon reference. Note that these asymptotic distributions are used as approximations to the true finite-sample distributions.

The WS test is a weighted double-length regression. First the variable being tested is regressed on the constant/trend variables (using the full current sample), and the residual from this is used as the dependent variable \( Y \) in the double-length regression. The data setup for the first half of this regression is the same as an augmented Engle-Granger test -- regress \( Y \) on lagged \( Y \) and lags of \( DY \). The weights are \((t-1)/T\), where \( T \) is @NOB in the original sample. In the second half, \( Y \) is regressed on \( Y(+1) \) and leads of \( Y-Y(+1) \), using weights \((t-(t-1))/T\). See Pantula et al (1994) for more details. P-values for the WS test are computed very roughly by interpolating between the asymptotic 5% and 10% level critical values given for the constant and no trend case in the reference. These P-values are fine for testing at the 5% and 10% levels, but they are not accurate for testing at other levels. The P-value for the case with a constant and a trend is only good for testing at the 5% level.
The regressions for the Dickey-Fuller tests are quite simple. See the example below which reproduces the Dickey-Fuller tests in the Examples section below.

```
SMPL 10,70; DY = LRGNP-LRGNP(-1);
? Sample for comparing AIC is the same for all lags.
? MAXLAG+1 observations are dropped.
SMPL 20,70;
TREND T;
DO LAG=1,10;
  SET MLAG = -LAG;
  OLSQ LRGNP LRGNP(-1) C T DY(-1)-DY(MLAG);
  SET alpha = @COEF(1);
  SET tauDF = (alpha - 1)/@SES(1);
  CDF(DICKEYF) tauDF;
ENDDO;
```

If you are computing this test by hand, it is easier to use:

```
OLSQ DY LRGNP(-1) C T DY(-1)-DY(MLAG) ;
CDF (DICKEYF) @T(1) ;
```

The Phillips-Perron test is done with the same Dickey-Fuller regression variables, using no augmenting lags. This test is given in Davidson and MacKinnon, equations (20.17) and (20.18) (see also the warnings there about the possibly poor finite-sample behavior of this test). These tests can be computed for 1 to 10 "lags" by using the following TSP commands (following the Dickey-Fuller example above):

```
OLSQ (silent) LRGNP C T ;
SET ssr =@SSR ;
? to ensure the same sample for each test:
SMPL 10,70;
TREND T;
OLSQ LRGNP LRGNP(-1) C T; Y = @RES;
SET alpha = @COEF(1); SET s2 = @S2; SET n = @NOB;
FRML EQPP Y = Y0; PARAM Y0;
DO LAG=1,10;
  GMM(INST=C,NMA=LAG,SILENT) EQPP;
  SET w2 = @COVOC;
  SET z = n*(alpha-1) - [n**2*(w2 - s2)]/[2*ssr];
  PRINT LAG,z,w2;
ENDDO;
```

The regressions for the Engle-Granger tests are just an extension of the Dickey-Fuller test, after an initial cointegrating regression:
TREND T;
OLSQ LGNP LEMPLOY C T; \( c \)ointegrating regression
\( E = @RES; \)
SMPL 10,70; DE = E-E(-1);
\( \) Estimation sample same for all lags -- MAXLAG+1 obs are dropped.
SMPL 20,70;
DO LAG=1,10;
SET MLAG = -LAG;
OLSO E E(-1) DE(-1)-DE(MLAG);
SET alpha = @COEF(1);
SET tauDF = (alpha - 1)/@SES(1); CDF(DICKEYF,NVAR=2) tauDF;
ENDDO;

The following equation defines the \( L+1 \) order VAR (Vector Auto Regression) that is used in the Johansen trace test:

\[
Y_t = Y_{t-1} \Pi_1 + Y_{t-2} \Pi_2 + \ldots + Y_{t-L+1} \Pi_{L+1} + CT_t \beta + U_t
\]

where \( Y(t) \) is \( 1 \) by \( G \) and \( CT(t) \) are (seasonal) constants and trends.

\( \Pi = I - \Pi_1 - \ldots - \Pi_{L+1} \) is a \( G \) by \( G \) impact matrix of rank \( r \)

\( \eta \alpha' \) where \( \eta \) and \( \alpha \) are \( G \) by \( r \) matrices

The Log Likelihood and AIC printed in the table are from the unrestricted version of this VAR. The restricted version is estimated with a \( 2G \)-equation VAR:

\[
dY_t = dY_{t-1} A_1 + dY_{t-2} A_2 + \ldots + dY_{t-L} A_L + CT_t A_0 + E_t
\]

\[
Y_{t-L-1} = dY_{t-1} B_1 + dY_{t-2} B_2 + \ldots + dY_{t-L} B_L + CT_t B_0 + F_t
\]

\[
S_{00} = \frac{\hat{E}^\prime \hat{E}}{T-L-1}, \quad S_{0k} = \frac{\hat{E}^\prime \hat{F}}{T-L-1}, \quad S_{kk} = \frac{\hat{F}^\prime \hat{F}}{T-L-1}
\]

\[
Q = S_{kk}^{-1/2} = \text{CHOL}(S_{kk})^t, \text{ that is, } QQ' = S_{kk}^{-1}
\]

\[
S = Q' S_{00} S_{0k} Q
\]

\( \lambda = \text{eigenvalues of } S, \text{ sorted high to low} \)

\( \eta = Q(\text{eigenvectors of } S) \)

\[
\text{trace}_i = -(T-L-1-(L+1)G) \sum_{j=1}^i \log(1 - \lambda_j)
\]
where \( T \) = number of observations in the current sample, \( L = \text{MAXLAG} \) = order of \( \text{VAR} \) beyond 1. The trace tests are labelled H0: \( r=0 \), H0: \( r<=1 \), etc. in the table of results. Note that the trace test includes a finite-sample correction (mentioned in Gregory (1994); originally given in Bartlett(1941)). These trace tests often have size distortions (the null of no cointegration or fewer cointegrating vectors is rejected when it is actually true). P-values are interpolated from the Osterwald-Lenum (1992) tables 0, 1.1*, and 2 (with no constant, constant, or constant & trend). These P-values are adequate for testing at the sizes given in the Osterwald-Lenum tables (.50, .20, .10, .05, .025, and .01). See Cushman et al (1995) for a detailed example of using Johansen tests in an applied setting. They illustrate the importance of the finite sample degrees of freedom correction, the size distortions of the P-values, lag length choice methods, and hypothesis testing.

Options

Unit Root Test Options:

ALL/NOALL perform all available types of unit root tests (WS, DF, and PP).

DF/NODF perform (augmented) Dickey-Fuller (tau) tests.

PP/NOPP perform the Phillips-Perron variation of the Dickey-Fuller (Z) test. For the PP test, the number of lags used is the order of the autocorrelation-robust T2 "long run variance" estimate (see the MAXLAG option).

WS/NOWS perform (augmented) Weighted Symmetric (tau) tests. This test seems to dominate the Dickey-Fuller test (and others) in terms of power, so it is performed by default. See Pantula et al (1994) or the Method section for details.

UNIT/NOUNIT use NOUNIT to skip all unit root tests (if you are only interested in cointegration tests, and you are sure which individual variables have unit roots).

Cointegration Test Options: (these apply only if you have more than one variable)

ALLORD/NOALLORD repeat the Engle-Granger tests, using each variable in turn on the left hand side of the cointegrating regression.
COINT

**EG/NOEG** perform (augmented) Engle-Granger tests (Dickey-Fuller test on residuals from the cointegrating regression). The Engle-Granger test is only valid if all the cointegrating variables are I(1); hence the default option to perform unit root tests on the individual series to confirm this before running the Engle-Granger test. Note that if you accept I(1) (i.e. reject I(0)), you will also want to difference the series and repeat the unit root test, to make sure you reject I(2) in favor of I(1). Note that you need to reduce the order of trends when testing such a differenced series -- for example, if the original series had a constant and trend in the equation, the differenced one will only have a constant.

**JOH/NOJOH** perform Johansen (trace) cointegration tests.

**COINT/NOCOINT** use NOCOINT to skip all cointegration tests (if you are only interested in unit root tests). You may prefer to use the UNIT or UNIT (NOCOINT) command for this. (UNIT and COINT are synonyms for the same command, except that UNIT has a default of NOCOINT; UNIT may also seem more appropriate if you are just testing one variable).

*General Options: (these apply to both unit root and cointegration tests)*

**CONST/NOCONST** include a constant term in the tests. NOCONST implies NOTREND.

**FINITE/NOFINITE** computes finite sample (vs. asymptotic) P-values when possible (augmented Dickey-Fuller and Engle-Granger tests). See the discussion and references under Method in the CDF entry of this manual. These are distinguished by different labels: P-valFin or P-valAsy ; normally the finite sample P-values will be slightly larger than the asymptotic ones.

**MINLAG** = smallest number of augmenting lags (default 0). This is denoted as $L$ in the equations under Method. Note that $p=L+1$ is the total AR order of the process generating $y$. So $L$ is the number of lags in excess of the first one. For the Phillips-Perron test, $L$ is the number of lags in the "autocorrelation-robust" covariance matrix.

**MAXLAG** = maximum number of augmenting lags. The default is min(10,2*@NOB(1/3)), which is 10 for 100 observations or below (the factor 2 was chosen arbitrarily to ensure this). If the number of observations in the current sample (@NOB) is extremely small, MAXLAG and MINLAG will be reduced automatically.
RULE= AIC2 or specifies the rule used to choose an optimal lag length (number of augmenting lags), assuming MINLAG < MAXLAG. The default is AIC2, which is described in Pantula et al (1994). If \( j \) is the number of lags which minimizes AIC (Akaike Information Criterion), then \( L = \text{MIN} (j + 2, \text{MAXLAG}) \) is used. Note that if \( j = \text{MAXLAG} \), you will probably want to increase MAXLAG. AIC2 apparently avoids size distortions for the WS and DF tests. AIC2 is also used here for EG tests. No direct rule is used for PP tests yet. Instead, the optimal lag from the DF test is also used for PP (if the DF test is performed at the same time). A plain AIC rule is used for JOH, i.e. \( L = j \) (this is not a very good rule for JOH; you may prefer to run the unconstrained VAR and test its residuals for serial correlation). These rules are a topic of current research, so as more useful rules are found, they will be added as options. For example, other possible rules are: (1) testing for remaining serial correlation in the residuals, (2) testing the significance of F-statistics for the last lag of (differenced) lagged variable(s), (3) SBIC (+2?), (4) automatic bandwidth selection for PP (not very encouraging in the current literature).

The current RULE=AIC uses a fixed number of observations for comparing regressions with different numbers of lags. Each regression is a column in the output table. If MINLAG<MAXLAG, then the RULE is used to select an "optimal" number of lags (\( j \)). A final column in the table is created for this, labelled "Opt:j". If \( j \) is less than MAXLAG, then the regression for this column is computed with the maximum available observations, so the test results may vary slightly from the original column for \( j \).

SEAS/NOSEAS include seasonal dummy variables, such as Q1-Q3. This option implies the CONST option. The SEAS option is available for FREQ Q, 2, or higher. The seasonal coefficients are only printed if PRINT is on.

SEAST/NOSEAST include seasonal trend variables (like Q1*TREND, Q2*TREND, Q3*TREND). This option implies the TREND option.

SEASTSQ/NOSEASTS include seasonal squared trend variables. SEASTSQ implies TSQ.

These are fairly simplistic trend terms, which may not be enough to adequately model a time series that has a change in its intercept and/or trend at some point in the sample. See Perron (1989) for more details. The "special exogenous trend variables" arguments described above may provide a crude examination of more detailed trends. If these variables are supplied, all series are regressed on these trend variables, and the residuals from this regression are used in all tests (instead of the original values of the series). No corrections to the P-values of the tests are made, however (other than in the degrees of freedom for calculating the t-statistics and s2).

TREND/NOTREND include a time trend in the tests.
**COINT**

TSQ/NOTSQ include a squared time trend in the tests.

*Output options*

PRINT/NOPRINT prints the options, and adds the coefficients and t-statistics of the augmenting lagged difference variables to the main tables.

TERSE/NOTERSE suppresses the main tables (only the summary tables are printed). Note that JOH and EG(NOALLORD) have no summary tables, so TERSE suppresses all their output.

SILENT/NOSILENT suppresses all output. This is useful for running tests for which you only want selected output (which can be obtained from the @ variables, that are stored - see the table below).

*Examples*

**FREQ A; SMPL 1909,1970; COINT LRGNP LEMPLOY;**

performs 11 augmented WS (tau) and Dickey-Fuller (tau) unit root tests with 0 to 10 lags. All tests are first done for LRGNP, then repeated for LEMPLOY. Eleven augmented Engle-Granger (tau) tests are constructed with 0 to 10 lags (with LRGNP as the dependent variable in the cointegrating regression). Optimal lag lengths for all tests are determined using the AIC2 rule. The test is recomputed for the optimal lag, using the maximum available observations, and this is stored in the final column of the table. All tests use a constant and trend variable.

**UNIT (ALL) LRGNP;**

performs the same unit root tests for LRGNP as the above example. Also performs the Phillips-Perron (z) tests computed separately for 0 to 10 lags in the autocorrelation-robust estimate.

**COINT (NOUNIT,ALLORD,MAXLAG=8) LRGNP LEMPLOY LCPI;**

performs 27 augmented Engle-Granger (tau) tests. That is, 9 tests with 0 to 8 lags, with LRGNP as the dependent variable in the cointegrating regression. Then repeat the tests, using LEMPLOY and later LCPI as the dependent variable in the cointegrating regression.

**SMPL 58:2 84:3; COINT(JOH,MAXLAG=2,SEAS,NOTREND,NOUNIT,NOEG) Y1-Y4;**

92
reproduces the Johansen-Juselius (1990) results for Finnish data (the chosen number of lags is 1, which matches the results from the paper). The test statistics are smaller than those in the paper, due to the finite-sample correction.

References


COLLECT (Interactive)

COLLECT allows a group of TSP statements to be entered before execution of the sequence.

**COLLECT** ;

**Usage**

This is particularly useful for introducing flow of control structures not directly allowed in interactive mode such as **PROC**s, **IF-THEN-ELSE** sequences, or **DO** or **DOT** loops. You will continue to be prompted for new lines in collect mode until you terminate the COLLECT mode.

Termination of the mode may be accomplished in two ways:

1. The **EXEC** command requests automatic execution of the entire range of commands just collected. Unlike its usage in interactive mode, the EXEC command takes no arguments when used to terminate collect mode.

2. The **EXIT** command may be used to return to interactive mode without executing the collected lines. The lines are stored and EXEC may be used on them interactively at any time.

While entering commands in collect mode, you may find you want to fix a typo, or modify something you've entered before requesting execution. For this reason, there are two commands that are always executed immediately, even while in collect mode - these are **EDIT** and **DELETE**. Of course, if you really mess things up, or simply change your mind, you can always abandon the effort with **EXIT**.

**HINT:** If there are **PROC**s or sequences of commands you find you use frequently (whether in collect mode or not) you may want to store them in external files to save having to type them more than once. Any group of TSP commands may be read from disk with the **INPUT** command. **INPUT** is functionally equivalent to collect mode, the only difference being that the commands are read from a file instead of your terminal. See the section on **INPUT** for more details on how to use this feature.

**WARNING:** You will confuse TSP if you begin a control structure in collect mode, and fail to end it properly before returning to interactive mode, e.g.:

```
1? **COLLECT**
```

Enter commands to be collected:
COLLECT

2> PROC X
3> (other statements)
4> ENDPXC <-- proper end to structure
5> EXIT

Collect mode is always entered from, and control always returned to the interactive mode.
COMPRESS

**Options**

COMPRESS removes unused TSP variables and frees up wasted working space.

**COMPRESS (PRINT);**

**Usage**

If you are trying to fit a particularly large TSP program onto a computer with limited working space, this command may be useful. It deletes internal variables such as old program lines and GENR formulas, variables whose names begin with @ (such as @RES), and variables which have been marked with the `DELETE` command. It also frees up the space associated with these variables and the space used by old copies of variables which have grown in size.

This command is not usually necessary, since TSP does an automatic compress operation when it runs out of working space or when too many TSP variables have been defined. However, in some cases this automatic compress will not work. For instance, no compression takes place inside a `PROC` which has arguments, and internal program lines are deleted only up to the first `DO` loop, first `PROC` or current `DOT` loop. Automatic compression also may not clear enough space when creating a large matrix. In these cases, use the COMPRESS command first before running a command which may need a large part of available memory.

An alternative way to compress out unused variables is to use the `SAVE` command, followed by restarting TSP and using the `RESTORE` command. This would remove all the PROCs and leave only variables like time series, scalars, lists, FRMLs, and matrices.

**Output**

The number of deleted variables (if any) is printed. The amount of recovered working space (if any) is also printed.

**Options**

PRINT/NOPRINT specifies whether a summary message is printed describing the space freed by compression. This message is always printed for automatic compression.
CONSTR

**Examples**

CONST defines scalar variables (constants) and assigns arithmetic values to them. To define scalars that will be estimated in one of the nonlinear procedures such as **LSQ**, use **PARAM** instead.

**CONST varname, value, varname, value,..... ;**

**Usage**

CONST may be followed by as many argument pairs as desired (limited only by TSPs argument limit). Each pair is the name of the scalar variable followed by the value it is to be given. The variable names may be new or previously defined variables. The value may be omitted, in which case the variable is either given the value zero if it is new or left unchanged if it has already been defined.

The use of the CONST procedure is primarily to suppress the estimation of some of the parameters in a nonlinear estimation: instead of using a **PARAM** statement to give the parameter a starting value, use a CONST statement to fix the parameter throughout the estimation.

**Output**

CONST produces no printed output; it stores the variables named in data storage.

**Examples**

**CONST DELTA .15 ;**
**CONST A1 A2 A3 A4 ;**
**LIST ALIST A1-44; CONST ALIST ;**
**PARAM ALPHA 1.0 BETA .5; CONST ALPHA BETA GAMMA .9;**

The second and third of these examples have the same effect. The fourth assigns a value only to the third variable GAMMA; the other two variables have the same value as they did previously, but their type is changed from PARAM to CONST.
CONVERT

Options  Examples

CONVERT changes series from one frequency to another. The options specify the method used for conversion: averaging the data, using the first, middle or last observation, or summing the data. Interpolation is optionally available for converting to higher frequencies.

CONVERT (AVERAGE or FIRST or MID or LAST or SUM, INTERPOL, MAP= <series>, SMPL) <list of series names> ;

or

<newseries> = <oldseries> ;

Usage

Use CONVERT after specifying the frequency you want to convert to with a FREQ statement. The frequency you convert from will be that of the series to be converted. The SMPL information is ignored so that the entire series is converted; this avoids the confusion of possibly mixing frequencies in the same series.

The first form of the command simply converts the series and stores it back in data storage under the same name; in this case more than one series can be converted at a time. The second form takes the old series on the right hand side of the equal sign, converts it to the new FREQ and stores it under the new series name; only one series may be converted in this way on each command.

Depending on the type of series you are converting, you can specify various methods of aggregating or “disaggregating” the series; if you do not say anything and you are converting to a lower frequency, CONVERT will average all the observations within an interval to produce a value for that interval. The default for converting to a higher frequency is to duplicate the value for all observations in the new interval; unless the INTERPOL or SUM option is used. With FREQ(PANEL), the individual sum, average, etc. is replicated to all observations for the individual.

Output

CONVERT produces no printed output. It stores one converted series in data storage.

Options

AVERAGE forms the new series by averaging all the observations within a period. This is the default.
FIRST forms the new series by choosing the first observation in the period.

MID forms the new series by choosing the middle observation in the period. If the number of observations per period is even, CONVERT uses the one before the halfway point.

LAST forms the new series by choosing the last observation in the period.

SUM forms the new series by summing all the observations in the period. If converting from a lower frequency to a higher, the new values are divided by the conversion ratio (e.g., by four, if converting from annual to quarterly).

Only one of the above options should be included.

INTERPOL/NOINTERP specifies linear interpolation when converting to a higher frequency (the default is to duplicate observations rather than interpolate). INTERPOL is used in conjunction with one of the other options to determine the placement of the peak value.

MAP= series computes SUM (default) or AVERAGE from an old series and stores it in a new series, using a MAP of pointers. This is helpful for aggregating grouped data, such as industries, states, or individuals with panel data. The rows of the map correspond to the rows in the old series. The values in the map correspond to the rows of the new series. Zero values mean the observation is not mapped to the new series. The SMPL option is the default when MAP is used, and it puts the map and old series under the control of the current SMPL, while the new output series will be FREQ N, starting at observation 1. If NOSMPL is used, the traditional CONVERT method is used, where the old and map series are used at their maximum defined lengths, and the current FREQ/SMPL are only used to determine the FREQ and starting point of the new series. With NOSMPL, the map and old series must be defined over exactly the same set of observations. The map cannot contain any missing values or contain only zeroes. The old series can contain missing values; they will result in missing values in the new series if the observations are mapped. If no observations of the old series are mapped to a given element of the new series, the element is given the value zero (for both sum and average). The length of the new series is equal to the maximum value in the map series.

SMPL/NOSMPL applies only when the MAP option is used. Otherwise the default is NOSMPL (use all the data in the series).

Examples

FREQ A;
CONVERT (AVERAGE) UNEMP;
CONVERT (SUM) SALES;
CONVERT (LAST) PCLOSE = PRICE;
Assume that UNEMP and SALES are quarterly variables and PRICE is a monthly variable. The statements above convert the unemployment rate by averaging the quarterly rates over the year, but convert sales from quarterly to annual by adding them, since they are a flow variable. The end of year price (PCLOSE) is obtained by using the December observation of the monthly price variable.

```
FREQ A ; SMPL 70, 72 ;
READ X ; 10 20 40 ;
FREQ Q ;
CONVERT DX = X ; CONVERT (SUM) SX=X ;
CONVERT (INTER,LAST) IX = X ;
```

results in

<table>
<thead>
<tr>
<th></th>
<th>DX</th>
<th>SX</th>
<th>IX</th>
</tr>
</thead>
<tbody>
<tr>
<td>70:1</td>
<td>10</td>
<td>2.5</td>
<td>2.5</td>
</tr>
<tr>
<td>70:2</td>
<td>10</td>
<td>2.5</td>
<td>5.0</td>
</tr>
<tr>
<td>70:3</td>
<td>10</td>
<td>2.5</td>
<td>7.5</td>
</tr>
<tr>
<td>70:4</td>
<td>10</td>
<td>2.5</td>
<td>10</td>
</tr>
<tr>
<td>.</td>
<td>.</td>
<td>.</td>
<td>.</td>
</tr>
<tr>
<td>.</td>
<td>.</td>
<td>.</td>
<td>.</td>
</tr>
<tr>
<td>72:1</td>
<td>40</td>
<td>10</td>
<td>25</td>
</tr>
<tr>
<td>72:2</td>
<td>40</td>
<td>10</td>
<td>30</td>
</tr>
<tr>
<td>72:3</td>
<td>40</td>
<td>10</td>
<td>35</td>
</tr>
<tr>
<td>72:4</td>
<td>40</td>
<td>10</td>
<td>40</td>
</tr>
</tbody>
</table>

Example of the MAP= option:

```
SMPL 1 5 ;  TREND T ;
READ MAPS; 0 1 2 2 3 ;
CONVERT (MAP=MAPS,AVE) AT = T ;
CONVERT (MAP=MAPS,SUM) ST = T ;
```

which yields the following:

<table>
<thead>
<tr>
<th></th>
<th>AT</th>
<th>ST</th>
</tr>
</thead>
<tbody>
<tr>
<td>2</td>
<td>2</td>
<td></td>
</tr>
<tr>
<td>3.5</td>
<td>7</td>
<td></td>
</tr>
<tr>
<td>5</td>
<td>5</td>
<td></td>
</tr>
</tbody>
</table>

```
FREQ(PANEL); CONVERT(SUM) NTI=C ;  ? NTI = obs per individual
```
COPY

Example

Makes a copy of an old TSP variable (series, matrix, constant, etc.) with a new name.

`COPY <old TSP variable> <new TSP variable>;`

Output

The new variable is stored in data storage. The old variable is left unchanged. If a variable with the new name already exists, it is deleted.

Example

Save the coefficients from a regression in the vector B1. Note that this can usually be done more efficiently with `RENAME`, unless the original @COEF needs to be saved for a procedure like `FORCST`.

`OLSQ Y C X;`
`COPY @COEF B1;`
CORR/COVA

See MSD

CORR (ALL, COVA, MOMENT, MSD, PAIRWISE, PRINT,
    WEIGHT=<seriesname>) <list of variables> ;
COVA (ALL, CORR, MOMENT, MSD, PAIRWISE, PRINT,
    WEIGHT=<seriesname>) <list of variables> ;
DATE

Examples

DATE prints the current time and date. This is useful when printing results from an interactive session.

If a variable name is supplied, nothing is printed, and the number of seconds since midnight are stored in the variable. This is useful for timing a group of TSP statements, such as a PROC.

DATE [<scalar variable name>];

Examples

DATE; ? print the current date/time, to "time stamp" the output file.

? print the elapsed time required by the procedure MYPROC
DATE SEC0;
MYPROC Y Z;
DATE SEC1;
SET NSEC = SEC1-SEC0;
WRITE (FORMAT=’(’’MYPROC took ’,G12.1,’ seconds.’’)’) NSEC;
DBCOMP (Databank)

DBCOMP compresses a TSP databank.

*DBCOMP* <filename> ;

**Usage**

Follow the word DBCOMP with the name of the TSP databank to be compressed. This can be useful if your disk storage is space-constrained and you have a great deal of data.
DBCOPY (Databank)

Options  Example

DBCOPY makes it possible to move a TSP databank from one type of computer to another, since the actual databank files are not compatible between different computers.

**DBCOPY (DOC) <list of filenames> ;**

Usage

Follow the word DBCOPY with the filenames of the TSP databanks to be moved. A file containing TSP commands and data is created for each databank. When this file is moved to another computer and run with TSP, the databank is created with all its original variables and values. The filename with the TSP commands will be the same as the databank name, except it will have filetype .TSP instead of .TLB. The .TSP file has record length of at most 80, so it can be easily moved to another computer. There are no restrictions on the sizes of the databanks, SMPLs and FREQs of the series, or types of the TSP variables (it handles CONST, PARAM, SERIES, MATRIX, FRML, and IDENT). The SMPLs and FREQs are determined by the series in the databank, not by the current SMPL or FREQ.

Options

**DOC/NODOC** controls the listing of documentation (created with the DOC command). Specify NODOC if you will be using TSP version 4.1 or earlier.

Example

Suppose that you have a databank called FOO.TLB which contains two time series X and Y, and a parameter B. The command

**DBCOPY FOO;**

would create the file FOO.TSP which would contain:

```
? Re-create TSP Databank
? FOO.TLB
END; OUT FOO;
PARAM B 3.14;
FREQ Q; SMPL 60:1,85:4;
LOAD X;
1 2 3 4 ... ;
LOAD Y;
11 22 33 44 ... ;
```
DBDEL (Databank)

Options  Example

DBDEL deletes one or more variables from a TSP databank.

**DBDEL (COMPRESS)** <filename> <list of variables> ;

Usage
Supply the filename and one or more variable names to delete. This is a way of getting rid of variables which were put in the databank by mistake, or which are no longer needed. It can also be used to crudely rename variables in a databank, if the variables are first copied to the new names and then the old names are deleted. An alternative for small databanks would be to use the **DBCOPY** command, and then use a text editor on the .TSP file to delete, rename, or generally edit variables and DOCUMENTATION.

Options

**COMPRESS**/**NOCOMPRESS** compresses the databank after the variables are deleted. This is the same operation as the DBCOMP command; the space formerly used by the variables is recovered for use by other files.

Example

Suppose that you have a databank called FOO.TLB which contains two time series X and Y, and a parameter B. The command

**DBDEL FOO X**;

would delete the series X from the databank.
DBLIST

DBLIST (Databank)

**Output**  **Options**  **Example**

DBLIST shows the contents of TSP databanks, with information on variable names, types, frequencies, and sample coverage.

**DBLIST (DATE,DOC,SILENT) <list of filenames> ;**

**Usage**

Follow the word DBLIST with the filenames of the TSP databanks to be listed. A list of all the variables contained in each databank is printed, along with brief information about each variable (in the format of the SHOW command). If the databank contains wasted space which could be removed with the DBCOMP command, the amount of wasted space is indicated. The FREQ and SMPL information on series may be useful for setting up a DBPRINT command.

**Output**

A list of the contents of the databanks will be printed. The list of names of the databank variables is stored in @RNMS for further use.

**Options**

DATE/NODATE specifies whether the date is to be printed.

DOC/NODOC specifies whether series documentation is to be printed.

SILENT/NOSILENT suppresses all printed output. @RNMS is stored.

**Example**

Suppose that you have a databank called FOO.TLB which contains two time series X and Y, and a parameter B. The command

**DBLIST FOO;**

would print the following information:

```
CONTENTS OF DATABANK FOO.TLB

<table>
<thead>
<tr>
<th>Class</th>
<th>Name</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>SCALAR</td>
<td>B</td>
<td>parameter 3.14000</td>
</tr>
<tr>
<td>SERIES</td>
<td>X</td>
<td>104 obs. from 1960:1-1985:4, quarterly</td>
</tr>
<tr>
<td></td>
<td>Y</td>
<td>104 obs. from 1960:1-1985:4, quarterly</td>
</tr>
</tbody>
</table>
```
DBPRINT (Databank)

Example

DBPRINT prints all the series in a TSP databank, under the control of the current FREQ and SMPL.

**DBPRINT <filename> ;**

Usage

Follow the word DBPRINT with the filename of the TSP databank whose series are to be printed. Make sure the FREQ (if any) and SMPL have been set. The FREQ and SMPL information on series from a DBLIST command may be useful for setting FREQ and the SMPL range. Any series not stored with the FREQ currently in effect will not be printed. TSP variables other than series, such as matrices and FRMLs, will not be printed. Parameters and constants can be printed with the DBLIST command, while matrices and equations can be printed using IN and PRINT together. The series printed are only temporarily brought into memory -- they are only stored when they have been accessed with an IN command. This protects any currently stored series from being overwritten by a series in the databank with the same name. Only one databank can be printed in a DBPRINT command.

Example

Suppose that you have a databank called FOO.TLB which contains two time series X and Y, and a parameter B. The commands

```
FREQ Q;
SMPL 60:1,60:4;
DBPRINT FOO;
```

would yield:

VALUES FOR ALL SERIES IN DATABANK FOO.TLB

<table>
<thead>
<tr>
<th></th>
<th>X</th>
<th>Y</th>
</tr>
</thead>
<tbody>
<tr>
<td>1960:1</td>
<td>1.00000</td>
<td>11.0000</td>
</tr>
<tr>
<td>1960:2</td>
<td>2.00000</td>
<td>22.0000</td>
</tr>
<tr>
<td>1960:3</td>
<td>3.00000</td>
<td>33.0000</td>
</tr>
<tr>
<td>1960:4</td>
<td>4.00000</td>
<td>44.0000</td>
</tr>
</tbody>
</table>
DEBUG

Examples

DEBUG turns the DEBUG switch on. When this switch is on, TSP produces a great deal more printed output than it usually does. This output is normally not of interest to users, but may be helpful to a TSP programmer or consultant.

DEBUG ;

Usage

Include the DEBUG statement in your TSP program directly before the command(s) for which you want additional output. The DEBUG switch will remain on until a NODBUG statement is encountered. For DEBUG output during the compile phase of the program, see the ASMBUG statement.

Output

DEBUG should be used with care, since it normally produces a great deal of printed output. For example, every fetch and store to data storage (VPUT/VGET) is printed, which facilitates following the progress of the program, but can be voluminous. In the nonlinear procedures, all the input data and the results of the differentiation of the equations will be printed. For any estimation procedure, the matrices involved in the computations will be printed at every iteration.

Every input command line will be printed, before and after it has been interpreted for dates, dots, and lists. Several of the non-estimation procedures also produce special debug output when this switch is on.

Example

DEBUG ;
LSQ (PRINT) EQNAME ;
NODBUG ;

This example causes debug output to be produced during the execution of a nonlinear least squares estimation.
DELETE

DELETE removes TSP variables from the symbol table.

`DELETE <list of variables> ;`

Usage

Useful for long or complex TSP programs, DELETE deletes variables from the symbol table. Data for the variables is not actually deleted from memory until an automatic compression occurs when space is needed to store a new variable (see the COMPRESS command).
DELETE (Interactive)

DELETE (Interactive)

DELETE followed by numbers instead of names, removes command lines.

DELETE <firstline>, [<lastline>] ;

Usage

When used in the interactive version of TSP, DELETE enables (re)execution of a range of lines with unnecessary steps eliminated. The lines are permanently lost. It is also useful in COLLECT mode for modifications to the range of lines just entered (before their execution), since DELETE is executed immediately regardless of mode. If both arguments are present, all lines from firstline through lastline will be deleted. If the second argument is omitted, only firstline will be deleted. Arguments must be valid line numbers (i.e., integers). Deleted line(s) will be absent from the backup file; DELETE will remain, however, to provide a more accurate record of actions taken during the session.
DIFFER differentiates a TSP equation analytically with respect to the list of arguments and stores each derivative in parsed form as another TSP equation (FRML). These equations may be evaluated using GENR, used in estimation, or printed just like any other TSP equation.

\[ \text{DIFFER (DEPVARPR=}<\text{new dependent variable name}>\text{, PRINT, PREFIX=}<\text{new equation name}>\text{) } \text{<equation name> } \text{<list of arguments> ;} \]

Usage

DIFFER requires the name of the equation to be differentiated, followed by one or more arguments for differentiation. If any of the arguments are not in the equation, a zero derivative will be stored (no error will be trapped). If the equation is unnormalized (no left hand side variable), the derivative equation will also be unnormalized.

Output

Ordinarily, DIFFER produces no printed output. It stores the derivatives as equations in data storage. If the PRINT option is specified, DIFFER prints each equation in symbolic form with a title specifying what the derivative is.

Options

\text{DEPVARPR=} \text{prefix for new dependent variable}. The default is Dlhsvar. If the original equation has no dependent variable (is unnormalized), then the default is to create unnormalized derivative equations.

\text{PRINT/NOPRINT} tells whether the resulting derivative equations are to be printed.

\text{PREFIX=} \text{the name to be given to the derivative equations if the names Deqname1, etc. are not wanted. The equation names will consist of the prefix name followed by the argument number for that derivative.}

Examples

1. Using the default options:

\[ \text{FRML EQ } Y = A + B*X + G*X**2; \]
\[ \text{DIFFER EQ A B X Q;} \]
DIFFER

creates the following FRMLs:

FRML DEQ1 DY1 = 1;  \  \ ? dY/dA
FRML DEQ2 DY2 = X;  \  \ ? dY/dB
FRML DEQ3 DY3 = B + 2*G*X;  \  \ ? dY/dX
FRML DEQ4 DY4 = 0;  \  \ ? dY/dQ

2. Using some prefix options for naming the results:

FRML EQ Y = A + B*X + G*X**2;
DIFFER (PREFIX=GYX,DEPVAR=MPX) EQ X;

creates the following FRML:

FRML GYX1 MPX1 = B + 2*G*X;  \  \ ? dY/dX

3. Unnormalized equation (residual from AR(1) model):

FRML E Y - (A + B*X) - RHO*(Y(-1) - (A+B*X(-1))); DIFFER E B;

creates the following (unnormalized) FRML:

FRML DE1 -X + RHO*X(-1);

4. Differentiate a Constant Elasticity of Substitution production function with respect to the two inputs K and L, compute the two marginal product series using the derivative equations, and print them.

FRML CES
Y=A*EXP(GAM*T)*(ALPHA*L**RHO+BETA*K**RHO)*(1/RHO);
DIFFER CES L K;
GENR DCES1 LMP ; GENR DCES2 KMP ;
PRINT LMP KMP ;

5. Differentiate the log likelihood for a PROBIT model with respect to its parameters and store the generated equations.

FRML PROBIT LOGL = LOG(D*CNORM(A+B*X) + (1-D)*(1-CNORM(A+B*X)));
DIFFER (PRINT,PREFIX=LOGL) PROBIT A B ;

The results are:

FRML LOG1 DLOG1 = [D*(NORM(A+B*X)+(1-D)*(-NORM(A+B*X))) / [D*CNORM(A+B*X)+(1-D)*(1-CNORM(A+B*X))];
FRML LOG2 DLOG2 = [D*(NORM(A+B*X)*X+(1-D)*(-NORM(A+B*X)*X)) / [D*CNORM(A+B*X)+(1-D)*(1-CNORM(A+B*X))];
This example illustrates the convenience of DIFFER when you use it to generate analytic derivatives of an objective function for use in another program or language, such as Fortran.
DIR

DIR (Interactive)

DIR examines a disk directory without interrupting the interactive TSP session.

**DIR [ * or <filename> ] ;**

Usage

DIR may be used in three different forms. The first,

**DIR ***

makes use of a wildcard specification, and produces a list of all files in your current directory with the extension .TSP. It is expected that the most frequent use of this command will be to locate files to INPUT.

The second form,

**DIR filename**

lists all files in the current directory that fit the description 'filename.*' . You may find it convenient to locate related input, output, and databank files in this way. 'Filename', however, must meet the requirements of a TSP variable name since it will be processed in the same manner. This means that no filename extension is possible (or needed) here.

The third form is the most flexible, and is simply the command with no arguments, prompting you for the file specification:

**DIR files: (computer response)**

in response to which you may type anything you would ordinarily include with a DOS/Windows directory command. In this way you may specify directories other than the current, lists of files, other wildcard combinations, or even command qualifiers such as /date, /size, etc...
DIVIND computes Divisia price and quantity indices from a set of n price and quantity series. A Divisia index of prices is obtained by cumulating the rate of change to the values of an index of price change, observation by observation. The index of price change is the weighted sum of the rates of change of the component prices. The weights are the current shares of the component goods in the total current expenditure on all the goods in the index.

A Divisia index is the ultimate extension of a chain index. A Divisia index of quantity can be obtained by applying the same strategy to quantities in place of prices, or, alternatively, by dividing total expenditure by the price index. However, the two quantity indices will not be exactly the same.

DIVIND (PNORM=<obs id>,PRINT,PVAL=<value>,QNORM=<obs id>,QVAL=<value>,TYPE=Q or P or N,WEIGHT=COMB or ARITH or GEOM) <name of output price index> <name of output quantity index> <list of pairs of input price and quantity series> ;

Usage

DIVIND has as its arguments the name to be given to the computed price index, then the name to be given to the computed quantity index, and finally the names of the series for prices and quantities of the components to be used as input to the calculations. The order is price for input one, quantity for input one, price for input two, quantity for input two, and so forth. No warning is given for non-positive prices for a quantity index, and vice versa (the formulas still hold unless WEIGHT=GEOM). When a quantity is zero for one or more periods, the series are spliced and the price is temporarily omitted from the index.

Output

Normally DIVIND produces no printed output, but stores the two computed index series in data storage. If the PRINT option is on, DIVIND prints a title, the options, the names of the input and output series, and a table of the two computed series labelled by the observation name.

Options

PNORM= identifies the observation where the price index is normalized. The index will have the value given by PVAL= at this observation. Note that there is no default for PNORM=.
PVAL= is the value to which the observation PNORM is to be normalized. The default is 1.0.

PRINT/NOPRINT tells whether the derived Divisia index series are to be printed. The default is no printing.

QNORM= identifies the observation where the quantity index is normalized. The index will have the value given by QVAL= at this observation. Note that there is no default for QNORM=.

QVAL= is the value to which the observation QNORM is to be normalized. The default is 1.0.

TYPE=Q specifies that the Divisia quantity index is to be computed and the price index is to be obtained by dividing total expenditure by the quantity index.

TYPE=P specifies that the Divisia price index is to be computed and the quantity index is to be obtained by dividing total expenditure by the price index.

TYPE=N specifies that both a Divisia price index and a Divisia quantity index are to be computed. Note that the product of the two indices will not be exactly proportional to total expenditure.

WEIGHT=ARITH specifies that the weights to be used in computing this period's rate of change of the index are the arithmetic averages of the shares in this period and the previous period.

WEIGHT=GEOM specifies that the weights are the geometric averages of the shares in this period and the previous period.

WEIGHT=COMB specifies that the weights are the geometric averages of (i) the arithmetic average, (ii) the share this period, and (iii) the share in the previous period.

Either QNORM= or PNORM= is required if TYPE=P,Q and both are required if TYPE=N.

The default values of the options are the following:

TYPE=Q,WEIGHT=COMB,NOPRINT,QVAL=1,PVAL=1

Examples

DIVIND (WEIGHT=ARITH, TYPE=P, PNORM=67) PRICEIN, QUANTIN, PS, QS, PND, QND, PD, QD;
FREQ Q;
DIVIND (WEIGHT=GEOM, TYPE=N, PNORM=75:1, PVAL=100,
QNORM=75:1, QVAL=100, PRINT) PI, QI, P1, Q1, P2, Q2, P3, Q3,
P4, Q4;

The first of these examples computes a Divisia price index as a weighted average of changes in PS, PND, and PD, using the shares of PS*QS, PND*QND, and PD*QD in total expenditure as weights. The series PRICEIN is normalized to have the value 1.0 in 1967 and QUANTIN is derived by dividing PRICEIN into total expenditure.

The second example computes a price index PI and quantity index QI independently from quarterly data. Both indices are normalized to have the value 100 in the first quarter of 1975. The weights are geometric averages of the shares in the adjacent years.

References


DO

Examples

DO specifies the beginning of a loop or grouped set of statements. The loop or group of statements must be terminated by an ENDDO ; statement.

DO ;
or
DO <index name> = <start value> TO <end value> [BY <increment>] ;
or
DO <index name> = <start value>, <end value> [, <increment>] ;

Usage

The first form of the DO statement (without arguments) is primarily used to specify the beginning of a block of statements which form a THEN or ELSE clause after an IF statement.

The other form of the DO statement specifies a conventional loop as in many programming languages. TSP executes the statements between the DO ... and ENDDO statement repetitively as many times as specified by the information given on the DO statement. The index or counter variable is set equal to the start value the first time through, and is changed each time through by the increment until the end value has been reached or exceeded. This test is done at the end of the loop, so the program always goes through once. DO loops can be nested with other DO loops or with DOT loops.

The start value, end value, and increment may be any real numbers (positive or negative), unlike some earlier versions of TSP which allowed only integers. If the increment is negative, obviously the index will be decremented by its absolute value, so the start value should be bigger than the end value.

The index variable is updated every time through the loop, so it may be used in computations or as a subscript. However, the DO loop in TSP is not a very efficient procedure, so that you should be wary of doing a substantial amount of variable transformation or computation with large DO loops. If you want the accumulated sum of a series, use a dynamic GENR -- for example,

\[ \text{ACSUM} = X; \]
\[ \text{SMPL} 2, N; \]
\[ \text{ACSUM} = \text{ACSUM}(-1) + X; \]

or

\[ \text{MSD (NOPRINT)} X; \]  \quad \text{? The result is in @SUM}
Loops with IFs are best done with logical expressions on the right hand side of a GENR, or with a `SMPLIF`.

Examples

```plaintext
DO I = 2 TO 7 BY 1 ;
  SET IM1 = I-1 ;
  SET X(I) = X(IM1) + X(I) ;
ENDDO ;
DO I = 2 TO 7 ;
  SET IM1 = I-1 ;
  SET X(I) = X(IM1) + X(I) ;
ENDDO ;
```

The first two examples here have the same effect, since the default value of the increment is one.

```plaintext
OLSQ Y C X1 X2 ;
IF ABS(@DW-2)>.5 ;
THEN ;
  DO ;
    AR1 Y C X1 X2 ;
    FORM EQ1 ;
  ENDDO ;
ELSE ;
  FORM EQ1 ;
```

This example runs OLS on an equation, checks the Durbin-Watson, and runs AR1 on the same equation if the Durbin-Watson is sufficiently different from two. The DO ... ENDDO statements bracket the set of statements which are to be executed if the Durbin-Watson test fails.
DOC

Options  Examples

DOC creates and maintains documentation for variables. Documentation can then be displayed with the SHOW command.

**DOC (ADD, REPLACE) <variable> 'description of variable';**

**Usage**

List the name of a variable (it doesn't have to exist yet) and give a description for it in quotes. There is no restriction on the length of the description, and you can separate lines in the description with backslashes (\). Note that the semicolon character (;) cannot be part of the description string due to TSP's rules regarding strings (see BASIC RULES, Number 2). TSP automatically maintains a date field for the variable if it has a description, and the DATE option in SHOW and DBLIST is used to display this.

SHOW and DBLIST will display as much of the description as they can fit on one line, following the other information on the variable. SHOW (DATE,DOC) will print the description on separate lines following the regular information. The DBLIST command also has the DATE and DOC options. The documentation is stored in .TLB (databank) files, and DBCOPY can be used to move it to other computers.

**Output**

No output is produced until the SHOW or DBLIST commands are used.

**Options**

ADD/NOADD specifies that the description should be added (appended) to any existing description.

REPLACE/NOREPLAC causes the description to replace any existing description.

**Examples**

**DOC CONS72 'Consumption in 1972 dollars\Source: ERP 1990';**

This is equivalent to

**DOC CONS72 'Consumption in 1972 dollars';**

**DOC (ADD) CONS72 \Source: ERP 1990';**
DOT

Options   Examples

DOT is the first statement in a DOT loop, which is like a regular DO loop, except that the values of the index are a series of character strings (names). Each of these names is substituted in turn each time through the loop wherever the symbol dot (.) appears in a variable name. The dot may appear anywhere in the variable name.

DOT (CHAR=<nesting level character>, INDEX=<variablename>, VALUE=<variablename>) <list of sector names or strings> ;

Usage

The primary use of DOT loops in TSP is the processing of multisectoral data, where the same group of statements is to be repeated on each sector. The names on the DOT command are the names of the sectors. They may also be integer numbers, which will be treated as the corresponding character string. Use the form `01', `02', etc. if you wish to include leading zeroes in the numbers.

Each DOT loop must be terminated by an ENDDOT statement. Any number of statements may appear between DOT and ENDDOT statements. TSP will cycle through them as many times as there are sectors on the DOT statement.

DOT loops may be nested up to ten deep and more than one dot included in each variable name (i.e., VAR...). The names substituted for the dots are taken in the order that the DOT statements appear, one from each statement. See the second example below. If there are fewer dots in a name than loops, the innermost loop is used first.

The DOT procedure cannot be used in a LOAD section; however, it can be used when reading data from files.

Legal dotted variable names are A., , .VAR , .D. , and AL.B .

Illegal dotted variable names are .1 , 2. (numbers), and .EQ. (logical operator).

Options

CHAR= special character to be used in conjunction with . in nested DOT loops, in order to specify explicitly which DOT statement is used to expand the period. It is normally used only to make a single . refer to the outer DOT loop when inside an inner loop.
INDEX= scalar variable name to hold the values 1,2,3, ... during the DOT loop. This is like having a DOT loop and a DO loop at the same time.

VALUE= scalar variable name to hold the values of numeric DOT sectors.

Examples

Suppose that you have data on the rate of change of wages (DW), the unemployment rate (U), and prices (P) for each of four countries: 1 (United States), 2 (England), 3 (Sweden), and 4 (Germany). When you create and load the data, give the series names like DW1, DW2, DW3, DW4, U1, U2, etc. Then you can easily run the same set of TSP statements on all four countries by means of a DOT loop. For example, to normalize the series to the same year compute the rate of change in prices, and run a regression of DW on the unemployment and rate of change of prices, use the following:

```
DOT 1-4 ;
NORMAL P. 72 100 ;
GENR DP. = LOG(P./P.(-1)) ;
OLSQ DW. C U. DP. ;
FRML EQ. DW. = A+B*U.+G*DP.*RHO ;
PARAM A B G RHO ;
LSQ EQ. ;
ENDDOT ;
```

If you wanted to work with data for the same four countries on prices and quantities of commodities, for example, food, housing, energy, etc., you could use the double dot construction:

```
DOT FOOD HOUS OIL ;
DOT 1-4 ;
GENR S..=P..*Q.. ;
ENDDOT ;
PRINT S.1-S.4 ;
ENDDOT ;
```

This example generates 12 series with the names SFOOD1, SFOOD2, SFOOD3, SFOOD4, SHOUS1-SHOUS4, and SOIL1-SOIL4, in that order. Each series is equal to the product of the corresponding price and quantity series. In the outer DOT loop, a table of the series for all the countries of each commodity is printed (note the use of the imbedded dot).

Here is another example with numbered sectors:

```
DOT (VALUE=J) 0-9 ;
SELECT COUNTRY=J;
OLSQ FISH C X ;
```

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FIT = @FIT; SET B. = @COEF(2);
ENDDOT;
SELECT 1;
PRINT COUNTRY FIT; PRINT B0-B9;

This example regresses the same dependent variable, FISH, on C and X in separate samples for each of 10 countries (same as PANEL (BYID,ID=COUNTRY) FISH C X;). The fitted values are saved and printed together.

Here is another example, showing the use of numeric character strings and double dots:

```
DOT `01` `10` `11`;
DOT FOOD HOUSE OIL;
   GENR S.. = P. * Q..
ENDDOT;
ENDDOT;
```

In this case the variables S01FOOD, S10FOOD, ..., S01HOUSE, etc. are created using a single price index PFOOD, PHOUSE, POIL, for each set of series Q01FOOD, Q10FOOD, etc. Note the use of a single dot for the P variable.

This example takes a list of variables called VARS and regresses each of them on the others in the list one by one. Note the use of the INDEX and CHAR options.

```
LIST VARS X Y Z W;
DOT (INDEX=I, CHAR=%) VARS;
   DOT (INDEX=J) VARS;
      IF I >= J; THEN; OLSQ .% C .
   ENDDOT;
ENDDOT;
```
DROP

DROP (Interactive)

Examples

DROP is used to drop a list of variables from the previous statement and re-execute it. It is the opposite of ADD.

\textit{DROP <list of variables>} ;

Usage

DROP is a convenient way to drop variables from a regression and perform a second estimation (without retyping the command). It is not restricted to this usage and may be used anywhere this type of command modification is needed.

The command

\textit{DROP var1 var2}

and the sequence

\textit{RETRY}
\textit{>> DELETE var1}
\textit{>> DELETE var2}
\textit{>> EXIT}

are identical in function since both permanently modify the previous command by deleting the first occurrences of \textit{var1} and \textit{var2}. The command is then automatically executed in both cases. The only potential difference between these approaches (besides the amount of typing) is in the definition of "previous". \textit{RETRY} with no line number argument assumes you want to modify the last line typed. DROP will not accept a line number argument, and always modifies the last line that \textit{is not itself} a DROP (or ADD) command.

Because of the way "previous" is defined for this procedure, you can execute a series of closely related regressions by entering the first estimation command, followed by a series of ADD and DROP commands. Since each ADD or DROP permanently alters the command, each new modification must take all previous modifications into account.

Note that it is not possible to combine ADD and DROP into one step to perform a REPLACE function, or to make compound modifications to a command. In these circumstances, RETRY must be used.

Output

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DROP echoes the modified command it will execute. Any further output will be a direct result of the command that is executed by it.

Examples

\texttt{OLSQ (WEIGHT=POP) YOUNG,C,RSALE,URBAN,CATHOLIC}

\texttt{DROP URBAN}

will run two regressions, the second of which is:

\texttt{OLSQ (WEIGHT=POP) YOUNG,C,RSALE,CATHOLIC}

This is also how the command will now look if you REVIEW it, since it has been modified and replaced in TSP's internal storage, and in your backup file.
DUMMY

DUMMY creates a set of zero-one variables which correspond to the different values taken by an input series. The number of dummy variables created is equal to the number of unique values of the input series (unless EXCLUDE is specified).

DUMMY (EXCLUDE, PREFIX=<name>) <series> [<listname> or <list of names>];

Usage

DUMMY followed by the name of a series will cause the variables NAME1, NAME2, etc. to be created, where NAME is the name of the series. To use another name as the prefix, include the PREFIX= option. If no input series is supplied and FREQ Q or FREQ M is in effect, Quarterly or Monthly dummies will be created, with the names Q1-Q4 or M1-M12. If a list of series is supplied, the dummies will be given the names in the list; be sure there are as many names as there are values of the variable.

Since the number of dummies created is usually equal to the number of unique values taken by the input series, care should be taken that the series used has a limited number of values. When a "continuous" rather than "discrete" variable is used for input, the number of dummies created could be equal to the number of observations, and storage allocation problems are likely to be the result.

It is well known that a complete set of dummy variables will be collinear with the constant term (intercept) in a linear regression. If you wish to use the dummies together with a constant in this way, you can create a set with one of the variables deleted by using the EXCLUDE option. In this case the number of variables created will be equal to the number of unique values taken by the input series less one.

Output

The set of dummy variables is stored in data storage, and the listname (if one was specified) is defined as the variable names for the set of dummies.

Options

EXCLUDE/NOEXCLUDE excludes the last dummy variable from the list. This option is useful if the list will be used in regressions with a constant term (to prevent multicollinearity).
**PREFIX**= Prefix for naming the dummy variables. The default prefix is the name of the input series.

**Examples**

```
FREQ Q; SMPL 75:1 85:4 ;
DUMMY;
```

creates Q1,Q2,Q3,Q4 (quarterly dummies). The series created have the following values:

<table>
<thead>
<tr>
<th>Obs</th>
<th>Q1</th>
<th>Q2</th>
<th>Q3</th>
<th>Q4</th>
</tr>
</thead>
<tbody>
<tr>
<td>75:1</td>
<td>1</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>75:2</td>
<td>0</td>
<td>1</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>75:3</td>
<td>0</td>
<td>0</td>
<td>1</td>
<td>0</td>
</tr>
<tr>
<td>75:4</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>1</td>
</tr>
<tr>
<td>76:1</td>
<td>1</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
</tbody>
</table>

and so forth.

```
FREQ M; SMPL 75:1 84:12 ; ? create M1-M11
DUMMY (EXCLUDE); ? (monthly dummies with M12 excluded).
```

The next example creates a list of dummies from a variable, SIZE, which takes on 3 values: 0, 2, and 3.5.

```
DUMMY SIZE SDLIST;
```

This is equivalent to the following statements:

```
SIZE1 = SIZE = 0;
SIZE2 = SIZE = 2;
SIZE3 = SIZE = 3.5;
LIST SDLIST SIZE1-SIZE3;
```

The next example creates a set of year dummies for panel data, assuming you have a variable YEAR which takes on values from 72 to 91:

```
DUMMY YEAR YEAR72-YEAR91 ;
```

This command creates 20 dummy variables: YEAR72, YEAR73, YEAR74, and so forth.

To create individual dummies for balanced data, using **TREND** and **INT()**, see the example under **AR1**.
EDIT

EDIT (Interactive)

Examples

EDIT is a simple editor providing the capability to perform argument modifications on a TSP command during an interactive session. If you are using the Windows or DOS versions of TSP you will not need this command, as you will be able to use arrow-key editing (for interactive use) or the Windows interface programs Oxmetrics or Through the Looking Glass (for editing batch files).

EDIT [<line number>] ;

Usage

Only one argument is allowed with this command, which must be a TSP line number. If this argument is omitted, EDIT will prompt you for modifications to the previous line. This command is generally used for correcting typos, or modifying lists of series, etc... before requesting re-execution of a procedure.

RETRY is identical to EDIT, except that execution of the modified command is automatic upon exit. Also, EDIT is treated as a special case in COLLECT mode (its execution is not suppressed), while RETRY is not.

The editor will first echo the command you wish to modify, then issue the prompt ">>": Responses to the prompt must consist of an editing command followed by appropriate arguments. Any unique abbreviation for an editing command will suffice (including a single letter). Complete arguments must be entered, even if you only wish to replace a single character. Only one modification per edit prompt may be made. Prompting will continue until an EXIT command or carriage return is given in response.

Arguments:

TSP breaks up everything you type (except data) into a string of "arguments". Each series or variable name is an argument, as is each operator. Statement terminators (semi-colons) and item separators (commas and blanks) are not considered arguments. This example has 10 arguments:

```
GENR GNPN = GNP / ( DELTA + R ) ;
```

<table>
<thead>
<tr>
<th>Argument</th>
<th>Contents</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>GENR</td>
</tr>
<tr>
<td>2</td>
<td>GNPN</td>
</tr>
<tr>
<td>3</td>
<td>=</td>
</tr>
<tr>
<td>4</td>
<td>GNP</td>
</tr>
<tr>
<td>5</td>
<td>/</td>
</tr>
</tbody>
</table>
The editing commands and their arguments are as follows:

**EXIT**

Editing completed (takes no arguments). A simple carriage return will also be interpreted as EXIT.

**DELETE arg n**

Delete the nth occurrence of "arg" in the command.

**REPLACE arg1 arg2 n**

Replace the nth occurrence of "arg1" with "arg2" in the command.

**INSERT arg1 arg2 n**

Insert "arg1" after the nth occurrence of "arg2" in the command. If there is only one argument provided, it is inserted at the end of the command.

**NOTE:** "n" is always assumed to be 1 if absent.

**Restrictions:**

1. Subscripts: you will not be able to edit successfully double subscripts, or subscripts with dates.

2. Parentheses: the editor will not successfully find specific parentheses in commands which contain lags, leads, or subscripts prior to the parenthesis you are specifying.

**Examples**

One type of modification you may wish to make is to change the list of options on a command without having to retype the whole thing (particularly if it is lengthy). As a simple example, here is how you might change a static forecast into a dynamic one, as well as request printing and plotting:

**5? FORCST (STATIC,DEPVAR = I) IFIT**

<no output since the print option is off>
6? EDIT 5
>> INS PRINT
>> REP STATIC DYNAMIC
>> EX
5. FORCST (PRINT,DYNAMIC,DEPVAR = I) IFIT ;

In this case, execution of the modified command would not take place until specifically requested.

Another common modification is to fix a typo. If you have made a typing error in line 10, for instance, and wish to correct it the sequence might look like this:

10? INT DP,DP1,LGNP,TIME,C INVR C,G,LM,TIME ;
<error message because procedure INST is misspelled>
11? EDIT 10
>>REPLACE INT INST 1
>>EXIT
10? INST DP,DP1,LGNP,TIME,C INVR C,G,LM,TIME ;
12? EXEC 10
10? INST DP,DP1,LGNP,TIME,C INVR C,G,LM,TIME ;
<output from the INST command.>

There are a number of ways that the previous example could be simplified to reduce typing. First of all, commands may be abbreviated. Note that INT was not a valid abbreviation for INST (see Basic Rules). Second, n=1 is the default on the REPLACE command as is carriage return for the EXIT command, so they may be omitted. Also since line 10 is the previous command, it may be omitted from the EDIT command. Lastly, statements 11 and 12 may be combined by substituting the RETRY command for EDIT. Assuming the same error on line 10, the correction would now look like this:

11? RET
>> R INT INST
>>
10? INST DP,DP1,LGNP,TIME,C INVR C,G,LM,TIME ;
<output from the INST command.>
ELSE

*Example*

ELSE signals that the statement or DO group of statements immediately following are to be executed if the last IF clause had a false result. The full syntax of the IF-THEN-ELSE sequence is IF expression; THEN; statement, or block of statements; ELSE ; statement, or block of statements ;.

ELSE ;

*Usage*

ELSE has no arguments. It is optional following an IF statement. If several ELSE statements appear in sequence, the first refers to the most recent IF clause, the second to the next most recent, and so forth. If more than one statement (or an INPUT command) is to be executed when the IF clause is false, enclose all the statements in a DO ; .... ENDDO ; group.

*Example*

```plaintext
IF @SSR>LIMIT ; THEN ;
   SET RESULT=@SSR ;
ELSE ; DO ;
   GENR Y = Y+INCR ;
   OLSQ Y C POP TIME ;
   SET RESULT=@SSR ;
ENDDO ;
```

In the above example, the statements in the DO group are executed once and only once when the original value of @SSR is less than or equal to the value of LIMIT.
**Example**

END terminates both the TSP program section and the TSP data section.

**Usage**

END has no arguments. END is also used as a delimiter. In the data section it is used to mark the end of the current section of data and to force a return to execution of the TSP program. If there is more than one LOAD; statement in the program section, each one will terminate at successive END statements in the data section.

**Example**

```
NAME USER;
LOAD;
........ execution with data from load section..............
STOP ; END ;
........ load section with data ................................
END ;
```
ENDDO

ENDDO is used to close DO loops.

ENDDO ;

Usage

ENDDO takes no arguments. Every DO statement must have an associated ENDDO statement somewhere following it in the program. The ENDDO statement always applies to the last DO which was encountered so that DO loops may be nested to any level.

ENDD or END DO are synonyms for ENDDO.
ENDDOT

ENDDOT is used to close DOT loops.

Usage

ENDDOT takes no arguments. Every DOT statement must have an associated ENDDOT statement somewhere following in the program. The ENDDOT statement always applies to the last DOT which was encountered so that DOT loops may be nested to any level.

END DOT is a synonym for ENDDOT.
ENDPROC

ENDPROC is used to end user PROCs.

**ENDPROC [<name of PROC>] ;**

**Usage**

ENDPROC takes one (optional) argument, the name of the PROC to which it belongs. Every PROC statement must have an associated ENDPROC statement somewhere following in the program. PROCs may be nested to any level, but if there is an ENDPROC statement missing for one of them, a fatal error will be trapped.

ENDP or END PROC are synonyms for ENDPROC.
ENTER

ENTER (Interactive)

Example

ENTER allows you to input data from the terminal in an interactive session. It will prompt you for observation as the data is entered. Use the READ command in batch mode.

ENTER <list of series names> ;

Usage

ENTER must have at least one argument; all prompting and storage will be defined by the most recent SMPL and FREQ prior to the ENTER command. In response to the prompt for data, you may enter as many items per line as you like -- the prompts will adjust accordingly. Prompting will cease when the current SMPL has been satisfied, and the series stored. If more than one series is being entered, you will be prompted to enter them sequentially.

If a numeric error is made in data entry, make a note of which observation(s) and finish entering the data. Then the UPDATE command may be used to correct the error. If a non-numeric entry is encountered you will be prompted to go back and correct it. Missing values may be stored by creating a SMPL with gaps (prompting will adjust) or using the missing value code "." .

Example

1? FREQ A
2? SMPL 1946,1975
3? ENTER GNP
Enter data for GNP
1946? 475.7 468.3 487.7 490.7
1950? 533.5 578.5 598.5 621.8 613.7
1955? 654.8 668.8 680.9 679.5 720.4 736.8 755.3
1962? 799.1 830.7 874.4 925.9 981.0 1007.7
1968? 1051.8 1078.8 1075.3 1107.5 1171.1
1973? 1233.4 1210.7 1186.4

GNP will be stored with 30 observations
EQSUB

**Options**    **Example**

EQSUB substitutes one or more equations into another. This is useful for estimation with several parameter restrictions, with long equations which have common terms, or for setting up a complicated model where the exogenous variables can be changed later by just changing one of the input equations.

**EQSUB** *(LAGS, NAME=<new output equation name>, PRINT) <main equation name> <list of input equation names> ;*

**Usage**

Define the main equation and the input equation(s) with **FRML** or **IDENT** statements. The EQSUB command substitutes each input equation into the main equation, in order from left to right. For each input equation, EQSUB looks for its dependent variable (or equation name, if there is no dependent variable) in the argument list of the main equation. If the dependent variable is found, the code from the input equation is inserted into the main equation, and the old variable name is deleted. For example:

**FRML EQ1 Y = A + XB;**
**FRML EXB XB = X1’B1 + X2’B2;**
**EQSUB EQ1 EXB;**

is equivalent to:

**FRML EQ1 Y = A + X1’B1 + X2’B2;**

The resulting equation replaces the main equation, unless the **NAME=** option is supplied. The **DOT** command is useful when there are several different main equations.

If you have many component input equations to define, it may be convenient to leave out the dependent variable name, so that you don't have to invent both a dependent variable and an equation name for each one. Such an equation is called "unnormalized" in TSP. For example,

**FRML E Y1 - XB; FRML XB B0 + B1’X1 + B2’X2;**
* change XB to add/delete exog. errors variables
**FRML TOBIT LOGL = YPOS*(LNORM(E/SIGMA) - LOG(SIGMA)) + YZERO*LCNORM(-XB/SIGMA);**
**EQSUB(NAME=TOBIT1) TOBIT E XB;**
Note that both TOBIT and E depend on XB, so XB is substituted in last. There is no need to substitute XB into E separately. A separate substitution would still operate correctly, but it would result in larger and less efficient code. The new FRML TOBIT1 is created, and the original TOBIT is left untouched for later use.

On the other hand, normalized input equations are recommended for parameter restrictions. The same FRMLs can usually be used to both impose parameter restrictions, and to evaluate the restricted parameters after estimation with ANALYZ. If the input equation is normalized, ANALYZ can store the restricted parameter name. For example, in a translog model with symmetry imposed:

```
FRML EQ1 SH1 = A1 + B11*LP1 + B12*LP2 + B13*LP3;
FRML EQ2 SH2 = A2 + B12*LP1 + B22*LP2 + B23*LP3;
```

\[\text{?Note: The last share equation is not used in estimation due to} \]
\[\text{? singularity.} \]

```
FRML EQ3 SH3 = A3 + B13*LP1 + B23*LP2 + B33*LP3;
? homogeneity/adding up constraints
FRML R13 B13 = -(B11+B12);
FRML R23 B23 = -(B12+B22);
EQSUB EQ1 R13; EQSUB EQ2 R23;
```

\[\text{? left out params from final equation} \]

```
FRML LO1 A3 = 1 - (A1+A2);
? note: this also depends on the restricted B23
FRML LO2 B33 = -(B12+B23);
EQSUB LO2 R23;
```

\[\text{LSQ EQ1 EQ2; ? Estimate model with restrictions in place} \]
\[\text{? Print and store values and standard errors for A3 and B33} \]

```
ANALYZ LO1 LO2;
EQSUB can also handle "lagged dependent variables"
FRML U Y - (A + B*X + G*Z(-2));
FRML E U - RHO*U(-1);
EQSUB U E;
```

is equivalent to

```
FRML E Y - (A+B*X+G*Z(-2)) - RHO*(Y(-1) - (A+B*X(-1)+G*Z(-3)));
```

Note that when the EQSUB command is given, all the variables in the equations must exist (either as series, PARAMs, CONSTs, or other FRMLs), so that EQSUB will know which ones need to be lagged (the series and FRMLs), and which ones don't need lags (the PARAMs and CONSTs).
Normally, the output equation is stored silently, replacing the input equation, or creating a new equation. If the PRINT option is on, the output equation is printed.

Options

LAGS/NOLAGS controls substitution for the dependent variable name when it is lagged. When the NOLAGS option is specified only the unlagged appearances of the dependent variable are substituted for.

NAME= new output equation name supplies a new name for the output equation. If this option is not present, the main equation is overwritten by the new one.

PRINT/NOPRINT controls whether the output equation is printed.

Example

See above. See also the TSP User's Guide for many additional examples of using EQSUB to set up log likelihood equations for estimation by ML. Here is one more example, illustrating the use of DOT to substitute input equations F1 to F20 into main equations E1 to E8:

DOT E1-E8;
   EQSUB . F1-F20;
ENDDOT;
EXEC

EXEC (Interactive)

EXEC forces execution (or re-execution) of a range of lines consisting of TSP commands that have already been entered in the interactive session via keyboard or input file.

EXEC [<first line number>], [<last line number>] ;

Note that in DOS/Win TSP, it is easier to use the up/down arrow keys to select and rerun a single command.

Usage

EXEC varies slightly depending upon the mode in which you are currently operating. In COLLECT mode, EXEC is used to execute the range of lines just collected, and return control to interactive mode. This is considered the standard exit from collect mode (the alternative is to suppress execution with the EXIT command). The whole range will be executed, so line number arguments will be ignored if you supply them. Lines in the range may be EDITed or DELETEd if necessary before EXECuting them.

In interactive mode, up to two arguments may be supplied with the EXEC command. If no argument is supplied, the previous command is re-executed. If only the first line number is supplied, a single line is executed, and if two line numbers are given, their inclusive range is executed. In any case, these lines may be commands that have been previously executed and edited, or commands that were entered but suppressed with an EXIT command in collect mode or at the end of an INPUT file.
EXIT (Interactive)

EXIT terminates the current operating mode of the program.

EXIT ;

Usage

If used in interactive mode, the interactive session will be terminated, returning control to the operating system; this is the same as typing STOP or END.

If used in collect mode, this command will return the user to the interactive level of the program WITHOUT executing the commands just collected (use the EXEC command to leave collect mode with automatic execution). EXIT may also be used to replace END at the very end of an INPUT file to suppress automatic execution upon completion of reading the file (see INPUT).
FETCH

Example  Reference

FETCH reads microTSP and EViews format databank files.

FETCH <list of series> ;
FETCH [disk:]seriesname [ [disk:]seriesname ] ;

Usage

MicroTSP-format databank files may be useful for transferring data between microTSP/EViews and TSP. They are plain (editable, non-binary) files containing comments, frequency, starting and ending dates, and data values (one per line). See the microTSP/EViews documentation for details. They are not efficient in terms of disk space usage or the time required to read or write them. However, they are easy to edit, for manual data revision. To fetch a series from a microTSP databank name series .DB, use the command

FETCH series ;

series.DB must be in the default directory.

To move regular TSP databanks between machines (such as VAX/VMS to a personal computer), use the DBCOPY command.

The STORE command creates the files read by FETCH.

Example

FETCH X Y;

reads the series X and Y (and any imbedded comments and documentation) from the files X.DB and Y.DB.

Reference

FIML invokes the Full Information Maximum Likelihood procedure. This procedure obtains maximum likelihood estimates of a nonlinear simultaneous equations model. The model should have N equations (some of which may be identities) in N endogenous variables and may be written in implicit form (equations without left-hand-side variables).

FIML is an asymptotically efficient estimator for simultaneous models with normally distributed errors. It is the only known efficient estimator for models that are nonlinear in their parameters.

For further details on this estimator and the method of estimation, see the references and the TSP User's Guide. See the LIML command for details on doing nonlinear Limited Information Maximum Likelihood (single equation ML) with the FIML command.

FIML (ENDOG=(<list of endogenous variables>, FEI), nonlinear options) <list of equation names> ;

Usage

FIML in its simplest form is invoked by listing the endogenous variables of the model in the options and following the options by the equation names:

FIML (ENDOG=(Y1,Y2,...,YN)) EQ1,EQ2,.....EQN ;

The equations must be previously defined by FRML and IDENT statements. Those which are IDENTs are assumed to hold exactly in the data and will not contribute to the covariance matrix of the disturbances. If the IDENTs can be substituted into the FRMLs, a FIML estimation of the resulting FRMLs will yield the same results as FIML on the original FRMLs and IDENTs.

IDENTs cannot be used to impose nonlinear (or linear) constraints on the parameters -- any constraints must be substituted directly into the structural equations. An attempt to use IDENTs in this way would fail because of an unequal number of endogenous variables and equations, and a singular Jacobian (the gradient of the equations with respect to endogenous variables).

The parameters to be estimated must be defined previously and starting values assigned by a PARAM statement or statements (or by being estimated in a previous nonlinear estimation).
If the equations are nonlinear in the endogenous variables, the Jacobian will not be constant over the series, and it will be evaluated at each observation. If the Jacobian is singular at the initial iteration, estimation halts. Usually this is caused by coefficients with zero starting values -- use 0.1 for these initial values instead. If the Jacobian is singular during the iterations, the parameter stepsize is automatically squeezed (this is treated as a numerical error).

FIML detects whether the Jacobian is constant or identity and speeds up the computations accordingly.

Output

Normal FIML output begins with a listing of the options (if the PRINT option is specified) and the equations. The model is checked for linearity in the parameters (which simplifies the computations) and for linearity in the variables (which greatly simplifies the derivatives). A message is printed if either form of linearity is found. The amount of working space used by FIML is also printed - this number can be compared with the amount printed at the end of the run to see how much extra room you have if you wish to expand the model.

Next FIML prints the starting conditions for the constants and parameters, and then iteration-by-iteration output. If the print option is off, this output consists of only one line, showing the beginning value of the (minus) log likelihood, the ending value, the number of squeezes in the stepsize search (ISQZ), the final stepsize, and a criterion which should go to zero rapidly if the iterations are well-behaved. This criterion is the norm of the gradient in the metric of the Hessian approximation. It will be close to zero at convergence.

When the print option is on, FIML also prints the value of the parameters at the beginning of the iteration and their direction vector. These are shown in a convenient table so that you can easily spot parameters with which you are having difficulty.

Finally FIML prints the results of the estimation (whether or not it converged); these results are printed when the print option PRINT, NOPRINT, or TERSE, but not when SILENT is specified. The names of the equations and endogenous variables are printed, the value of the log likelihood at the maximum, and the corresponding estimate of the covariance of the structural disturbances.

Following this is a table of parameter estimates and asymptotic standard errors, as well as their estimated variance-covariance matrix (if it has been unsuppressed).
For the default HCOV=B option, this is computed by summing the outer product of the gradient vector for all structural parameters and error covariance parameters over all observations, inverting this matrix (the BHHH matrix), and taking the submatrix corresponding to the structural parameters. This is a consistent estimate of the information matrix, with good small sample properties. Note that the BHHH matrix is not block diagonal between the structural and error covariance parameters at the maximum even though the second derivative matrix is block diagonal at the maximum, so this procedure is necessary. See Calzolari and Panattoni (1988) for details.

HITER=U and HCOV=U use numeric second derivatives for iteration and computing the variance estimate respectively, using equations (25.3.23) and (25.3.26) in Abramovitz and Stegun (1972). Numeric second derivatives can provide a very close approximation to the true Hessian. The drawback is that computing them is relatively slow, requiring 2*K*K function evaluations for a model with K parameters. HITER=U yields quadratic convergence during iterations, which can be faster than HITER=F (BFGS method) if the number of parameters is less than about 7. HCOV=U provides standard errors which are more reliable than BFGS (HCOV=F often produces "false zero" standard errors). They match HCOV=N standard errors to 3-5 digits in the tests we've performed.

HITER=C and HCOV=C use a discrete Hessian for iteration and computing the variance estimate respectively. This is a numeric difference of analytic first derivatives. This is even more accurate than HCOV=U in terms of matching HCOV=N results -- it's usually good to 6+ digits in standard errors. It also requires 2*K derivative evaluations for a model with K parameters. This option is only useful in a command that has analytic first derivatives but not analytic second derivatives. The most important such command in TSP is FIML. However, note that most models estimated by LSQ can also be estimated using FIML; typically convergence will be much faster using FIML (HITER=C rather than LSQ with the default iteration).

FIML also stores its results in data storage. The estimated values of the parameters are stored under the parameter names. In addition, the following results are stored:

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@RNMS</td>
<td>list</td>
<td>#params</td>
<td>Parameter names.</td>
</tr>
<tr>
<td>@LOGL</td>
<td>scalar</td>
<td>1</td>
<td>Log of likelihood function.</td>
</tr>
<tr>
<td>@SBIC</td>
<td>scalar</td>
<td>1</td>
<td>Schwarz-Bayes information criterion with nobs=</td>
</tr>
<tr>
<td>@AIC</td>
<td>scalar</td>
<td>1</td>
<td>Akaike information criterion</td>
</tr>
<tr>
<td>@NCOEF</td>
<td>scalar</td>
<td>1</td>
<td>Number of parameters in model</td>
</tr>
<tr>
<td>@NCID</td>
<td>scalar</td>
<td>1</td>
<td>Number of identified parameters in model</td>
</tr>
<tr>
<td>Variable</td>
<td>Type</td>
<td>Description</td>
<td></td>
</tr>
<tr>
<td>----------</td>
<td>------</td>
<td>-------------</td>
<td></td>
</tr>
<tr>
<td>@IFCONV</td>
<td>scalar</td>
<td>1</td>
<td>Convergence status (1 = success).</td>
</tr>
<tr>
<td>@GRAD</td>
<td>vector</td>
<td>#params</td>
<td>Gradient of likelihood at maximum.</td>
</tr>
<tr>
<td>@COEF</td>
<td>vector</td>
<td>#params</td>
<td>Estimated values of parameters (also stored under their names).</td>
</tr>
<tr>
<td>@SES</td>
<td>vector</td>
<td>#params</td>
<td>Standard errors of estimated parameters.</td>
</tr>
<tr>
<td>@T</td>
<td>vector</td>
<td>#params</td>
<td>asymptotic T-statistics</td>
</tr>
<tr>
<td>%T</td>
<td>vector</td>
<td>#params</td>
<td>p-values for asymptotic T-statistics (based on normal distribution)</td>
</tr>
<tr>
<td>@SSR</td>
<td>vector</td>
<td>#eqs</td>
<td>Sum of squared residuals for each equation, stored in a vector.</td>
</tr>
<tr>
<td>@S</td>
<td>vector</td>
<td>#eqs</td>
<td>Standard error of each equation, stored in a vector.</td>
</tr>
<tr>
<td>@DW</td>
<td>vector</td>
<td>#eqs</td>
<td>Durbin-Watson statistic for each equation, stored in a vector.</td>
</tr>
<tr>
<td>@RSQ</td>
<td>vector</td>
<td>#eqs</td>
<td>R-squared for each equation, stored in a vector (if eqs are normalized)</td>
</tr>
<tr>
<td>@ARSQ</td>
<td>vector</td>
<td>#eqs</td>
<td>Adjusted R-squared for each equation (if eqs are normalized)</td>
</tr>
<tr>
<td>@YMEAN</td>
<td>vector</td>
<td>#eqs</td>
<td>Vector of means of dependent variables (if eqs are normalized)</td>
</tr>
<tr>
<td>@SDEV</td>
<td>vector</td>
<td>#eqs</td>
<td>Vector of standard deviations of dependent variables (if eqs are normalized)</td>
</tr>
<tr>
<td>@VCOV</td>
<td>matrix</td>
<td>#par* #par</td>
<td>Estimated variance-covariance of estimated parameters.</td>
</tr>
<tr>
<td>@COVU</td>
<td>matrix</td>
<td>#eqs*#eqs</td>
<td>Residual covariance matrix (the # of eqs is the number of structural equations).</td>
</tr>
<tr>
<td>@FIT</td>
<td>matrix</td>
<td>#obs*#eqs</td>
<td>Matrix of fitted values (if equations are normalized)</td>
</tr>
<tr>
<td>@RES</td>
<td>matrix</td>
<td>#obs*#eqs</td>
<td>Matrix of residuals</td>
</tr>
</tbody>
</table>

**Options**

**ENDOG** (list of endogenous variables). This defines the endogenous variables, including those that do not appear on the left hand side of any behavioral equation. There is no requirement that endogenous variables appear on the left side of any equation, since FIML estimates are invariant to this normalization. However, the number of endogenous variables must be equal to the number of equations.
FIML/NOFEI specifies that models with additive individual fixed effects are to be estimated. The panel structure must have been defined previously with the FREQ (PANEL) command. The equations specified must be linear in the parameters (this will be checked) and variables.

**Nonlinear options** These options control the iteration methods and printing. They are explained in the NONLINEAR section of this manual. Some of the common options are MAXIT, MAXSQZ, PRINT/NOPRINT, and SILENT/NOSILENT.

The legal choices for HITER= are G (Gauss, the default), B (BHHH), and D (DFP -- numeric derivatives). HCOV=B (BHHH) is the default method for calculating standard errors, and D is legal when HITER=D is used.

**Examples**

To obtain full information maximum likelihood estimates of the illustrative model from the User’s Manual, use the following statement:

```plaintext
FIML (ENDOG=(GNP,CONS,I,R,LP)) GNPID, CONSEQ, INVEQ, INTRSTEQ, PRICEQ ;
```

To do Box-Cox regression properly, use FIML to include the Jacobian term in the likelihood (note unnormalized equation):

```plaintext
FRML EQ1 (Y**LAM-1)/LAM - (A + B*X);
PARAM A B LAM 1 ;
FIML(ENDOG=Y) EQ1;
```

Klein-I model (see Calzolari and Panattoni for correct results):

```plaintext
FORM (VARPREF=C_) CONS CX C P(-1) W ;
FORM (VARPREF=I_) INV I C P(-1) K(-1) ;
FORM (VARPREF=W_) WAGES W1 C E E(-1) TM ;
IDENT WAGE W = W1+W2 ;
IDENT BALANCE CX+I+G - (TX+W+P) ;
IDENT PPROD E = P+TX+W1 ;
FIML (ENDOG=(CX,I,W1,W,P,E)) CONS INV WAGES WAGE BALANCE PPROD ;
```

**References**


FIND (Interactive)

Example

FIND lists all lines entered in the session so far that begin with the specified TSP command.

FIND <TSP command> ;

Usage

FIND is useful in conjunction with other interactive commands that use line number as arguments. You may want to EDIT or EXEC a particular command, but cannot remember its line number. The line could be found with REVIEW, but if it has been a long session, this could take some hunting.

FIND will accept only one TSP command as an argument.

Example

FIND OLSQ

will list all OLSQ commands you have entered or read in during the session along with their line numbers.
FORCST

Output   Options   Examples   Reference

FORCST allows you to use the results of any linear equation estimation routine in TSP to compute predicted values of the dependent variable over observations which may be the same or different from those used for estimation. You may also do a forecast using a different set of exogenous variables. The model for the forecast is either that specified on the previous linear (index) estimation procedure (OLSQ, INST, PROBIT, TOBIT, ORDPROB, POISSON, NEGBIN, ARCH or AR1) or you may supply the model yourself using various options. To compute predicted values after a nonlinear estimation procedure (LSQ or FIML) use the GENR, SIML, or SOLVE commands. Use BJFRCST if your model was estimated by Box-Jenkins techniques in BJEST.

\[\text{FORCST (COEF=\text{<vector name>}, DEPVAR=\text{<var name>}, DYNAM or STATIC, PRINT, RHO=\text{<scalar>}) <predvarname> [\text{<list of indep variables>}] ;}\]

Usage

The simplest form of FORCST follows the estimation command which specifies the model to be used for prediction; no other estimation commands can intervene. You should change the sample with a SMPL statement between the estimation and the forecast if you want to forecast for a different time period. The statement in this case is:

\[\text{FORCST <name to be given to predicted variable> ;}\]

Include a PRINT option with the command to have the results printed and plotted.

The other way to use the FORCST command does not require it to appear immediately following the estimation. However, you must specify the coefficient vector, the names of the right hand side variables, and, if a serial correlation correction is desired, the value of RHO.

If STATIC or NODYNAM is specified, FORCST will treat a lagged dependent variable like any other exogenous variable in computing the forecast. However, if you specify the (default) DYNAM option on the FORCST statement, FORCST will feed back the fitted values into the lagged dependent variable dynamically. As an initial condition the actual lagged dependent variable is used in the first period.
When FORCST follows an AR1 regression, the forecast is computed including a serial correlation correction using the estimated value of rho from the regression. This means that the dependent variable $y$ must be available to the program so that

$$
\hat{y}_t = X_t \hat{\beta} \quad \text{if } t = 1
$$

$$
\hat{y}_t = X_t \hat{\beta} + y_{t-1} - \hat{\rho} X_{t-1} \hat{\beta} \quad \text{if } t > 1
$$

may be computed. This "static" forecast is that computed by FORCST when STATIC is specified. To obtain a true dynamic forecast or extrapolation, use the (default) DYNAM option. In this case, FORCST will look for presample data to calculate a presample residual. In either case, unless the FORCST statement immediately follows the AR1 estimation of interest, the name of the dependent variable must appear in the DEPVAR= option.

If there are any gaps in the SMPL vector, FORCST will treat the observations between each pair of SMPL numbers separately. If the DYNAM option is off, the output will be identical to that of a single SMPL over the entire period except for the indicated gaps. With RHO not zero or the DYNAM option, forecasting will start anew with each pair of SMPL numbers (that is, it will start with new initial conditions), and thus the results will be different than a single forecast over the entire sample.

Output

When the PRINT option is off, no output is printed by FORCST and only the single forecasted series is stored in data storage.

When the PRINT option is on, the procedure prints a title, the vector of coefficients used in the forecast, the serial correlation parameter, and whether the forecast is static or dynamic. A plot of the forecasted series is printed which has the observation's name down the left hand side and the values of the series printed on the right hand side. The series is also stored.

Options

**COEF**= the name of a vector containing the coefficient estimates to be used in the forecast. This could be the vector of coefficients stored under the name @COEF after a previous estimation. The order of the coefficients in this vector should match the order of the names of the right hand side variables in the forecast model.

**DEPVAR**= the name of the dependent variable in the estimation model. This option is necessary to obtain correct dynamic forecasts when the FORCST procedure is not executed immediately following the estimation procedure.
FORCST

DYNAM/STATIC specifies whether the forecast is to be dynamic or static. A static forecast uses historical (supplied by the user) values for the lagged endogenous variable(s) throughout the forecast period, while a dynamic forecast uses the lagged forecasted values whenever it can (i.e., in the second observation for lag one, third observation for lag two, and so forth). This also applies to the lagged endogenous variable which appears in the residual in an AR(1) forecast. Obviously, STATIC is the default for non-AR1 forecasting when there is no lagged dependent variable.

PRINT/NOPRINT specifies whether or not the forecast is to be printed and plotted. If this option is on, a title and a description of the forecasting model are also printed.

RHO= the value of the serial correlation parameter if an AR1 forecast is being requested and the value from the immediately preceding estimation is not wanted.

Examples

In the example below, an OLSQ equation is used to extrapolate into a future time period; values of the exogenous variables for that time period should already have been loaded. The extrapolated values of CONS are stored under the name CONSP; they are also printed and plotted (because of the PRINT option).

SMPL 1 20;
OLSQ CONS C GNP;
SMPL 21 30;
FORCST (PRINT) CONSP;

The example below computes a series CONSP using the equation estimated by OLSQ, but with a different series for GNP on the right hand side. If an AR1 forecast of this type were desired, the serial correlation coefficient would be specified as a RHO= option in the parentheses also.

SMPL 1 20;
OLSQ CONS C GNP;
COPY @COEF B ;

......
FORCST (COEF=B) CONSP C GNPNEW;

This example shows the use of the DYNAM option to obtain a dynamic forecast with a lagged endogenous variable on the right hand side of the equation (I(-1)).

SMPL 1 20;
OLSQ I I(-1) GNP ;
This example shows both uses of the FORCST procedure; the dynamic option was specified on the first statement so that an extrapolation of the estimating equation will be produced. The statements which save R and B were required only for the second forecast, which is over the same sample as the original estimation, but uses a different right hand side variable, GNPNEW.

The example below shows how to compute a set of forecasts for a sales variable based on three slightly different GNP projections:

Reference

FORM

Options   Examples

FORM makes a TSP equation (FRML) from the results of a linear estimation procedure or from a list of names (such as regression variables). The FRML can then be used in an estimation or simulation model; it can have either names or constants as its coefficients. The NAR option makes it easy to estimate linear regression equations with AR(p) errors -- just use the resulting FRML in LSQ for direct nonlinear estimation.

FORM (COEFPR=<coefficient prefix>, NAR=<number of AR terms>, PARAM, PRINT, RESIDUAL, RHOPREF=<rho prefix>, SUM, VALUE, VARPR=<coefficient prefix>) <equation name> [<list of names>];

or
FORM (VALUE) <list of equation names>;

Usage

The first argument to FORM is required; it is the name you wish to give to the equation. If there is only one argument, FORM must follow the linear estimation command (OLSQ, INST, LIML, VAR, or AR1) which created the results you wish to save as an equation. In this case, the equation created by FORM is the same equation that would be used by FORCST to generate a single equation forecast. If you GENRed this equation, you would obtain a static forecast just as if you had run the FORCST procedure (but with a slight loss of precision). The variable names are retrieved from @RNMS and @LHV, while the coefficient values are taken from @COEF and @RHO.

If there are three or more arguments, FORM does not look for a previous estimation; instead it creates a new equation. In this case, the variable names are the remaining arguments and the coefficient names are constructed mechanically (zero starting values are used). Note: having exactly two arguments is invalid unless the previous estimation was a VAR with two equations, because it implies a dependent variable but no right hand side.
The output equation can be used for estimation or for simulation, or in any application which can use equations (such as ANALYZ, DIFFER, EQSUB, etc.). The PARAM option is used for estimation; in this case, parameter names are inserted into the FRML and values are also stored under these names (as if a PARAM statement had been issued). These names are normally created by appending 0, 1, 2, etc. to a coefficient prefix such as B. If there is a constant term (C), the number start with 0; otherwise they start with 1. PARAM names can also be created from the original variable names, using the VARPREF option. In either case, the PARAM names are stored in the list @RNMSF, which can be used in commands like UNMAKE to fill them with new starting values.

For simulation, use the FORM command after you estimate a linear equation; constant values are inserted into the FRML. FRMLs with parameter names can also be used for simulation; using constant values is just slightly more compact and it also prevents the values from being changed inadvertently.

Output

Normally FORM produces no printed output. If the PRINT option is on, the options and the output equation are printed. A TSP equation is stored in data storage under the name supplied by the user. If the PARAM option is on, scalar PARAMeters will be stored, and a list of the parameter names is stored under @RNMSF.

Options

COEFPREF= a prefix for the coefficient names. The default is B plus the equation name. Specify this carefully to avoid overwriting existing variables. For example, avoid creating A0-A11 and then A11-A13.

NAR= number of autocorrelation terms. The default is zero unless the previous estimation was AR1.

PARAM/NOPARAM specifies whether the coefficients will be names or constant values. PARAM is the default if there are three or more arguments, or if COEFPREF or VARPREF have been specified.

PRINT/NOPRINT specifies whether the options and output equation will be printed.

RESIDUAL/NORESIDUAL specifies whether the FRML should be unnormalized (have no dependent variable).
**FORM**

**RHOPREF** = a prefix for the autocorrelation coefficient names. The default is RHO plus the equation name, if NAR=1. Otherwise it is PHI, plus the equation name, plus the order of the autoregressive lag.

**SUM/NOSUM** specifies that the equation is to be formed as a sum of terms specified in a list. This is useful when constructing a log likelihood for ML, when you do not know how many terms you will need. See the examples below.

**VALUE/NOVALUE** is used to fix the values of any PARAMs in a list of existing FRMLs. Then they can be printed to show the values with the other variables, or stored in a databank for later use with SIML or SOLVE (without having to worry about storing the associated PARAMs). VALUE is the default if there is only one argument, or if there are several arguments, and they are all existing FRMLs. This option was formerly called CONST.

**VARPR** = a prefix for the coefficient names, to be used in conjunction with the variable names. The default is to use a COEFPR instead. Special coefficient names are used for C (intercept), lags, and leads. 0 (zero) is used in place of C, positive numbers are appended for lags, and L plus positive numbers are appended for leads.

**Examples**

1. This example causes an equation named CONSEQ to be printed and stored.

   ```
   OLSQ CONS C GNP GNP(-1) ;
   FORM (PRINT) CONSEQ ;
   ```

   prints

   ```
   FRML CONSEQ CONS = (123.0 + 0.9*GNP) + 0.05*GNP(-1)
   ```

   (assuming that the coefficient estimates from the OLSQ were (123., .9, .05).)

2. This example saves the same equation as in example 1 with names instead of values.

   ```
   OLSQ CONS C GNP GNP(-1) ;
   FORM (COEFPR=B) CONSEQ ;
   ```

   is equivalent to the following commands:

   ```
   FRML CONSEQ CONS = B0 + B1*GNP + B2*GNP(-1);
   PARAM B0 123 B1 .9 B2 .05 ;
   LIST @RNMSF B0-B2 ;
   ```
3. This example uses a VARPREF instead of a COEFPREF.

\[ \text{OLSQ CONS C GNP GNP(-1);} \]
\[ \text{FORM (VARPREF=B) CONSEQ ;} \]

is equivalent to the following commands:

\[ \text{FRML CONSEQ CONS = B0 + BGNP*GNP + BGNP1*GNP(-1);} \]
\[ \text{PARAM B0 123 BGNP .9 BGNP1 .05 ;} \]
\[ \text{LIST @RNMSF B0 BGNP BGNP1 ;} \]

4. Same as example 3, but the RESID option is used to make the equation unnormalized. This is a form which is most useful for use with the ML command, but could also be used with commands like LSQ, GMM, or FIML.

\[ \text{OLSQ CONS C GNP GNP(-1);} \]
\[ \text{FORM (VARPREF=B,RESID) EC;} \]

is equivalent to the following commands:

\[ \text{FRML EC CONS - ( B0 + BGNP*GNP + BGNP1*GNP(-1) );} \]
\[ \text{LIST @RNMSF B0 BGNP BGNP ;} \]
\[ \text{PARAM @RNMSF ;} \]
\[ \text{UNMAKE @COEF @RNMSF ;} \]

5. This example creates a similar equation and estimates it with LSQ.

\[ \text{FORM CE CONS C GNP GNP(-1); LSQ CE;} \]

is equivalent to: (Note the default coefficient names based on the equation name.)

\[ \text{FRML CE CONS = BCE0 + BCE1*GNP + BCE2*GNP(-1);} \]
\[ \text{PARAM BCE0-BCE2; LSQ CE;} \]

6. This example illustrates including the lagged residual term from AR1.

\[ \text{AR1 IMPT C GNP RELP ;} \]
\[ \text{FORM (PARAM) IE;} \]

creates the equation

\[ \text{FRML IE IMPT = IE0 + IE1*GNP + IE2*RELP + RHOIE*(IMPT(-1) - IE0 - IE1*GNP(-1) - IE2*GNP(-2));} \]
\[ \text{PARAM IE0 -251.8369 IE1 2.7984 IE2 .2636 RHOIE .8328 ;} \]
7. In this example, we estimate the same equation as in example 6, except with AR(4) errors. This FRML could also be used in a system of equations with LSQ or FIML.

FORM (NAR=4) IE IMPT C GNP RELP;
LSQ IE;

8. FORM can be used after a VAR estimation:

VAR Y1 Y2 | C T ;
FORM (VARPR=H) EQ1 EQ2 ;

creates

FRML EQ1 Y1 = HY1_0 + HY1_T*T ;
FRML EQ2 Y2 = HY2_0 + HY2_T*T ;

where the HY1_0, HY1_T, HY2_0, HY2_T parameters are created and set to estimated values from the VAR.

9. Example of the SUM option:

FORM (SUM) EQ S X1-X3 ;

creates

FRML EQ S = X1+X2+X3 ;
FORMAT

FORMAT is an option used with the READ and WRITE commands. It supplies the format for reading and writing data within a TSP program. This section describes how to construct a FORMAT string. See READ and WRITE for a description of where to use it.

**FORMAT= FREE or BINARY or RB4 or RB8 or DATABANK or EXCEL or LABELS or LOTUS or '(format text string)'**

Usage

FORMAT has several alternatives:

1. **FORMAT=FREE** specifies that the numbers are to be read in free format, that is, they are delimited by one or more blanks but may be of varying lengths and mixed formats.

2. **FORMAT=BINARY** specifies that the data are to be read in binary (machine) format, where each variable occupies a single precision floating point word. Binary data may not be mixed with data in other formats, nor can it be moved from one computer type to another. **FORMAT=RB4** (Real Binary 4-byte) is the same thing.

3. **FORMAT=RB8** is double precision binary (8-byte).

4. **FORMAT=DATABANK** specifies that the file being used is a TSP databank. This is an alternative to the IN or OUT/KEEP statements.

5. **FORMAT=EXCEL** reads an Excel spreadsheet file. If the filename ends with .XLS, this is the default.

6. **FORMAT=LABELS** is for WRITE only -- it means that labels like those of the standard PRINT command are to be used.

7. **FORMAT=LOTUS** reads Lotus 123 worksheet files (.WK1, .WKS), with column names at the top and optional dates in the first column. This is the default if the filename includes .WK.

8. **FORMAT=RB4** is the same as **FORMAT=BINARY** (single precision binary).

9. **FORMAT=RB8** is used for double precision binary.
10. **FORMAT=’(format text string)’** specifies a format with which the data will be read. The format text string in TSP is very similar to the format statement in Fortran, since the Fortran format processor is used on it. However, you do not need to know Fortran to construct a simple format string; and consequently, a description of the features you will need is given here.

**Technical note**: Since all data in TSP are floating point, do not use integer or alphabetic formats (unless you're using OPTIONS CHARID). Also, avoid parenthetical groupings in the format unless you are sure you know what they do, because the results can be unpredictable with different operating systems.

A format string starts and ends with a ’ or “, with the parentheses immediately inside these quotes. TSP checks that these parentheses exist and inserts them if they are missing. Note that it is possible to use quotes within the format string – just use double quotes outside the parentheses and single quotes within them. For example: WRITE(FORMAT=’( SE(beta)=’,G12.5)’) SEB; . Alternatives for character strings are WRITE (FORMAT=LABELS), the TITLE command, and the H (Hollerith) format type (if you like to count characters).

Format types within the parentheses describe how long numbers are and how many decimal places they have in the data record. The format types useful to you in a TSP program are usually X, F, E, and G. X specifies columns to be skipped on reading; F the format of floating point numbers, and E the format of floating point numbers in exponential (scientific) notation. G is used for output and specifies the most suitable format (F or E) to be used.

Here is a very simple example using the format (F5.2) to read 5 columns of data:

```
Col: 12345
   10000
```

The number read will be 100.00 since the format specified a 5 digit field with 2 digits after the decimal point.

Here is an example of data for the format (F10.5,5X,E10.3,F8.0):

```
Col: 12345678901234567890123456789012345678901234567890123456789
     343.5  .98765E-01  200
```
The first format specifies a field of length 10 with 5 digits to the right of the decimal point, but since a decimal point was explicitly included in the data, the number will be read as 343.5 from columns 1 to 10. Then 5X specifies that 5 columns (11 to 15) are to be skipped.

E10.3 reads a number in exponential format, which is used when a number is too big or too small for normal notation. Once again, the 10 specifies a field length of 10 columns (16 to 25) and the 3 that there are 3 digits to the left of the decimal point. Since the decimal point is specifically included, the number is read as .98765 times 10^{-1} or .098765.

The final format is F8.0, which specifies a field length of 8 columns (26 to 33) and no digits to the right of the implied decimal point. In this case the number to be input has no decimal point and will be read as 200.

Formats can be combined in many ways: for example, an integer number prefacing a format specification tells how many such fields should be read. Here are several examples:

\((8F10.5)\)

specifies an 80 column record with eight numbers of ten columns each.

\((10X,5E12.6,10X,8F3.1)\)

specifies a 104 column record with 13 variables, 5 in E-format and 8 in F-format. Columns 1 to 10 and 71 to 80 will be skipped when reading.

\((20F5.2/20F5.2)\)

specifies two records per observation (the / means to skip to a new record), each with 20 variables.

This last example demonstrates an important feature of formatted data loading in TSP. In general, one observation will be read or written with each pass through the format statement. That is, the format statement should allow for exactly as many numbers as there are variables to be loaded. Usually one record will be read for each observation which contains all the variables for that observation, but it is possible to have more than one record per observation by use of the slash (/) format descriptor. The records do not necessarily have to have identical formats, although they will usually be of the same length.
FREQ

Examples

FREQ sets the frequency for the series in your TSP run. It may be changed during the course of a TSP run, but series of different frequencies cannot be mixed in the same command (see CONVERT for an exception to this rule). The PANEL options are used to interpret any FREQ N series as panel (time series-cross section) data, that is to tell TSP how to identify one individual from the next. These options are used by any subsequent TSP commands which support panel data, such as PANEL, AR1, and PRINT.

**FREQ NONE or ANNUAL or MONTHLY or QUARTER or WEEKLY;**

or

**FREQ <value>;**

or

**FREQ (PANEL, ID=<ID series>, T=<value>, N=<value>, TIME=<series>, START=<date>) N or A or Q or M or W or <value>;**

Usage

FREQ is very simple: FREQ followed by one of the choices above. Single letter abbreviations are allowed.

The annual, monthly, quarterly, and weekly frequencies imply one, 12, 4, or 52 periods per year respectively. The year is assumed to be base 1900 if it has two digits or base 0 if it has four. You can reset the base using the BASEYEAR= option (see the OPTIONS command entry for details). The format of dates in TSP is always YYYY:PP where YYYY is the year and PP is the period. PP can be any number between 1 and the frequency. The period is suppressed when the frequency is annual. The weekly frequency assumes exactly 52 weeks per year and CONVERT's to quarterly but not to monthly. (TSP does not have a calendar.) If the frequency specified is a number, it represents the number of periods per year.

For convenience in SMPL, quarterly or monthly dates may be specified with decimal points rather than colons. This allows the use of computation and symbolic names for the dates. See the examples below.

The default frequency is none. This frequency is provided for convenience in dealing with non-time series data; the data are assumed to be numbered from observation one, unless you specify the sample otherwise.

Output
The scalar variable @FREQ is stored, with the value of the current frequency. This variable can be used to restore a frequency and sample by a PROC. For example,

COPY @SMPL SMPSAV; COPY @FREQ FRQSAV;

can be used at the start of the PROC, and

FREQ FRQSAV; SMPL SMPSAV;

can be used at the end.

Examples

FREQ A;
SMPL 1890 1920;

specifies data with annual frequency running from 1890 to 1920.

FREQ QUARTER;
SMPL 47:1 82:4;

specifies data with quarterly frequency running from the first quarter of 1947 to the fourth quarter of 1982. The above statements are equivalent to

FREQ QUARTER; SMPL 47.1 82.4;

or

FREQ Q; CONST BEGDATE 47.1 ENDDATE 82.4; SMPL BEGDATE ENDDATE;

FREQ 26;

specifies data with a biweekly frequency.

FREQ (PANEL,T=5);

specifies balanced panel data with 5 observations for each individual, and no particular time series frequency.

FREQ (PANEL,ID=CUSIP,START=1974) A;

specifies possibly unbalanced panel data with an ID series CUSIP that distinguishes each individual, and annual data starting in 1974 for each individual.
Examples

FRML defines equations for TSP. These equations can be used later in the program for estimation, simulation, or they may simply be saved for later computation. The ANALYZ, DIFFER, LSQ, FIML, SIML, ML, EQSUB, and SOLVE procedures all require equations specified by FRML or FORM (PARAM) as input. GENR and SET can also take a FRML name as an argument.

\[
\text{FRML } \langle \text{equation name} \rangle \langle \text{variable name} \rangle = \langle \text{algebraic expression} \rangle ;
\]

or

\[
\text{FRML } \langle \text{equation name} \rangle \langle \text{algebraic expression} \rangle ;
\]

Usage

There are two forms of the FRML statement: the first has the name to be given to the equation, followed by an equation in normalized form, that is, with the name of the dependent variable on the left hand side of the equal sign and an algebraic expression for that variable on the right hand side. The expression must be composed according to the rules given in the Basic Rules section which introduces this manual. These rules are the same wherever an equation is used in TSP: in IF statements, GENR, SET, FRML, IDENT, SMPLIF, SELECT, and GOTO. DOTTed variables can be used in FRMLs.

The second form of the FRML statement is "implicit": there is no equal sign but simply an algebraic expression. This is used for fully simultaneous models, where it might not even be possible to normalize the equations. The ANALYZ, FIML, LSQ and SIML procedures can process implicit equations.

Equations defined by FRML are the same as those defined by IDENT except that the estimation procedures assume that a FRML has an implied additive disturbance tacked on the end, while an IDENT does not. If the FRML is not normalized, it is treated as being equal to the implied disturbance. The distinction is useful only in FIML, where identities may be necessary to complete the Jacobian (to ensure that it is a square matrix).

An equation defined by a FRML statement can contain numbers, parameters, constants, and series. The equation can always be computed at any point by use of GENR (see GENR for the form of the statement). When it is computed, the parameters and constants are supplied with their current values before computation, and the equation is computed for all the values of the series in the current sample.

Examples
These are the equations which are used to estimate the illustrative model by three stage least squares in LSQ:

\[
\begin{align*}
\text{FRML CONSEQ CONS} &= A + B \cdot \text{GNP} ; \\
\text{FRML INVEQ I} &= \text{LAMBDA}(-1) + \text{ALPHA} \cdot \text{GNP} / (\text{DELTA} + \text{R}) ; \\
\text{FRML INTRSTEQ R} &= D + F \cdot (\log(\text{GNP}) + \text{LP} - \text{LM}) ; \\
\text{FRML PRICEQ LP} &= \text{LP}(-1) + \text{PSI} \cdot (\text{LP}(-1) + \text{LP}(-2)) + \text{PHI} \cdot \log(\text{GNP}) + \text{TREND} \cdot \text{TIME} + P0 ;
\end{align*}
\]

In these equations, the dependent variables are CONS, I, R, and LP. The other series are GNP, LM, and TIME. Note the use of lagged series in the equation also. The other variables, A, B, LAMBDA, ALPHA, DELTA, D, F, PSI, PHI, TREND, and P0, are parameters and constants. The first FRML could be written in unnormalized (implicit) form as its residual:

\[
\text{FRML CONSEI CONS} - (A + B \cdot \text{GNP}) ;
\]

When the dependent variable is actually an expression, the unnormalized form is required. The following FRML is invalid:

\[
\text{FRML EQNL LOG(Y) = A + B \cdot X} ;
\]

It should be rewritten as an implicit FRML (for use in \texttt{FIML} or \texttt{SIML}):

\[
\text{FRML EQNL LOG(Y) - (A + B \cdot X)} ;
\]

Here are some more examples; see the \texttt{DIFFER} section also for examples using the normal density and cumulative normal function.

\[
\begin{align*}
\text{FRML ZERO A72-A73} ; \\
\text{FRML TRIGEQ COSX} &= \cos(x) ; \\
\text{FRML TRIGEQ2 COSXY} &= x \cdot y / (x \cdot x + y \cdot y)^{0.5} ; \\
\text{FRML RCONSTR RHO} &= (2/\pi) \cdot \text{atan} (\text{PARAM}) ;
\end{align*}
\]

See the \texttt{DOT} command for an example of defining several similar FRMLs in a DOT loop.
GENR

Options  Examples

GENR computes transformations of one or more series over the current SMPL and stores the result as a new series with the name specified. A previously created TSP equation may also be GENRed and the result stored as a series. TSP equations can be created with FRML, EQSUB, IDENT, FORM or DIFFER.

<new series name> = <algebraic formula> ;
GENR (SILENT, STATIC) <new series name> = <algebraic formula> ;
GENR (SILENT, STATIC) <equation name> [<new series name>] ;

Usage

The first two forms of GENR are the most commonly used: GENR followed by the new variable name, an equal (=) sign, and a formula which should be composed according the rules for TSP equations given in the Basic Rules section of this Help System. Note that the GENR keyword is not required. This formula can involve any of the legal TSP functions and as many variables as desired (subject to overall TSP limits on space).

The third form of GENR is usually used to compute predicted values after a nonlinear estimation (LSQ or FIML). It consists of GENR followed by an equation name and then the name of the variable where you want to place the computed values of the equation. If no such name appears, GENR will put the series in data storage under the name given on the left hand side of the equation. If the equation was implicit (there is no left hand side variable), you must supply a name for GENR to store the results.

If there are any missing values in the input series for GENR or arithmetic errors during computation, missing values will be stored for the affected observations of the output series and warnings will be printed (unless the missing data is part of the argument to a MISS() function). Missing values can be generated directly with the internal names @MVAL, @MISS, @NA or @MV.
If the series appearing on the left hand side of the equation also appears on the right with a lag or lags or leads, it may be updated dynamically as it is computed. A warning message "dynamic GENR" is printed, (TSP Versions prior to 4.1 did not update the right hand side lagged dependent variables dynamically). Use the SILENT option or the SUPRES SMPL; command to suppress this message. This is much more efficient than SET with subscripts. For example, U = RHO*U(-1) + E; creates an AR(1) variable U. PDV = REV + PDV(1)/(1+R); does a reverse dynamic GENR (future observations are evaluated first). The STATIC option can be used to prevent such dynamic evaluation.

Expressions like X(-1) and X(I) are treated as lags/leads if X is a series; otherwise they are treated as subscripts (like @COEF(1)). Expressions like X(76:2) [a date] and M(1,2) [a matrix element] are subscripts (evaluate to scalars).

Output

GENR produces no printed output, except a note when a dynamic GENR is being performed. It stores one new or replacement series in data storage.

Options

SILENT/NOSILENT suppresses the "dynamic GENR" message. It does not suppress error or warning messages.

STATIC/NOSTATIC prevents dynamic evaluation of equations with lagged or led dependent variables.

Examples

\[
\begin{align*}
GNPL1 & = GNP(-1) ; \\
GENR \quad DP & = \text{LOG}(\text{PRICE}/\text{PRICE}(-1)) ; \\
GENR \quad WAVE & = \text{GAMMA} \times \text{SIN}(\text{TREND}) ; \\
DUM & = X > 0; \\
\text{SCLEVEL} & = 1 \times (\text{SC} \leq 6) + 2 \times (\text{SC} > 6 \text{ and } \text{SC} \leq 9) + 3 \times (\text{SC} > 9 \text{ and } \text{SC} \leq 12) + 4 \times (\text{SC} > 12) ;
\end{align*}
\]

This makes a dummy variable equal to one when X is greater than zero (the logical expression has a value of one when true), and equal to zero otherwise.

\[
\text{SCLEVEL} = 1 \times (\text{SC} \leq 6) + 2 \times (\text{SC} > 6 \text{ and } \text{SC} \leq 9) + 3 \times (\text{SC} > 9 \text{ and } \text{SC} \leq 12) + 4 \times (\text{SC} > 12) ;
\]

This makes SCLEVEL equal to 1, 2, 3, or 4, depending on which range the value of SC falls into (this is essentially a recode operation).

\[
\text{GENR CONSEQ CONSFIG} ;
\]
The last two examples compute the series defined by equations; the first computes a fitted consumption from a previously defined (FRML or IDENT) or estimated consumption equation (FORM) and the second generates an identity to define the variable on the left hand side. This ensures that the identity will hold in the data, which is necessary for proper estimation.

SMPL 1,10;
A = 1; B = 1;
SMPL 2,10;
GENR A = A(-1) + 1;
GENR(STATIC) B = B(-1) + 1;

This creates A as a time trend (just like the TREND A; command). B is 1,2,2,2,2,2,2,2.
GMM does General Methods of Moments estimation on a set of orthogonality conditions which are the products of equations and instruments. Initial conditions for estimation are obtained using three-stage least squares. The instrument list may be different for each equation (see the MASK option or the INST option) and the form of the covariance matrix used for weighting the estimator is under user control (the HETERO option for heteroskedastic-consistency and the NMA= option for moving average disturbances).

GMM (FEI, HETERO, ITEROC, ITERU, LSQSTART, COVOC=OWN or <covariance matrix of orthogonality conditions>, COVU=<covariance matrix of residuals>, INST=(<list of instruments>), KERNEL=<spectral density kernel type>, MASK=<matrix of zeros and ones>, NMA=<number of autocorrelation terms>, OPTCOV, nonlinear options) <list of equations> ;

Usage

List the instruments in the INST= option and list the equations after the options; the products of these two are the orthogonality conditions, which are minimized in the metric of an estimate of their expected covariance. This estimate is computed using 3SLS estimates of the parameters, unless the NOLSQSTART option has been specified. If the HETERO and NMA= options are not used, the estimation method coincides with conventional 3SLS estimation. The usual GMM estimator that allows for heteroskedasticity in addition to cross-equation correlation is the default. If you wish to use different instruments for each equation, supply each set in a series of separate lists separated by | to the INST= option (see the examples). Alternatively you can use the MASK= option to drop instruments selectively.

The GMM estimator prints the Sargan or J of overidentifying restrictions if the degrees of freedom are greater than zero (the model is exactly identified if they are equal to zero). If you want to nest overidentifying tests of a series of models, be sure to specify the NOLSQSTART option so that the variance-covariance matrix of the OC's will be held fixed across the tests (otherwise the chi-squared for the difference between two nested models may have the wrong sign).

Method
GMM

See Hansen (1982) for most of the details. If the equations are nonlinear, the iteration method is the usual LSQ method with analytical derivatives (a variant of the method of scoring).

Output

The following results are printed and stored:

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@PHI</td>
<td>scalar</td>
<td>1</td>
<td>E'HH'E, the objective function for instrumental variable estimation.</td>
</tr>
<tr>
<td>@GMMOVID</td>
<td>scalar</td>
<td>1</td>
<td>test of overidentifying restrictions (@PHI*@NOB)</td>
</tr>
<tr>
<td>%GMMOVID</td>
<td>scalar</td>
<td>1</td>
<td>P-value of the above test (using degrees of freedom)</td>
</tr>
<tr>
<td>@NOVID</td>
<td>scalar</td>
<td>1</td>
<td>number of overidentifying restrictions (degrees of freedom)</td>
</tr>
<tr>
<td>@RNMS</td>
<td>list</td>
<td>#params</td>
<td>Parameter names.</td>
</tr>
<tr>
<td>@COEF</td>
<td>vector</td>
<td>#params</td>
<td>Estimated values of parameters, also stored under their names.</td>
</tr>
<tr>
<td>@SES</td>
<td>vector</td>
<td>#params</td>
<td>Standard Errors of estimated parameters.</td>
</tr>
<tr>
<td>@T</td>
<td>vector</td>
<td>#params</td>
<td>T-statistics.</td>
</tr>
<tr>
<td>@SSR</td>
<td>vector</td>
<td>#eqs</td>
<td>Vector of sum of the squared residuals for each of the equations</td>
</tr>
<tr>
<td>@YMEAN</td>
<td>vector</td>
<td>#eqs</td>
<td>Vector of means of the dependent variable for each of the equations</td>
</tr>
<tr>
<td>@SDEV</td>
<td>vector</td>
<td>#eqs</td>
<td>Vector of standard deviations of the dependent variable for each of the equations</td>
</tr>
<tr>
<td>@S</td>
<td>vector</td>
<td>#eqs</td>
<td>Vector of standard errors for each of the equations</td>
</tr>
<tr>
<td>@DW</td>
<td>vector</td>
<td>#eqs</td>
<td>Vector of Durbin-Watson statistics for each of the equations</td>
</tr>
<tr>
<td>@RSQ</td>
<td>vector</td>
<td>#eqs</td>
<td>Vector of R-squared for each of the equations</td>
</tr>
<tr>
<td>@ARSQ</td>
<td>vector</td>
<td>#eqs</td>
<td>Vector of adjusted R-squared for each of the equations</td>
</tr>
<tr>
<td>@OC</td>
<td>vector</td>
<td>#eqs*#inst</td>
<td>Estimated orthogonality conditions</td>
</tr>
<tr>
<td>@COVOC</td>
<td>matrix</td>
<td>#eqs*#inst</td>
<td>Estimated covariance of orthogonality conditions</td>
</tr>
<tr>
<td>@COVU</td>
<td>matrix</td>
<td>#eqs*#eqs</td>
<td>Residual covariance matrix</td>
</tr>
</tbody>
</table>
Inverse square root of \@COVU, the upper triangular weighting matrix

Covariance matrix of the transformed residuals. This is equal to the number of observations times the identity matrix if estimation is by ML

Estimated variance-covariance of estimated parameters

Matrix of residuals = actual - fitted values of the dependent variable

Matrix of fitted values of the dependent variables

Options

\textbf{COVOC=} covariance matrix of the orthogonality conditions. The default is to compute starting values with 3SLS and form the covariance matrix from these.

\textbf{COVOC=OWN} computes residuals from the current starting values and forms the covariance matrix from these.

\textbf{COVU=} covariance matrix of residuals. This is used for the initial 3SLS estimates if the default LSQSTART option is in effect. The default is the identity matrix. This option is the same as the old WNAME= option in \texttt{LSQ}.

\textbf{FEI/NFEI} specifies whether a model with individual fixed effects is to be estimated. \texttt{FREQ (PANEL)} must be in effect for the FEI option.

\textbf{HETERO/NOHETERO} specifies conditional heteroskedasticity of the residuals, and causes the COVOC matrix to include interaction terms of the residuals and the derivatives with respect to the parameters. Specify this option to obtain the usual Hansen or Chamberlain estimator. When HETERO is on, GMM checks that the number of OC's is less than the number of observations so that COVOC will be positive definite (if not, an error message is printed).

\textbf{INST=} (<list of instrumental variables>), assumed to be orthogonal to the residuals of the supplied equations by assumption. In some models these variables are referred to as the "information set." Don’t forget to include C, the constant, unless your model does not require one.

\textbf{INST=} (list1 | list2 | ... ) specifies a list of different instruments for each equation. There must be as many lists as there are equations.
ITEROC/NOITEROC causes iteration on the COVOC matrix. Normally it is left fixed at its initial estimate. If MASK and LSQSTART are used, one iteration is made on the COVOC matrix. This also occurs if NOLSQSTART is used and COVOC= is not specified.

ITERU/NOITERU causes iteration on the COVU matrix. This is the same as the old MAXITW= option in LSQ.

KERNEL=BARTLETT or PARZEN. The spectral density kernel used to insure positive definiteness of the COVOC matrix when NMA > 0. BARTLETT is discussed by Newey and West (1987), while PARZEN is discussed by Gallant (1987). Both are reviewed by Andrews (1991).

LSQSTART/NOLSQSTART specifies if 3SLS should be used to obtain starting values for the parameters and COVOC. NOLSQSTART should be specified if you are restarting iterations with old parameter values and a COVOC matrix. This will be important for testing (see the discussion above).

MASK= a matrix of zeroes and ones which specifies which instruments are to be used for which equations. The matrix is # of instruments by # of equations and the default is a matrix of ones (all instruments used for all equations).

NMA= number of autocorrelation terms (AR and/or MA) to be used in computing COVOC. Some forecasting-type models imply a given NMA value, but other models have no natural choice. See the Andrews reference for automatic bandwidth selection procedures. When there are missing values in the series, NMA does not include terms which cross the gaps in the data. This is useful in panel data estimation.

OPTCOV/NOOPTCOV specifies whether or not the COVOC matrix is optimal. Under the default NOOPTCOV, the @VCOV matrix is computed using the "sandwich" formula of Hansen's 1982 Theorem 3.1 (p.1042). This is appropriate, for example, if the user has supplied a COVOC matrix, but has not scaled it properly. Note: in this (improperly scaled COVOC) case, the @GMMOVID statistic will be invalid. When OPTCOV is in effect, formula (10) of Hansen's Theorem 3.2 (p.1048) is used for the @VCOV matrix. When the user has not supplied a COVOC matrix, the OPTCOV and NOOPTCOV options produce almost exactly the same results. The only difference is due to the difference between the COVOC matrix that was used for iterations, and the COVOC matrix evaluated at the final parameter/residual values. This difference is usually small.

Examples

\textit{GMM (INST=(C,Z1-Z10),NMA=2,HET) EQ1 EQ2;}
To exclude Z2 as instrument for EQ1, Z1 as instrument for EQ2:

\[ \text{READ(NROW=3,NCOL=2) SEL; } \]
\[ \begin{array}{cc}
1 & 1 \\
1 & 0 \\
0 & 1 \\
\end{array} \]

? C,Z1 enter EQ1 and C,Z2 enter EQ2

\[ \text{GMM (INST=(C,Z1,Z2),MASK=SEL) EQ1 EQ2;} \]

The same thing can be done using a different list for each equation:

\[ \text{LIST INST1 C Z1 ; } \]
\[ \text{LIST INST2 C Z2 ; } \]
\[ \text{GMM (INST=(INST1|INST2)) EQ1 EQ2 ; } \]

References


GOTO

**Example**

GOTO provides a method of transferring control within your TSP program, that is, of changing the order in which the statements are executed. However, modern programming theory suggests that the use of GOTO statements should be avoided since they tend to produce unreadable and hard to debug programs. In TSP, the **DO** loop facility and **IF-THEN-ELSE** syntax can be used to avoid the use of GOTO statements.

```
GOTO <statement number> ;
```

or

```
GO TO <statement number> ;
```

**Usage**

A GOTO statement consists of GOTO followed by a statement number which must been defined somewhere in your program. The effect of the statement is to transfer control immediately to that statement. [Note that the GOTO statement cannot be used to transfer to the data section of your program; you can use a series of **LOAD** statements instead.]

**Example**

```
IF A; THEN; GO TO 100 ;
   B = K*K ;
GOTO 200 ;
100 B = K*K*10000 ;
200 OLSQ Y C B ;
```

This example shows the GOTO statement being used to create two branches of the program, one to be executed if A is false, and one to be executed if A is true. The same thing can be done with an **IF-THEN-ELSE** sequence:

```
IF A ; THEN ;
   B = K*K*10000 ;
ELSE ;
   B = K*K ;
OLSQ Y C B ;
```
GRAPH

See also graphics version

Output  Options  Example

GRAPH plots one series against another, using a scale determined by the range of each series. Use the PLOT command to graph series against time.

GRAPH <series for y-axis> <series for x-axis> ;

Usage

Supply the series names in the order y-axis series followed by the x-axis series.

Output

GRAPH produces a one page graph of the second series against the first. The graph is labeled and surrounded by a box. If more than one point is plotted in the same place on the graph, the number of such points will be printed if it is less than or equal to 9. If there are more than 10 to 35 observations at one point, A to Z will be plotted. If there are more than 35 observations at one point, an X will be plotted and the number of such points will be listed after the graph, sorted by the X values. Any observations with missing values will not be plotted (and a warning message will be printed).

The size of the graph is controlled by the line printer width option LIMPRN and the page length option LINLIM. These options are automatically set at the beginning of your TSP run, but they may be changed by using the OPTIONS command. If you have already issued an OPTIONS CRT ; command they will be set at 80 characters wide by 24 characters high.

GRAPH does not store any data in data storage.

Options

Examples

GRAPH TIME COSINE ;
GRAPH Y X ;
GRAPH (graphics version)

**Output**  **Options**  **Examples**

GRAPH plots one series versus another, using a scale determined by the range of each series. The plot may be printed as well as displayed if a hardcopy device such as a Laserjet or dot matrix printer is available. This section describes the graphics version of GRAPH, which is available only for TSP/Oxmetrics, Macintosh TSP, unix, and DOS/Win TSP. For other versions, see the non-graphics GRAPH command.

```
GRAPH (A4, CIRCLE=<radius series>, DASH, DEVICE=<name of printer>, FILE=<name of file>, HEIGHT=<height of letters>, HIRES, LANDSCAP or PORTRAIT, LINE, ORIGIN, PAIR, PREVIEW, SORT, SURFACE, SYMBOL, TITLE='text string to be used as title', WIDTH, XMIN=<x axis minimum>, XMAX=<x axis maximum>, YMIN=<y axis minimum>, YMAX=<y axis maximum>) <x-axis series> <list of y-axis series> ;
```

or

```
<x-series 1> <y-series 1> <x-series 2> <y-series 2> ... ;  (for the PAIR option)
```

or

```
<x series> <y series> <z series> ;  (for the SURFACE option)
```

**Usage**

GRAPH has several forms, depending on the options specified. If GRAPH is followed by a pair of series names, a scatter plot will be produced using the x-axis for the first series and the y-axis for the second. If there are more than two series names and NOPAIR (the default) is specified, the others will also be plotted on the y-axis, and their points will be connected with a line. This makes it easy to plot fitted and actual values from a regression using the default options (see the example below). If the PAIR option is specified, there must be an even number of series names, and each pair will be plotted versus each other, the first series on the x-axis and the second on the y-axis.

TSP/Oxmetrics automatically displays graphs in different windows by default. If there are gaps in the SMPL, observations with missing data, or the FREQ (PANEL) option is set, any lines being graphed will contain breaks.

In the DOS/Win or MAC version of TSP, the graph will be displayed on the screen; if a DEVICE= is specified, a prompt is also displayed which instructs you to type "P" if you wish to print the graph. If you type anything else, the graph will not be printed; this is useful if you decide you do not like its appearance after you have seen the screen.
GRAPH (graphics version)

Output

The graphics version of GRAPH produces a multi-color scatterplot by default. The first series is graphed with points and the remaining series with lines. The LINE option can be used to graph all series with lines. GRAPH differentiates the series using different colors.

Options

General  Oxmetrics only  DOS/Win only  MAC only

DASH/NODASH specifies whether the lines for different series on the screen are to be distinguished by using different dash patterns. The default is no dashes (just color) on the screen and dashes on printed output. There are 7 dash patterns.

LINE/NOLINE specifies whether the first series should be plotted with a line rather than as a scatter of points. The default is to use dots to represent the series.

ORIGIN/NOORIGIN causes a horizontal line to be drawn starting at zero on the vertical axis.

PAIR/NOPAIR specifies whether all the series except the first are to be plotted on the y-axis (the default) or whether the series are to be used in pairs of x versus y. Since GRAPH always draws a line for every pair of series except the first, it implicitly assumes that the x-axis series will be sorted by the user before the procedure is executed.

PREVIEW/NOPREVIE specifies whether the graph is to be shown on the screen before printing or saving. The default is PREVIEW for interactive use and NOPREVIE for batch use.

SORT/NOSORT controls the ordering of the data on the X-axis so that a line connecting the points will not cross itself.

SYMBOL/NOSYMBOL specifies that symbols are to be used for plotting. When the LINE option is on, NOSYMBOL is the default.

TITLE='a string which will be printed across the top of the graph'.

XMAX= maximum value for the x-axis. This value must be greater than or equal to the maximum value of the x series.

XMIN= minimum value for the x-axis. This value must be less than or equal to the minimum value of the x series.
GRAPH (graphics version)

YMAX= maximum value for the y-axis. This value must be greater than or equal to the maximum value of the y series.

YMIN= minimum value for the y-axis. This value must be less than or equal to the minimum value of the y series.

For convenience, the DEVICE=, FILE=, and HEIGHT= options of GRAPH are retained for the next PLOT(s) or GRAPH(s) until they are overridden explicitly. These options may also be set in a LOGIN.TSP file with a GRAPH statement which does not specify any series to graph.

Oxmetrics only

CIRCLE= radius series. This places a circle symbol at each point, with area based on the radius. Values in the range 100 to 1000 are visible and not too large. If you have a weight proportional to area, use its square root and then scale it to the range 100 to 1000.

SURFACE/NOSURFACE specifies that a three-dimensional surface plot is to be created. This option requires 3 variables as arguments.

DOS/Win only

A4/NOA4 specifies A4 paper size. Available for DEVICE=LJ3 or POSTSCRIPT only.

DEVICE= CHAR or EPSON or LJ2 or LJ3 or LJET or LJP75 or LJR100 or LJR150 or LJR300 or POSTSCRI or PS specifies the hardcopy device to be used for printer output. LJ means HP LaserJet or compatible, EPSON is EPSON dot matrix or compatible, POSTSCRI and PS are Postscript output, and CHAR is the old line printer output (characters instead of graphics). The LJ suffixes specify models of the printer and the LJR suffixes specify the resolution of the LaserJet printer directly, rather than giving the printer type. The maximum resolutions for the LJET, LJP75, and LJ2 printers are 100, 150, and 300 respectively. Note that default larger resolutions imply larger file sizes and printing times.

FILE= the name of a file to which the graphics image is to be written. This file can be printed later. For example, if you are running under DOS and your printer device is LPT1, use the command:

copy/b file LPT1;

HEIGHT= letter height in inches. The default* is .25. Values in the range (0,1] are valid.
HIRES/NOHIRES controls how graphs are printed in batch mode (when PREVIEW is not being used). Normally (NOHIRES), graphs are printed in character mode to the batch output file. When the HIRES option is used, the patched DEVICE= and FILE= will be used; usually this will send a page to LPT1 for each graph.

LANDSCAP/PORTRAIT specifies the orientation of the plot. On the Mac, specify this option in the dialog box.

SURFACE/NOSURFACE specifies that a 3-dimensional surface is to be plotted (TSP/Oxmetrics only). There must be three arguments (x, y, z axis variables). Once you see the plot, you can use ALT>, ALT<, ALT+, and ALT- to rotate the view.

MAC only

WIDTH/NOWIDTH specifies whether varying width sizes are to be used to distinguish the lines corresponding to different series on the graph.

Examples

This graph displays Student's t distribution for 2, 4, and 10 degrees of freedom, a Cauchy distribution, and the Laplace and Normal distributions (chosen to have the same variance as the others):

GRAPH(DEVICE=LJ3,LINE,XMIN=-5,XMAX=5,TITLE='Some Fat-Tailed Distributions') T T2 T4 T10 CAUCHY LAPLACE NORMAL;

The following example plots (and stores on disk) the graph shown in the user's guide:

SMPL 1,1000;
RANDOM (SEEDIN=49813) X E;
Y = 1 + X + E;
OLSQ Y C X;
GRAPH(DEV=LJ3,PREVIEW,FILE='GRAPH.PLT',TITLE='Actual and Fitted Values') X Y @FIT;
HELP

HELP

Examples

HELP provides basic syntax information on commands, and it will also list command by various functional groups. This is useful for checking the syntax on an unfamiliar command or for checking related commands, but it does not replace the manuals (or the HELP system, if you are using the Windows version). The syntax summary may also be printed if you supply the wrong number of arguments or an unrecognized command option.

HELP;
or HELP COMMANDS;
or HELP <command name>;
or HELP FUNCTION;
or HELP NONLINEAR;
or HELP ALL;
or HELP GROUP;
or HELP <group number>;

Usage

HELP by itself lists the various ways of using the HELP command, which are given below:

HELP COMMANDS lists all the TSP commands, ten per line.
HELP command name gives details on a particular command.
HELP FUNCTION lists functions and operators (both general and matrix).
HELP NONLINEAR lists the options for nonlinear estimation procedures.
HELP ALL gives a one-line description of each TSP command, from A to Z.
HELP GROUP gives the same description, but sorted by functional groups.
HELP group number gives the same description for a single group. The group numbers can be obtained using the HELP command with no arguments.

Examples
HELP AR1;
summarizes the arguments and options of the AR1 command.

HELP 1;
gives a one-line description of each command in the linear estimation group.
HIST

See also graphics version

Output  Options  Examples

HIST produces histograms (bar charts or frequency distributions) of series. It is convenient for obtaining a rough picture of the univariate distribution of your data.

\[ \text{HIST (BOT, DISCRETE, MAX=\text{<maximum for x-axis>}, MIN=\text{<minimum for x-axis>}, NBINS=\text{<number of bins>}, PERCENT, PRINT, WIDTH=\text{<width of bin>}) <list of series>;} \]

Usage

Follow HIST with the names of one or more series for which you would like to see a frequency distribution. The default options for output yield a histogram with ten equally spaced bins or cells running from the minimum value of the series to the maximum value. The bars for each cell have a width of two lines on the printed page, and are based on the left hand axis of the graph.

Output

If the PRINT option is on, a plot of the histogram for each series is produced.

The following is stored in data storage:

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@HIST</td>
<td>matrix</td>
<td>#bins*#series</td>
<td>matrix with observation counts for each series</td>
</tr>
<tr>
<td>@HISTVAL</td>
<td>matrix</td>
<td>#bins*#series</td>
<td>matrix with bin lower bounds for each histogram</td>
</tr>
</tbody>
</table>

Options

BOT/NOBOT causes the printed histogram to be based on the left side of the page. The default is the middle of the page.

DISCRETE/NODISCRE specifies whether the series are discrete or continuous. If the series are discrete, there will be one cell for each unique value (limited by NBINS).

MAX= upper bound on the last cell. The default is the maximum value of the series.
**MIN**= lower bound on the first cell. The default is the minimum value of the series.

**NBINS**= the number of bins or cells (The default is 10 for NODISCRETE and 20 for DISCRETE).

**PERCENT**/**NOPERCENT** causes the percent in each cell rather than the absolute number to be printed (the graph looks the same, but the labels on the horizontal axis are different).

**PRINT**/**NOPRINT** tells whether the histogram is to be printed or just stored.

**WIDTH**= the width of the bars (the default is 2 printer lines).

**Examples**

**HIST X ;**

produces a plot with the vertical axis containing ten cells running from the minimum value of X to the maximum value of X, and the horizontal axis showing the number of observations of X which take on values within each of the cells.

**HIST (MAX=100,MIN=0,NBINS=40,PERCENT) Y1 Y2 ;**

produces two histograms, each with 40 cells which have a width equal to 2.5. The percent of observations of Y1 (or Y2) which fall in each cell are shown.

Suppose the variable REASON takes on the values 0,1,2, and 3. The command

**HIST (DISCRETE) REASON ;**

will produce a histogram with four cells, containing the number of observations taking on each one of the four values of REASON.
HIST (graphics version)

HIST (graphics version)

Output  Options  Examples

HIST produces histograms (bar charts or frequency distributions) of series. It is convenient for obtaining a rough picture of the univariate distribution of your data. The graphics version (options NOPRINT, PREVIEW) is the default for TSP/Oxmetrics.

\[
\text{HIST (CDF, DENSITY, DISCRETE, HIST, MAX=<maximum X value>, MIN=<minimum X value>, NBINS=<number of bins>, NORMAL, PRINT, PREVIEW, STANDARD, TITLE=<text string>) <list of series> ;}
\]

Usage

Follow HIST with the names of one or more series for which you would like to see a frequency distribution. The default options for output yield a histogram with ten equally spaced bins or cells running from the minimum value of the series to the maximum value.

Output

Plots of the histogram for each series are produced, each in a separate window.

The following is stored in data storage:

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@HIST</td>
<td>matrix</td>
<td>nbins*#series</td>
<td>matrix with observation counts for each series</td>
</tr>
<tr>
<td>@HISTVAL</td>
<td>matrix</td>
<td>nbins*#series</td>
<td>matrix with bin lower bounds for each histogram</td>
</tr>
</tbody>
</table>

Options

CDF/NOCDF includes a normal QQ plot below the histogram.

DENSITY/NODENSITY superimposes a smooth density on the histogram.

DISCRETE/NODISCRE specifies whether the series are discrete or continuous. If the series are discrete, there will be one cell for each unique value (limited by NBINS). Cells with zero counts will be omitted.

HIST/NOHIST specifies whether to include a bar type histogram in the printout.
HIST (graphics version)

**MAX=** upper bound on the last cell. The default is the maximum value of the series.

**MIN=** lower bound on the first cell. The default is the minimum value of the series.

**NBINS=** the number of bins or cells (The default is 10 for NODISCRETE and 20 for DISCRETE).

**NORMAL/NONORMAL** superimposes a normal density on the histogram.

**PRINT/NOPRINT** tells whether the histogram is to be printed or just stored.

**STANDARD/NOSTANDARD** standardizes the data before plotting.

**TITLE=** 'title string' labels the plot.

**Examples**

**HIST X ;**

produces a plot with the vertical axis containing ten cells running from the minimum value of X to the maximum value of X, and the horizontal axis showing the number of observations of X which take on values within each of the cells.

**HIST (MAX=100,MIN=0,NBINS=40) Y1 Y2 ;**

produces two histograms, each with 40 cells and a width equal to 2.5. The fraction of observations of Y1 (or Y2) which fall in each cell are shown.

Suppose the variable REASON takes on the values 0,1,2, and 3, with the following counts:

<table>
<thead>
<tr>
<th>Value</th>
<th>Number of observations</th>
</tr>
</thead>
<tbody>
<tr>
<td>0</td>
<td>10</td>
</tr>
<tr>
<td>1</td>
<td>0</td>
</tr>
<tr>
<td>2</td>
<td>34</td>
</tr>
<tr>
<td>3</td>
<td>42</td>
</tr>
</tbody>
</table>

The command

**HIST (DISCRETE) REASON ;**

will produce a histogram with three cells, containing the number of observations taking on the values of REASON = 0, 2, 3.
On the other hand, the command

\textit{HIST REASON ;}

will default to the INTEGER mode and produce a histogram with four cells, containing the number of observations taking on the four values of REASON.

If SIZE is a variable containing the log sales or employment of a cross section of firms,

\textit{HIST (DENSITY,TITLE="Size distribution")}

produces a graph of the size distribution for the firms with a smooth approximation to the density superimposed.
IDENT

Example

IDENT defines identities for TSP. These identities can be used to complete simultaneous equations model for full information maximum likelihood estimation with FIML or for simulation with SIML or SOLVE.

**IDENT** `<equation name> <variable name>=<algebraic expression> ;`

or

**IDENT** `<equation name> <algebraic expression> ;`

Usage

There are two forms of the IDENT statement: the first has the name to be given to the equation, followed by an equation in normalized form, that is, with the name of the dependent variable on the left hand side of the equal sign and an algebraic expression for that variable on the right hand side. The expression must be composed according to the rules given in the Basic Rules section in this manual. These rules are the same wherever an equation is used in TSP: in IF statements, GENR, SET, FRML, and IDENT.

The second form of the IDENT statement is "implicit": there is no equal sign but simply an algebraic expression. This is used for fully simultaneous models, where it might not be possible to normalize the equations. The FIML and SIML procedures can process implicit equations, although the SOLVE procedure cannot.

Equations defined by IDENT are the same as those defined by FRML except that the estimation procedures assume that a FRML has an implied additive disturbance tacked on the end, while an IDENT does not. The distinction is useful only in FIML, where identities may be necessary to complete (square) the Jacobian.

An equation defined by a IDENT statement can contain numbers, parameters, constants, and series. The equation can always be computed at any point by use of GENR (see GENR for the form of the statement). When it is computed, the parameters and constants are supplied with their current values before computation, and the identity is computed for all the values of the series in the current sample.

Output

IDENT produces no output. A single equation is stored in data storage.

Example
Here is the identity that completes the five equation illustrative model in the User's Guide:

\[ \text{IDENT GNPID GNP} = \text{CONS} + I + G \]

This identity states that GNP (Gross National Product) is always equal to the sum of CONS (consumption), I (investment), and G (government expenditures).
IF

Examples

IF provides conditional execution of commands in TSP. It requires a subsequent THEN statement, and may also be followed by an ELSE statement if you wish to have a branch for a false result.

IF <scalar expression> ;

Usage

The result of the scalar expression following IF is interpreted as true if it is larger than zero and false otherwise. If the result of the expression is a logical value itself (for example, TEST < 2.0) it will be given the value one for true and zero for false by TSP. The expression must be formulated using TSP's Basic Rules for formula construction.

Only if the expression gives a true result will the statement following the next THEN ; statement be executed. If that statement is a DO statement, all the statements up to the closing ENDDO ; statement will be executed. If there is an ELSE ; statement subsequently, the statements following it will be executed in the same manner if the result of the IF was false.
IN (Databank)

Examples

IN specifies a list of external databank files to be searched automatically for variables not found in TSP's working data storage.

\[
\text{IN <list of filenames> ;}
\]

or

\[
\text{IN 'filename strings'};
\]

Usage

Follow the word IN with the names of the TSP databanks to be searched for your variables. Most systems use binary .TLB files for databanks.

After the IN statement appears in your program, TSP searches each of the files in the order in which you specified for any variables which are not already loaded. The IN statement remains in effect until another IN statement is encountered. If you wish to stop searching any files, include an IN statement with no arguments to cancel the previous statement. Up to 8 IN databanks can be active at one time.

IN sets the FREQ and SMPL from the first series in a databank if no SMPL is present.

Examples

Suppose you have created a TSP databank called TSPDATA.TLB on disk with the members GNP, CONS, etc. using the OUT statement.

Then you can access this databank as shown below. It is important that the frequency specified match the frequency of the series you want from the databank or a warning will be issued. The sample does not necessarily have to be the same since TSP will remove observations or fill in missing values as appropriate.

\[
\text{FREQ A ; SMPL 46 75 ;}
\text{IN TSPDATA ;}
\text{GENR CONSL1 = CONS(-1) ;}
\text{OLSQ CONS C GNP CONSL1 ;}
\]

Other examples of legal IN statements:

\[
\text{IN BANK1 BANK2 BANK3 ;
IN ; ? cancel the previous databanks.}
\text{IN USDATA UKDATA DLDATA SWDATA ;}
\]
INPUT

Example

INPUT reads a stream of TSP commands from an external (disk) file and executes them as a unit.

INPUT [filename] ;
or
INPUT 'filename string' ;

Usage

With INPUT you can use a file of TSP commands you have already written. The input file does not have to be a complete TSP program; you may use disk files as a convenient way to store frequently used pieces of your programs, such as user defined procedures (PROCs), or tables of data to load.

INPUT takes one filename only as an argument. If it is not enclosed in quotes, the filename must conform to restrictions placed on TSP variable names: it must be limited to eight characters and the filename extension must be omitted (.tsp will be assumed). With quotes, the filename can include directory information and extensions and can be up to 128 characters long.

In interactive mode, if the filename is absent, you will be prompted for it. In this case you may also specify a directory as well as an extension or disk unit, but the whole name must be 128 characters or less. Again, if the extension is omitted, .tsp will be assumed.

You will also be prompted to have the file's output displayed on screen as it executes, or to have it sent to another file. Either way, the commands read and executed become part of your TSP session, and may be REVIEWed, EDITed, EXECed, etc.... If you send the output to a disk file, you will be prompted for an output filename. This name may also be up to 128 characters long. If no extension is given, .out will be assumed. If you do not provide a filename, it will default to the same name as the input file except with the .out extension. If another file already has this name, it will be overwritten, unless your system allows multiple versions of the same file.

If you want the command stream to be read and stored but NOT executed, use an EXIT command at the end of the input file instead of END. This is the same as using the EXIT command in collect mode. You may then REVIEW the commands read, and EXEC selected sections.
INPUT commands can be nested (they may appear in files used for input). login.tsp is a special INPUT file; it is read automatically at the start of interactive sessions and batch jobs (this is useful for setting default options).

Examples

1? INPUT ILLUS
Do you want the output printed at the terminal (y/n)? [y]N
Enter name of TSP output file:

This example reads the illustrative example from ILLUS.TSP in the current directory, and places the output in ILLUS.OUT. When execution is finished, the prompt will reappear on screen; the exact line number will depend on how many lines were read from the INPUT file. REVIEW would display everything that had been read.
INST obtains single equation instrumental variable estimates. INST is a synonym for 2SLS. By choosing an appropriate list of instrumental variables, INST will obtain conventional two stage least squares estimates. Options allow you to obtain weighted estimates to correct for heteroskedasticity, or to obtain standard errors which are robust in the presence of heteroskedasticity of the disturbances.

**Usage**

In the basic INST statement, list the dependent variable first and then the independent variables which are in the equation. Include an option INST containing a list of variables to be included as instruments in parentheses. The list of instruments must include any exogenous variables in the equation, in particular the constant, \( c \), as well as any additional instruments you may wish to specify. There must be at least as many instrumental variables as there are independent variables in the equation to meet the rank condition for identification. Any observations with missing values will be dropped from the sample.

Two stage least squares is INST with all the exogenous variables in the complete model listed as instruments and no other variables. Valid estimation can be based on fewer instruments when a complete model involves a large number of exogenous variables, or estimates can be made even when the rest of a simultaneous model is not fully specified. In these cases, the estimator is instrumental variables, but not really two stage least squares.

If there are exactly as many instruments as independent variables specified, the resulting estimator is classic instrumental variables, that is,

\[
b = (Z'X)^{-1}Z'y
\]

where \( Z \) is the matrix of instruments, \( X \) the matrix of independent variables, and \( y \) the dependent variable. (For the more general case, see the formulas below).
Instrumental variable estimation can also be done using the AR1 procedure (for models with first order serial correlation) and the LSQ procedure (for nonlinear and multi-equation models). In these cases, include the list of instruments in the INST option. See those commands for further information.

The FEI options specifies that a model including individual fixed effects is to be estimated. FREQ (PANEL) must be in effect when using this option. The estimates are computed by removing individual means from all the variables, which implies that the effects are treated as exogenous variables. The WEIGHT option is not available with FEI.

The list of independent variables on the INST command may include PDL variables, however, you are responsible for seeing that there are enough instruments for these variables after the constraints implied by PDL are imposed. If the PDL variable is exogenous, the most complete list would include all the lags of the variable over which the PDL is defined. These variables may be highly collinear, but will cause no problems due to TSP's use of the generalized inverse when computing regressions - a subset of the variables which contains all the information will be used. If the PDL variable is endogenous, you must include enough instruments to satisfy the order condition for identification. The number required can be computed as

\[ \#\text{inst} = \#\text{lags} \less (\text{order of polynomial}) \less (\text{number of endpoint constraints}) \]

Method

Let \( y \) be the dependent variable, \( X \) be the \((T \times k)\) matrix of independent variables, and \( Z \) be the \((T \times m)\) matrix of instrumental variables (the included and excluded exogenous variables for two stage least squares). Then the formulas used to compute the coefficients, their standard errors, and the objective function are the following:

\[
P_z = Z(Z'Z)^{-1}Z' \\
b = (X'P_zX)^{-1}(X'P_zy) \\
= [X'Z(Z'Z)^{-1}Z'X]^{-1}X'Z(Z'Z)^{-1}Z'y \\
e = y - Xb \\
s^2 = e'e/(T-k) \\
V(b) = s^2[X'P_zX]^{-1} \\
\hat{PHI} = e'P_z e
\]
The structural residuals $e$ are used to compute all the usual goodness-of-fit statistics.

**Output**

The output of INST begins with an equation title, the name of the dependent variable and the list of instruments. This is followed by statistics on goodness-of-fit: the sum of squared residuals, the standard error of the regression, the R-squared, the Durbin-Watson statistic for autocorrelation of the residuals, and an F-statistic for the hypothesis that all coefficients in the regression except the constant are zero. The objective function $e'P(Z)e$ (stored as @PHI) is the length of the residual vector projected onto the space of the instruments. This is analogous to the sum of squared residuals in OLSQ -- it can be used to construct a pseudo-F test of nested models. Note that it is zero for exactly identified models (if they have full rank). A test of overidentifying restrictions (@FOVERID) is also printed when the number of instruments is greater than the number of right hand side variables. It is given by $\frac{\@PHI}{(\@S2^{(m-k)})}$

All the above statistics are based on the "structural" residuals, that is residuals computed with the actual values of the right hand side endogenous variables in the model rather than the "fitted" values from a hypothetical first stage regression.

Following this is a table of right hand side variable names, estimated coefficients, standard errors and associated t-statistics. If the variance-covariance matrix has not been suppressed (see the SUPRES command), it is printed after this table. Finally, if the RESID and PLOTS options are on, a table and plot of the actual and fitted values of the dependent variable and the residuals is printed.

INST also stores most of these results in data storage for later use. The table below lists the results available after an INST command. The fitted values and residuals will only be stored if the RESID option is on (the default).

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@LHV</td>
<td>list</td>
<td>1</td>
<td>Name of the dependent variable</td>
</tr>
<tr>
<td>@RNMS</td>
<td>list</td>
<td>#vars</td>
<td>Names of right hand side variables</td>
</tr>
<tr>
<td>@SSR</td>
<td>scalar</td>
<td>1</td>
<td>Sum of squared residuals</td>
</tr>
<tr>
<td>@S</td>
<td>scalar</td>
<td>1</td>
<td>Standard error of the regression</td>
</tr>
<tr>
<td>@S2</td>
<td>scalar</td>
<td>1</td>
<td>Standard error squared</td>
</tr>
<tr>
<td>@YMEAN</td>
<td>scalar</td>
<td>1</td>
<td>Mean of the dependent variable</td>
</tr>
<tr>
<td>@SDEV</td>
<td>scalar</td>
<td>1</td>
<td>Standard deviation of the dependent variable</td>
</tr>
<tr>
<td>@NOB</td>
<td>scalar</td>
<td>1</td>
<td>Number of observations</td>
</tr>
</tbody>
</table>
@DW scalar 1 Durbin-Watson statistic
@RSQ scalar 1 R-squared
@ARSQ scalar 1 Adjusted R-squared
@FST scalar 1 pseudo F-statistic for zero slopes
@FOVERID scalar 1 test of overidentifying restrictions when #inst>@vars
@PHI scalar 1 The objective function \( e'P(Z)e \)
@COEF vector #vars Coefficient estimates
@SES vector #vars Standard errors
@T vector #vars T-statistics
@VCOV matrix #vars*#vars Variance-covariance of estimated coefficients
@RES series #obs Residuals = actual - fitted values of the dependent variable
@FIT series #obs Fitted values of the dependent variable
@AI series #obs estimated fixed effects stored as a series (for FEI)
@COEFAI vector #individuals estimated fixed effects (for FEI)
@SESAI vector #individuals standard errors of fixed effects (for FEI)
@TAI vector #individuals T-statistics for fixed effects (FEI)
%TAI vector #individuals p-values for T-statistics of fixed effects (FEI)

If the regression includes a PDL variable, the following will also be stored:

@SLAG scalar 1 Sum of the lag coefficients
@MLAG scalar 1 Mean lag coefficient (number of time periods)
@LAGF vector #lags Estimated lag coefficients, after "unscrambling"

**REGOPT (NOPRINT) LAGF;**

will turn off the lag plot for PDL variables.

**Options**

FEI/NOFEI specifies that a model with individual-specific effects is to be computed. FREQ(PANEL) must be in effect.

FEPRINT/NOFEPRINT specifies that the fixed effect estimates are to be printed as well as stored.
**HCOMEGA** = BLOCK or **DIAGONAL** specifies the form of the $\Omega = E[uu']$ matrix to use when computing ROBUST standard errors. Ordinarily, the default is diagonal, which yields the usual robust standard errors. When **FREQ** (PANEL) is in effect, the default is BLOCK, which allows for cross-time correlation of the disturbances within individuals. This feature can be used for any kind of grouped data, simply by ensuring that the relevant PANEL setup has been defined.

**INST**=(list of instrumental variables) or the name of a list containing instrumental variables.

**NORM/UNNORM** tells whether the weights are to be normalized so that they sum to the number of observations. This has no effect on the coefficient estimates and most of the statistics, but it makes the magnitude of the unweighted and weighted data the same on average, which may help in interpreting the results. The NORM option has no effect if the WEIGHT option has not been specified.

**ROBUSTSE/NOROBUST** causes the variance of the coefficient estimates, the standard errors, and associated t-statistics to be computed using the formulas suggested by White, among others. These estimates of the variance are consistent even when the disturbances are not homoskedastic (although they must be independent), and when their variances are correlated with the independent variables in the model. See the references for the exact formulas. When FEI is specified with ROBUST, the standard errors for the fixed effects will still be conventional estimates.

**SILENT/NOSILENT** suppresses all output. The results are still stored.

**TERSE/NOTERSE** suppresses printing of everything but the $e'P(Z)e$ objective function and the table of coefficients.

**WEIGHT**= the name of a series which will be used to weight the observations. The data and the instruments are multiplied by the square roots of the weighting series before the regression is computed, so that the weighting series should be proportional to the inverses of the variances of the disturbances. If the weight is zero for a particular observation, that observation is not included in the computations nor is it counted in determining degrees of freedom. This option is not available with FEI.

**Examples**

This example estimates the consumption function for the illustrative model, using the constant, trend, government expenditures (G), and the log of the money supply (LM) as instruments:

```plaintext
INST INST=(C,G,TIME,LM) CONS,C,GNP;
```
Using population as weights, the following example regresses the fraction of young people living alone on various other demographic characteristics across states. Population is proportional to the inverse of the variance of per capita figures.

\[ \text{INST (WEIGHT=POP INST=(C, URBAN, CATHOLIC, SERVEMP, SOUTH) YOUNG, C, RSALE, URBAN, CATHOLIC ;} \]

Other examples of the INST/2SLS command:

\[ \text{INST (ROBUSTSE, INST= (C LOGR LOGR(-1) LOGR(-2) LOGR(-3))) LOGP C LOGP(-1) LOGR ;} \]
\[ 2SLS(INST=(C,LM(-1)-LM(-3))) TBILL C RATE(4,12,FAR) ;} \]

Note that the constant (C) must always be named explicitly as an instrument if it is needed.

References


INTERVAL estimates a model like the linear Ordered Probit model, but where the limits are known. Unlike Ordered Probit, the limits may be different for different observations. INTERVAL is also similar to two-limit Tobit, with the difference that when the dependent variable is between the upper and lower bounds, only that fact is observed and not its actual value. INTERVAL is useful when the dependent variable is in a known range, but the actual value has been censored for confidentiality reasons.

\[
\text{INTERVAL (LOWER=<lowerlimit>,UPPER=<upperlimit>, nonlinear options) <dependent variable> <list of independent variables> ;}
\]

Usage

The basic INTERVAL statement is like the OLSQ statement: first list the dependent variable and then the independent variables. If you wish to have an intercept term in the regression (usually recommended), include the special variable C or CONSTANT in your list of independent variables. You may have as many independent variables as you like subject to the overall limits on the number of arguments per statement and the amount of working space, as well as the number of data observations you have available.

The LOWER= and UPPER= options are required. Normally these will be series with the lower and upper limits for each observation. The dependent variable should be coded so that it lies between its two bounds. For example, if category=3 means that the dependent variable (Y) is between 10 and 20, then Y should be coded so that \( 10 < Y < 20 \). The lower and upper limits for this observation will take the values 10 and 20. See the examples below.

The observations over which the regression is computed are determined by the current sample. If any of the observations have missing values within the current sample, INTERVAL will print a warning message and will drop those observations.

The list of independent variables on the INTERVAL command may include variables with explicit lags and leads as well as PDL (Polynomial Distributed Lag) variables. These distributed lag variables are a way to reduce the number of free coefficients when entering a large number of lagged variables in a regression by imposing smoothness on the coefficients. See the PDL section for a description of how to specify such variables.

Output
INTERVAL

The output of INTERVAL begins with an equation title and the usual starting values and diagnostic output from the iterations. Final convergence status is printed. After convergence, the number of observations, the value of the log likelihood, and the Schwarz-Bayes information criterion are printed. This is followed by observation counts for lower, upper, and double bounded observations, and the usual table of right hand side variable names, estimated coefficients, standard errors and associated t-statistics.

INTERVAL also stores some of these results in data storage for later use. The table below lists the results available after an INTERVAL command.

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@LHV</td>
<td>list</td>
<td>1</td>
<td>Name of dependent variable</td>
</tr>
<tr>
<td>@RNMS</td>
<td>list</td>
<td>#vars</td>
<td>List of names of right hand side variables</td>
</tr>
<tr>
<td>@IFCONV</td>
<td>scalar</td>
<td>1</td>
<td>=1 if convergence achieved, 0 otherwise</td>
</tr>
<tr>
<td>@YMEAN</td>
<td>scalar</td>
<td>1</td>
<td>Mean of the dependent variable</td>
</tr>
<tr>
<td>@NOB</td>
<td>scalar</td>
<td>1</td>
<td>Number of observations</td>
</tr>
<tr>
<td>@LOGL</td>
<td>scalar</td>
<td>1</td>
<td>Log of likelihood function</td>
</tr>
<tr>
<td>@AIC</td>
<td>scalar</td>
<td>1</td>
<td>Akaike information criterion</td>
</tr>
<tr>
<td>@SBIC</td>
<td>scalar</td>
<td>1</td>
<td>Schwarz Bayesian information criterion</td>
</tr>
<tr>
<td>@NCOEF</td>
<td>scalar</td>
<td>1</td>
<td>Number of independent variables (#vars)</td>
</tr>
<tr>
<td>@NCID</td>
<td>scalar</td>
<td>1</td>
<td>Number of identified coefficients</td>
</tr>
<tr>
<td>@COEF</td>
<td>vector</td>
<td>#vars</td>
<td>Coefficient estimates</td>
</tr>
<tr>
<td>@SES</td>
<td>vector</td>
<td>#vars</td>
<td>Standard errors</td>
</tr>
<tr>
<td>@T</td>
<td>vector</td>
<td>#vars</td>
<td>T-statistics</td>
</tr>
<tr>
<td>%T</td>
<td>vector</td>
<td>#vars</td>
<td>p-values for T-statistics</td>
</tr>
<tr>
<td>@GRAD</td>
<td>vector</td>
<td>#vars</td>
<td>Gradient of log likelihood at convergence</td>
</tr>
<tr>
<td>@VCOV</td>
<td>matrix</td>
<td>#vars*</td>
<td>Variance-covariance of estimated coefficients</td>
</tr>
<tr>
<td>@FIT</td>
<td>series</td>
<td>#obs</td>
<td>Fitted values of dependent variable</td>
</tr>
</tbody>
</table>

If the regression includes a PDL or SDL variable, the following will also be stored:

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@SLAG</td>
<td>scalar</td>
<td>1</td>
<td>Sum of the lag coefficients</td>
</tr>
<tr>
<td>@MLAG</td>
<td>scalar</td>
<td>1</td>
<td>Mean lag coefficient (number of time periods)</td>
</tr>
<tr>
<td>@LAGF</td>
<td>vector</td>
<td>#lags</td>
<td>Estimated lag coefficients, after &quot;unscrambling&quot;</td>
</tr>
</tbody>
</table>

Method

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Like the binary and ordered Probit models, the Interval model is based on an unobserved continuous dependent variable ($y^*$). The model is

$$y^* = XB + e.$$  

Instead of $y^*$, we observe a category value $Y$, which implies that $y^*$ lies between known limits, where the limits may include minus or plus infinity. In the usual application the set of possible limits are the same for all observations but this is not necessary. The underlying model is the same as that used for Ordered Probit (that is, $e$ is assumed to be normally distributed), but with known limits.

For example, suppose there are 3 categories, with category values 0, 1, and 2, where the first and the last are open-ended. The model is

$$Y = 0 \text{ if } MU0 <= XB + e < MU1 \text{ (MU0 = } -\text{infinity, MU1 a known value)}$$

$$Y = 1 \text{ if } MU1 <= XB + e < MU2 \text{ (MU2 a known value)}$$

$$Y = 2 \text{ if } MU2 <= XB + e < MU3 \text{ (Note: MU3 = infinity)}$$

The terms in the likelihood function for observations with each of the values 0, 1, or 2 are the following:

$$\log L(Y = 0) = \log [\Phi(\mu_0 - X\beta)]$$

$$\log L(Y = 1) = \log \left[ \int_{\mu_0 - X\beta}^{\mu_1 - X\beta} \phi(u) \, du \right]$$

$$\log L(Y = 2) = \log [1 - \Phi(\mu_2 - X\beta)]$$

where $\Phi(.)$ and $\phi(.)$ denote the cumulative normal distribution and normal density respectively. INTERVAL uses analytic first and second derivatives to obtain maximum likelihood estimates via the Newton-Raphson algorithm. This algorithm usually converges fairly quickly. TSP uses zeros for starting parameter values. @START can be used to provide different starting values (see NONLINEAR). Multicollinearity of the independent variables is handled with generalized inverses, as in the other linear and nonlinear regression procedures in TSP.

If you wish to estimate a nonstandard ordered probit model (e.g. adjusted for heteroskedasticity or with a nonlinear regression function), use the ML command. See the website for examples of how to do this.

**Options**

**LOWER=** scalar or series containing lower bounds (required).
INTERVAL

UPPER= scalar or series containing upper bounds (required).

The usual nonlinear options are available - see the NONLINEAR section of this manual.

Example

A simple example, showing how to estimate a binary Probit model using PROBIT and INTERVAL with scalars as the lower and upper bounds for the dependent variable.

\begin{verbatim}
PROBIT D C X1-X8 ;  \textit{Probit estimation, }D=0 \textit{ or }1.
Q = 2*D-1 ;  \textit{redefine dep variable to be }(-1,0)
INTERVAL (LOWER=0,UPPER=0) Q C X1-X8 ;
\end{verbatim}

A more complex example, where there are 4 categories (<40, 40 to 50, 50 to 60, and >60), showing how to code the lower and upper bounds and the dependent variables. YCAT takes on the values 1 to 4 corresponding to the four categories.

\begin{verbatim}
yrec = 35*(ycat=1)+45*(ycat=2)+55*(ycat=3)+65*(ycat=4) ;
ylo = 40*(ycat=1)+40*(ycat=2)+50*(ycat=3)+60*(ycat=4) ;
yhi = 40*(ycat=1)+50*(ycat=2)+60*(ycat=3)+60*(ycat=4) ;
interval (lower=ylo,upper=yhi) yrec c x1-x8 ;
\end{verbatim}

Note that by coding the upper and lower limits to be equal for YCAT=1 and YCAT=4 we have specified that they represent a single bound (upper in the case of YCAT=1 and lower in the case of YCAT=4).

Reference

KALMAN

Output  Options  Examples  References

KALMAN estimates linear models using the Kalman filter method. It can handle fairly general State Space models, but it is typically used to estimate regression-type models where the coefficients follow a random process over time.

```
KALMAN (BPRIOR=<prior vector>, BTRANS=<matrix of coefficients in transition equation>, EMEAS, ETRANS, PRINT, SILENT, SMOOTH, VBPRIOR=<variance of prior>, VMEAS=<variance factor in measurement equation>, VTRANS=<variance factor in transition equation>, XFIXED=<X matrix for measurement equation>) <list of dependent variables> [| <list of independent variables>];
```

Usage

The Kalman filter model consists of two parts (the state space form):

**measurement equation:**

\[ y_t = \beta_t + \varepsilon_t, \; \varepsilon_t \sim N(0, \sigma^2_H) \]

**transition equation:**

\[ \beta_t = \beta_{t-1} + \eta_t, \; \eta_t \sim N(0, \sigma^2_Q) \]

**initial conditions:**

\[ \beta_0 \sim N(\overline{\beta}, \sigma^2_P) \]

The matrices \( T, H, \) and \( Q \) are assumed to be known, and they each default to the identity matrix if they are not supplied by the user in the KALMAN options list. Note that they are not allowed to vary over time, but this constraint can be easily relaxed by running KALMAN within a loop over the sample. The NOEMEAS and NOETRANS options are used to zero the variances of the measurement and transition equations respectively.
The \( y(t) \) and \( X(t) \) variables are the dependent and independent variables, just as in ordinary least squares. If you want to use more than one dependent variable, list all the \( y \) variables first, then a \( | \), and then list the \( X \) variables for each \( y \) (duplicate the \( X \) list if they are the same for every \( y \)). You may want to insert zeros along with the \( X \) variables to prevent cross-equation restrictions (see the Examples). If \( X(t) \) is fixed over the sample, use the XFIXED option.

To get a time-varying parameter model, specify VTRANS=Q (the noise-to-signal ratio) and BTRANS=T (if it is not the identity matrix).

To evaluate the likelihood function for general ARMA(p,q) models, fill the BTRANS and VTRANS matrices with the estimated coefficients for the model; see Harvey, p.103 for the general form.

Output

A standard table of coefficients and standard errors is printed for the final state vector, along with the log likelihood. The following items are stored (and may be printed):

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@COEF</td>
<td>vector</td>
<td>m</td>
<td>Final state vector</td>
</tr>
<tr>
<td>@SES</td>
<td>vector</td>
<td>m</td>
<td>Standard errors</td>
</tr>
<tr>
<td>@T</td>
<td>vector</td>
<td>m</td>
<td>T-statistics</td>
</tr>
<tr>
<td>@VCOV</td>
<td>matrix</td>
<td>m*m</td>
<td>Variance-covariance matrix</td>
</tr>
<tr>
<td>@LOGL</td>
<td>scalar</td>
<td>1</td>
<td>Log of likelihood function</td>
</tr>
<tr>
<td>@SSR</td>
<td>scalar</td>
<td>1</td>
<td>Sum of squared recursive residuals</td>
</tr>
<tr>
<td>@S2</td>
<td>scalar</td>
<td>1</td>
<td>Variance of recursive residuals</td>
</tr>
<tr>
<td>@RES1</td>
<td>matrix</td>
<td>#obs*n</td>
<td>Prediction errors (one step ahead)</td>
</tr>
<tr>
<td>@RECRES</td>
<td>matrix</td>
<td>#obs*n</td>
<td>Recursive residuals (i.i.d.)</td>
</tr>
<tr>
<td>@STATE</td>
<td>matrix</td>
<td>#obs*m</td>
<td>Evolving state vectors</td>
</tr>
<tr>
<td>@RESD</td>
<td>matrix</td>
<td>#obs*n</td>
<td>Direct residuals (if SMOOTH is on)</td>
</tr>
<tr>
<td>@SMOOTH</td>
<td>matrix</td>
<td>#obs*m</td>
<td>Smoothed state vectors</td>
</tr>
</tbody>
</table>

Note that the first few rows in the residuals or state vectors may be zero if those observations were used to calculate priors. Note also that recursive residuals for OLS regressions can be obtained using OLSQ with the following options set:

\[ \text{REGOPT (CALC) RECRES;} \]

Method
The Kalman filter recursively updates the estimate of beta(t) (and its variance), using the new information in $y(t)$ and $X(t)$ for each observation, so it can be viewed as a Bayesian method. However, the user does not have to supply priors; they are calculated automatically for regression-type models from the initial observations of the sample. See the Harvey reference for the actual updating formulas. If the default prior is singular, KALMAN adds one more observation to the calculation. The smoothed state vectors (if requested) are estimates based on the full sample; again, see Harvey for the details.

The method of estimation for each time period is maximum likelihood conditional on the data observed to that point. An orthonormalizing transformation of X is used to improve accuracy.

The variance sigma squared and the log likelihood are computed from the recursive residuals. The recursive residuals are

$$e_t = F_t^{-1/2} v_t$$

in Harvey's notation, so that $E[e(t)'e(t)] = I$. If you do not factor sigma squared out of $H$, $Q$, and $P(0)$, the estimated $@S2$ should be close to unity. For $m > 1$, if the prediction errors are collinear, there may be problems estimating sigma squared, the standard errors, recursive residuals, and log likelihood.

**Options**

- **BPRIOR**= the vector of prior coefficients $b(0)$ for measurement equation. Required if XFIXED is used; otherwise it will be calculated by default from a regression in the initial observations of the sample. If the first m observations are not sufficient to identify the prior, one observation is added and BPRIOR is estimated again.

- **BTRANS**= T, the matrix of coefficients in the transition equation (default identity matrix).

- **EMEAS**/NOEMEAS indicates the presence of an error term in the measurement equation (NOEMEAS or NOEM is the same as VMEAS = zero).

- **ETRANS**/NOETRANS indicates the presence of an error term in the transition equation.

- **PRINT**/NOPRINT prints the prior, @STATE, @RES1, @RECRES, @SMOOTH, @RESD.

- **SILENT**/NOSILENT turns off most of the output.
KALMAN

SMOOTH/NOSMOOTH computes fixed-interval smoothed estimates of the state vector b(t) (stored in @SMOOTH) and the direct residuals @RESD.

VBPRIOR= P(0), the variance of the prior (symmetric matrix). Required if BPRIOR is specified. Note: sigma squared is factored out of this matrix.

VMEAS= H, the variance of the measurement equation (symmetric matrix). Default: identity matrix. In Harvey's notation, this is SHS'.

VTRANS= Q, the variance of the transition equation (symmetric matrix). Default: identity matrix. Specifies the "noise-to-signal ratio" if H = identity matrix. In Harvey's notation, this is RQR'.

XFIXED= X matrix for measurement equation, when it is fixed over time.

Examples

One of the simplest Kalman filter models is equivalent to OLSQ (using a transition equation of b(t) =b(t-1) =b). This model can be estimated with the command:

**KALMAN (NOETRANS) Y C X;**

which produces the same coefficient estimates as OLSQ Y C X ;, but calculates them recursively, along with the recursive residuals.

To estimate a Cooley-Prescott "adaptive regression" model where b(t) follows random walk with a nondiagonal variance matrix:

**KALMAN (VTRANS=NSRMAT) Y C X1 X2;**

A "stochastically convergent parameter" model (convergent towards zero in this case, since the transition matrix has roots less than one):

**MFORM (TYPE=DIAG,NROW=3) TMAT=.9; KALMAN (BTRANS=TMAT,VTRANS=NSRMAT) Y C X1 X2;**

Here is an example with two dependent variables; note the two lists of exogenous variables, which must be of the same length. In this case, both equations are forced to have the same two coefficients.

**KALMAN (NOET) Y1 Y2 | C1 X1, C2 X2;**

The example below has two dependent variables, but in this case the equations have separate coefficients; note the use of zero variables. This specification is still somewhat unrealistic because H is identity (same variance and no correlation between errors in the equations):
ZERO = 0;
KALMAN (NOET) Y1 Y2 | C1 X1 ZERO ZERO, ZERO ZERO C2 X2;

Harvey’s Example 2 (p.116-117) ("signal plus noise" or Cooley-Prescott):

READ Y; 4.4 4.0 3.5 4.6;
SET Q = 4; SET A0 = 4; SET P0 = 12;
KALMAN (VT=Q,BPRIOR=A0,VBPRIOR=P0) Y C;

Harvey’s Exercise 1 (p.119) (stochastically convergent):

READ Y; 4.4 4.0 3.5 4.6;
Y4 = Y-4;
SET RHO=.5; SET Q = 4; SET A0 = .2; SET P0 = 3;
KALMAN (BT=RHO,VT=Q,BP=A0,VBP=P0) Y4 C;

Bootstrapping a variance for the transition equation:

SMPL 1,100;
KALMAN (NOETRANS) Y C X1 X2;
UNMAKE @STATE B1-B3;
SMPL 4,100;
DOT 1-3;
D.=B.-B.(-1);
ENDDOT;
COVA D1-D3;
MAT VTOS=@COVA/@S2;
KALMAN (VTRANS=VTOS) Y C X1 X2;

Hyperparameter estimation using ML PROC. This example estimates the variances of the transition matrix Q, using the ML PROC.

MFORM(NROW=2,TYPE=SYM) Q;
PARAM Q11,1 Q22,2;
ML KFQ Q11,Q22;
? KFQ evaluates log likelihood for ML PROC
PROC KFQ;
   IF (Q11<=0 .OR. Q22<=0); THEN; ? Check constraints
     SET @LOGL=@MISS; ? Are variances >0?
   ELSE; DO; ? yes, evaluate.
     SET Q(1,1) = Q11; SET Q(2,2) = Q22;
     SUPRES @COEF;
     KALMAN(SILENT,VTRANS=Q) Y C X;
     NOSUPRES @COEF;
   ENDDO;
ENDDPROC;
References


KEEP (Databank)

**Examples**

KEEP causes variables to be saved in the currently open TSP databank(s).

```plaintext
KEEP <list of variable names> ;
or
KEEP ALL ;
```

**Usage**

KEEP is followed by a list of variable names to be saved in the databank(s) named in the last OUT statement executed. It marks the variables for storage at that point in your program. Variables are automatically saved when they are changed or created (with GENR or SET statements, by matrix computations, or as the output of CAPITL, SAMA, etc.), but it may be convenient to specify explicitly the storing of such variables as equations. Any TSP variable may be named in a KEEP statement: including equations, constants, parameters, matrices, series, or models.

If you want to avoid cluttering up your databank with intermediate results or when you have errors in your run, store data by using an OUT statement followed by a KEEP statement explicitly naming everything you want stored at the very end of your run. Set MAXERR to zero, and TSP aborts before storing the data if it encounters any errors.

Supplying the keyword ALL forces all variables to be stored in any open databanks. Note that PROCs cannot be stored in a databank.

**Output**

KEEP produces no printed output. The variables named are placed in data storage with flags so they will be stored in the appropriate databank at the end of the TSP run.

**Examples**

```plaintext
OUT PDATA ;
FRML EQ1 Y1 = A1+B11*LNP1+B12*LNP2/(A1+G11*LNP1+G12*LNP2) ;
FRML EQ2 Y2 = A2+B12*LNP1+B22*LNP2/(A1+G12*LNP1+G22*LNP2) ;
PARAM A1 A2 B11 B12 B22 G11 G12 G22 ;
LSQ EQ1 EQ2 ;
KEEP EQ1 EQ2 A1 A2 B11 B12 B22 G11 G12 G22 ;
```
This example specifies and estimates a two-equation nonlinear least squares model and saves the equations and parameter estimates on the databank PDATA. Note that the LSQ statement causes the current parameter values to be stored automatically when it finishes, so that it is redundant to specify them on the KEEP statement. However, the equations EQ1 and EQ2 will not be stored unless you name them on a KEEP statement.

```
DOT US UK SW D ;
   OUT DB. ;
   KEEP GNP. CONS. P. DP. DW. U. ;
ENDDOT ;
```

This example shows how data can be placed in different databanks in one run. Identical databanks have been created to hold the series of each of four countries separately: DBUS (United States), DBUK (United Kingdom), DBSW (Sweden), and DBD (Germany). The same set of series is stored in each of the databanks using the KEEP statement.
KERNEL

Options References

KERNEL computes a kernel density estimation or regression. Kernel estimation is a semi-parametric method for approximating a probability distribution.

\[
\text{KERNEL (BANDWIDTH=<bandwidth>,RELBAND=<relative \ bandwidth>,IQR) <variable> ;}
\]

or

\[
\text{KERNEL (BANDWIDTH=<bandwidth>,RELBAND=<relative \ bandwidth>,IQR) <dependent variable> <independent variable> ;}
\]

Usage

When KERNEL is used with a single argument, a Gaussian kernel density of the variable is computed and stored in @DENSITY. You can display the result using a GRAPH command with the variable and @DENSITY as arguments.

When KERNEL is used with two arguments, a Gaussian kernel regression of the first variable on the second is computed; the smoothed values of the dependent variable are stored in @FIT.

The default bandwidth for both estimators is RELBAND=1, which uses Silverman's default bandwidth:

\[
h = \text{RELBAND} \cdot h_0 \cdot 0.9^{\text{NOB}^{(-0.2)}},
\]

where \( h_0 \) is the standard deviation of the independent variable for the default NOIQR option, and the minimum of the standard deviation and the interquartile range divided by 1.349 for IQR. When the number of observations is one, \( h=1 \) is used. For values of RELBAND < 1, the fit is closer to the data (less smooth), while values of RELBAND > 1 fit less closely to the data (more smooth).

Output

KERNEL produces no printed output. A series called @DENSITY is stored when there is one argument and @FIT is stored when there are two arguments.

Options

BANDWIDTH= specifies the absolute value of the bandwidth.
KERNEL

**RELBand** specifies the bandwidth relative to $h$, the default bandwidth.

**IQR/NIOQR** specifies whether the interquartile range is to be used to compute the bandwidth.

**Method**

Given the observed data series $x(i)$, $i=1,...,N$, the Kernel estimator $f(x)$ of the density of $x$ may be obtained using the following equation:

$$ f(x) = \frac{1}{Nh} \sum_{i=1}^{N} K \left( \frac{x - x_i}{h} \right) $$

where $K$ is the kernel function and $h$ is a 'band width' or smoothing parameter TSP uses the Normal or Gaussian kernel and a method based on a Fast Fourier Transform to evaluate this density.

The Kernel regression of $y$ conditional on $x$ is computed using the following equation:

$$ f(y | x) = \frac{\sum_{i=1}^{N} y_i K \left( \frac{x - x_i}{h} \right)}{\sum_{i=1}^{N} K \left( \frac{x - x_i}{h} \right)} $$

Neither estimator is very sensitive to the choice of kernel function, but both are sensitive to the choice of bandwidth $h$. The options allow the user to control the bandwidth either in absolute size or in size relative to the variance or interquartile range (if IQR is used) of the series. The default value of $h$ is given by

$$ h = h_0 0.9 N^{-0.2} $$

where $h_0$ is the standard deviation of the $x$ series. Silverman (1986) shows that this choice has good mean squared error properties.

**References**


LAD computes least absolute deviations regression, also known as L1 regression or Laplace regression. This type of regression is optimal when the disturbances have the Laplace distribution and is better than least squares (L2) regression for many other leptokurtic (fat-tailed) distributions such as Cauchy or Student's t.

LAD (LOWER=<lower censoring limit>, METHOD=<iteration method>, NBOOT=<# replications>, QUANTILE=<value>, RESAMPLE=<method for computing s.e.s>, SILENT, TERSE, UPPER=<upper censoring limit>) <dependent variable> <list of independent variables> ;

Usage

To estimate by least absolute deviations in TSP, use the LAD command just like the OLSQ command. For example,

LAD CONS,C,GNP ;

estimates the consumption function using L1 (median) regression instead of L2 (least squares) regression.

Various options allow you to perform regression for any quantile and censored L1 regression. Standard errors may be obtained using the bootstrap - see the options below.

Output

The usual regression output is printed and stored (see OLSQ for a table). The likelihood function (@LOGL) and standard error estimates are computed as though the true distribution of the disturbances was Laplace; this is by analogy to least squares, where the likelihood function and conventional standard error estimates assume that the true distribution is normal (with a small sample correction to the standard errors). The additional statistics are shown in the table below.

@PHI contains the sum of the absolute values of the residuals. This quantity divided by the number of observations and squared is an estimate of the variance of the disturbances, and is proportional to the scaling factor used in computing the variances of the coefficient estimates.
The LM test for heteroskedasticity is a modified Glejser test due to Machado and Santos-Silva (2000). This test is the result of regressing weighted absolute values of the residuals on the independent variables.

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@PHI</td>
<td>scalar</td>
<td>1</td>
<td>sum of abs value of residuals</td>
</tr>
<tr>
<td>@IFCONV</td>
<td>scalar</td>
<td>1</td>
<td>=1 if there were no simplex iteration problems, 0 otherwise</td>
</tr>
<tr>
<td>@UNIQUE</td>
<td>scalar</td>
<td>1</td>
<td>=1 if the solution is unique, 0 otherwise</td>
</tr>
</tbody>
</table>

**Method**

The LAD estimator minimizes the sum of the absolute values of the residuals with respect to the coefficient vector $b$:

$$
\min_b \sum_{i=1}^{N} |y_i - X_i b|
$$

The estimates are computed using the Barrodale-Roberts modified Simplex algorithm. A property of the LAD estimator is that there are $K$ residuals that are exactly zero (for $K$ right-hand-side variables); this is analogous to the least squares property that there are only $N-K$ linearly independent residuals. In addition, the LAD estimator occasionally produces a non-unique estimate of the coefficient vector $b$; TSP issues a warning message in this case.

When a quantile other than 0.5 is requested, the formula above is modified slightly.

When the number of observations is greater than 100 and the model is not censored, the estimated variance-covariance of the estimated coefficients is computed as though the true distribution were Laplace:

$$
V(b) = \alpha^2 (X'X)^{-1}
$$

where $\alpha^2$ depends on $\tau$, the quantile used.

- $\alpha^2 = \lambda^2$ when $\tau = 0.5$
- $\alpha^2 = \frac{\tau(1-\tau)}{f[F^{-1}(\tau)]^2}$ in general

When the quantile tau is less than the median,
\[ f[F^{-1}(\tau)] = \frac{\tau}{\lambda} \quad \text{and} \quad \sigma^2 = \lambda^2 \frac{1-\tau}{\tau} \]

and when it is greater than the median,

\[ f[F^{-1}(\tau)] = \frac{1-\tau}{\lambda} \quad \text{and} \quad \sigma^2 = \lambda^2 \frac{\tau}{1-\tau} \]

The variance parameter lambda is estimated as though the data has the Laplace distribution,

\[ f(e) = \exp(-|e|/2\lambda) \]

\[ \hat{\lambda} = \frac{1}{n} \sum_{i=1}^{n} |e_i| \]

This formula can also be derived as the BHHH estimate of the variance-covariance matrix if the first derivative of |e| is defined to be unity at zero, as it is everywhere else. The outer product of the gradients of the likelihood function will then yield the above estimate.

The alternative to these Laplace standard errors is to use the NBOOT= option to obtain bootstrap standard errors based on the empirical density. This is the default when the model is censored, or the number of observations is less than 100. In the case of censored estimation, the Bilias et al (2000) resampling method is used to speed up the computations; this can be changed using the RESAMPLE option.


**Options**

- **LOWER=** the value below which the dependent variable is not observed. The default is no limit.
METHOD= BRCENS/SUBSET. The default method is the Barrodale-Roberts or BRCENS simplex method. The alternative SUBSET method evaluates the objective function for all possible subsets of K observations, where K is the number of RHS variables. (The K observations are fitted exactly to the objective function to yield a vector of parameter estimates, then the objective function is computed for all observations using this vector). For censored LAD or quantile regression, this is guaranteed way of finding the global optimum, but it may take a very long time if K is large.

For uncensored LAD or quantile regression, this is a way of investigating possible multiple optima. There may be multiple global optima (different parameter vectors which yield the same objective function value). If this occurs, the different solutions are stored in @COEFD. To provide a reproducible result, instead of choosing one of the multiple solutions at random, TSP uses the following rules to choose a solution:

1. Try an arithmetic average of the parameter vectors. If this yields the same objective function value, this is the reported solution. (This is equivalent to using the average of the two middle values for the median, when there is an even number of observations).

2. Choose the parameter vector with minimum L1 norm (sum of absolute values of the coefficients). If this yields a tie, use rule 3.

3. Choose the parameter vector with minimum absolute value of the first coefficient. If this yields a tie, pick the first vector of those tied.

Note that METHOD=SUBSETS is not used when doing bootstrapping of the SEs.

NBOOT= number of replications for bootstrap standard errors. For the uncensored model, the default is zero if there are 100 or more observations (conventional standard errors under the assumption that the disturbances are Laplace). Otherwise NBOOT=200. The coefficient values from the bootstrap are stored in a @BOOT, an NBOOT by NCOEF matrix, for use in computing other statistics, such as the 95% confidence interval.

QUANTILE= quantile to fit. The default is 0.5 (the median).
RESAMPLE=BILIAS/DIRECT specifies the resampling method to be used for the bootstrap standard error estimates for the censored model. DIRECT resamples from the original data and runs the censored regression estimator to compute the SEs. BILIAS (the default) zeros the observations where the predicted dependent variable is censored, then resamples from the partially zeroed observations, and runs the uncensored LAD/quantile regression to compute the bootstrap SEs. This method is faster and avoids possible convergence or local optima problems with the DIRECT method. See the Bilias et al (2000) reference for details.

SILENT/NOSILENT suppresses all printed output. The results are stored.

TERSE/NOTERSE suppresses all printed output except the table of coefficient estimates and the value of the likelihood function.

UPPER= the value above which the dependent variable is not observed. The default is no limit. This option cannot be used at the same time as the LOWER= option.

Example

Here is an example that computes the median of the numbers 1-10:

```
SMPL 1,10 ; TREND T ;
LAD T C ; ? reports solution=5, but any solution from 5 to 6 is valid
LAD (METHOD=SUBSETS) T C ; ? reports solution=5.5, average of 5 and 6
```

References


LENGTH

Examples

LENGTH determines the length of a list. It is useful in PROCs which are passed lists as arguments, since the length may be required in matrix dimensions or degrees of freedom calculations. This command counts arguments; it does not compute the length of a series or a matrix.

\[ \text{LENGTH <list of variables or lists> <length of list> ;} \]

Output

The list(s) on the command line are expanded, and the final total number of items is stored in the last argument.

Examples

Count the number of parameters from an estimation:

\[ \text{LSQ EQ1-EQ5;} \]
\[ \text{LENGTH @RNMS NOPAR;} \]

The following commands cause LENA to be stored with a value of 3, and LENAB to be stored with a value of 5:

\[ \text{LIST LA X Y Z;} \]
\[ \text{LENGTH LA LENA;} \]
\[ \text{LIST BB B1-B2;} \]
\[ \text{LENGTH LA BB LENAB;} \]
LIML

Output  Options  Examples  References

LIML computes the limited information maximum likelihood estimator for a single equation linear structural model. To estimate a single equation nonlinear model via LIML, use FIML with unrestricted reduced-form equations for the included endogenous variables (those which appear on the right hand side of the primary equation).

*LIML (BEKKER, FEI, FEPRINT, FULLER=<scalar value>, INST=(<list of instruments>), SILENT, TERSE) <dependent variable> <list of rhs endogenous and exogenous variables> ;*

Usage

LIML's form is identical to the 2SLS/INST command -- specify a list of instruments and the variables in the equation. LIML determines the list of endogenous variables, included exogenous variables, and excluded exogenous variables by comparing the instrument list with the variables in the equation. If there are no endogenous variables, OLSQ is used and a warning is printed. If the right hand side equation is exactly identified (number of endogenous variables equals number of excluded exogenous variables), LIML is equivalent to 2SLS, so 2SLS is used, and a warning is printed. If the equation is under-identified, an error message is printed (just like 2SLS).

Output

The output of LIML begins with an equation title, the name of the dependent variable and the lists of endogenous, included exogenous and excluded exogenous variables. The LIML eigenvalue and an F-test of the overidentifying restrictions are printed. The log likelihood at the solution is stored, but not printed.

If FULLER is used, the FULLER constant and the computed K-class value are also printed.

This is followed by various statistics on goodness-of-fit: the sum of squared residuals, the standard error of the regression, the R-squared, and the Durbin-Watson statistic for autocorrelation of the residuals.
The estimated concentration parameter (\textit{mu squared}) and Cragg-Donald F-statistic (CDF) are also shown and stored. When there is a single right hand side endogenous variable, CDF is an F-statistic which tests if the excluded exogenous variables (called Z2) have zero coefficients in the reduced form; the concentration parameter is equal to CDF times the number of excluded exogenous variables. These statistics are also valid for multiple RHS endogenous variables. They can be used to assess whether the model has a "weak instruments" or "many instruments" problem. For a single RHS endogenous variable, the bias of 2SLS is proportional to rho/CDF (Nelson and Startz 1990) where rho is the correlation between the reduced form and structural equations), so low values of the CDF imply a high bias. For a single RHS endogenous variable, values of CDF lower than 10 are considered to be problematic (Staiger and Stock (1997), refined in Stock and Yogo (2004). Unrealistically low computed standard errors for 2SLS also occur in such a situation and the use of LIML(FULLER=1) or plain LIML with the Bekker SEs instead of 2SLS is generally advised because these estimators have much lower bias and properly sized standard errors, especially when the number of excluded instruments is large.

LIML(FULLER=1) and LIML can still suffer from a "weak instruments" problem. For a single RHS endogenous variable, values of mu squared lower than 10~15 suggest a problem (bias and/or standard errors that are too small - see Hansen, Hausman and Newey (2004). For larger values of mu squared, LIML(FULLER=1) and LIML have low bias and properly sized standard errors. LIML(FULLER=1) has a slightly higher median bias than plain LIML, but it is mean unbiased, and it has a smaller MSE than LIML (it has finite moments).

Following this is a table of right hand side variable names, estimated coefficients, standard errors and associated t-statistics. If the variance-covariance matrix has not been suppressed (see the \texttt{SUPRES} command), it is printed after this table. Finally, if the RESID and \texttt{PLOTS} options are on, a table and plot of the actual and fitted values of the dependent variable and the residuals is printed.

LIML also stores most of these results in data storage for later use. The table below lists the results available after a LIML command. The fitted values and residuals will only be stored if the RESID option is on (the default).

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@LHV</td>
<td>list</td>
<td>1</td>
<td>Name of the dependent variable</td>
</tr>
<tr>
<td>@RNMS</td>
<td>list</td>
<td>#vars</td>
<td>Names of right hand side variables</td>
</tr>
<tr>
<td>@SSR</td>
<td>scalar</td>
<td>1</td>
<td>Sum of squared residuals</td>
</tr>
<tr>
<td>@S</td>
<td>scalar</td>
<td>1</td>
<td>Standard error of the regression</td>
</tr>
<tr>
<td>@S2</td>
<td>scalar</td>
<td>1</td>
<td>Standard error squared</td>
</tr>
</tbody>
</table>
If the regression includes PDL variables, the following will also be stored:

@SLAG scalar 1 Sum of the lag coefficients
@MLAG scalar 1 Mean lag coefficient (number of time periods)
@LAGF vector #lags Estimated lag coefficients, after "unscrambling"

Method

The LIML eigenvalue is the minimum eigenvalue of the following matrix:
\[ H^{-1}H \]

\( H \) is the residual covariance matrix of the endogenous and dependent variables regressed on all of the instruments. \( H_1 \) is the residual covariance matrix of the same variables regressed on just the included instruments. This eigenvalue is calculated by CACM algorithm 384. The FULLER constant term (if non-zero) is subtracted from this eigenvalue to yield \( K \), which is then used in the standard K-class formula to compute the coefficients. The standard errors for the NOBEKKER option are computed from the K-class inverse matrix times the sum of squared residuals divided by (number of observations minus number of estimated coefficients).

The F-test for overidentifying restrictions is given by

\[
FSTAT = (LAMBDA - 1)(T - NZ)(K2 - G1)
\]

where \( LAMBDA \) is the LIML eigenvalue, \( T \) is the number of observations, \( NZ \) is the number of instruments, and \( K2 - G1 \) is the number of overidentifying restrictions = number of excluded exogenous variables \( K2 \) minus the number of included endogenous variables \( G1 \). See Anderson et al (1986).

The reported log likelihood is the same as would be computed by FIML on the model plus additional unconstrained equations for the right hand side variables as functions of the instruments.

Options

BEKKER/NOBEKKER specifies that Bekker standard errors are to be computed (see Hansen, Hausman, and Newey 2004). These standard errors are better for small samples and/or when there are large numbers of excluded instruments.

FEI/NOFEI specifies whether a model with individual fixed effects is to be estimated. FREQ (PANEL) must be in effect.

FEPRINT/NOFEPRINT specifies whether the estimated effects and their standard errors are to be printed.

FULLER= value used to weight the eigenvalue towards zero. The formula used is

\[
K = LAMBDA - FULLER/(T - NZ),
\]
where $K$ is the K-class constant, $\text{LAMBDA}$ is the LIML eigenvalue, $T$ is the number of observations, and $NZ$ is the number of instruments. FULLER=0 (default) is the standard LIML estimator, which is median-unbiased. FULLER=1 yields a mean-unbiased estimator. FULLER values between 0 and $8-16/(T-NZ-2)$ dominate LIML in small-sigma efficiency. The LIML estimator modified in this way has smaller tails than the standard LIML estimator, which gives it good small-sample properties (see the references for details).

**INST**= (list of instruments). This list should include all the exogenous variables in the equation being estimated as well as the other exogenous variables in the model. Do not forget to include a constant if there is one in the model. Weights are not supported at present.

SILENT/NOSILENT suppresses all output.

TERSE/NOTERSE prints minimal output (estimated coefficients and a summary statistic).

**Examples**

This example estimates the consumption function for the illustrative model in the TSP User's Manual, using the constant, trend, government expenditures (G), and the log of the money supply (LM) as instruments:

```
LIML (INST=(C,G,TIME,LM)) CONS C GNP ;
```

Other examples:

```
LIML (INST=(C,LOGR,LOGR(-1),LOGR(-2),LOGR(-3)) LOGP C LOGP(-1) LOGR ;
LIML (FULLER=1,INST=(C,LOGR,LOGR(-1),LOGR(-2),LOGR(-3)) LOGP C LOGP(-1) LOGR ;
```

**References**


LIST gives a single name to a list of TSP variables for use later in the program wherever that list of variables is needed. It provides a convenient way to handle long lists of variables repeatedly used. After you define a list, any time it appears in a command, the contents of the list replace the listname. Lists may also be subscripted or lagged. PROC arguments are also treated as lists (the same type of replacement occurs).

\textbf{LIST (FIRST=<number>, LAST=<number>, PREFIX=<name>, SUFFIX=<name> or <value>) <list name> [=] <list of variable names> ; or}
\textbf{LIST (DROP) <list name> <list of variable names> ; or}
\textbf{LIST (PRINT, DELETE) <list of listnames> ;}

\textbf{Usage}

The LIST name can be any legal TSP variable name. Follow this with any number of legal TSP variable names (they may include lags or leads) including the names of other lists.

Lists may be nested indefinitely, that is, the names following the LIST name may be LISTS. The only limitation on length is that on the ultimate list of variables. LISTS may not be defined recursively -- the list name cannot appear in its own list when a list is first defined. An example of an illegal recursive definition is \texttt{LIST LL A B LL;} (if \texttt{LL} has not already been defined as a list).

A LIST of variables may be specified as a range, i.e., \texttt{VAR1-VAR20}. A list like this is interpreted as \texttt{VAR1, VAR2, VAR3, ..., VAR19, VAR20}. You can also use a range expression for numbers or for lags (\texttt{X(-1)-X(-20)}). Such an implicit list can also appear directly in a command (see the DOT command below). You can combine this type of list with a DOT loop to operate on individual elements of the list very conveniently:

\texttt{LIST VARLIST VAR1-VAR20 ;}
\texttt{DOT 1-20 ;}
\texttt{------- computations on VAR. -------}
\texttt{ENDDOT ;}
\texttt{------- operations on VARLIST -------}
This enables you to use the variables as a group or as individual variables in GENR (since listnames are not allowed in GENR, SET, or equation specifications).

A special list named @ALL is always available; it contains the names of all the series in the current TSP run.

Output

LIST produces no printed output. A TSP variable list is stored in data storage. SHOW LIST; can be used to see the currently defined lists, and LIST (PRINT) can be used to view the contents of one or more lists.

Options

DELETE/NODELETE deletes existing lists.

DROP/NODROP drops variables from an existing list.

FIRST= starting integer for a sequence of names in a list. The default is 1. FIRST can be larger than last, to make a list in decreasing order. It cannot be negative.

LAST= ending integer for a sequence of names in a list. The default is 1, and it cannot be negative.

PREFIX= name to be used as the prefix in constructing a list in the form: namelfirst-namelast. See the examples. PREFIX can also be used to append a list of different names to a common prefix.

SUFFIX= name or number to be added to all the variable names in the list.

PRINT/NOPRINT prints the contents (i.e. included names) of existing lists.

Examples

A few simple examples:

```
LIST EQ EQ1-EQ4 ;  creates LIST EQ EQ1 EQ2 EQ3 EQ4 ;
LIST EQS EQ01-EQ04 ; ? creates LIST EQS EQ01 EQ)@ EQ03 EQ04 ;
LIST LAGS X(-1)-X(-20) ;
LIST STATES 1-50 ;
MSD @ALL ;            ?prints simple statistics for all data series.
```

```
LIST INSTVAR C TIME LM G ;
LIST ENDOGVAR GNP CONS I R LP ;
LIST ALLVARS INSTVAR ENDOGVAR ;
```
The list ALLVARS consists of the series C, TIME, LM, G, GNP, CONS, I, R, and LP.

The following example shows the power of implicit lists when combined with DOT loops to eliminate repetitive typing:

```
PRINT PAT72-PAT76 RND72-RND76 ;
PARAM A72-A76 DELT72-DELT76 BETA ;
DOT 72-76 ;
FRML EQ. PAT. = EXP(A.+BETA*RND.+DELT72*RND72 +
DELT73*RND73 +
DELT74*RND74 + DELT75*RND75 +DELT76*RND76) ;
PARAM A. 1.0 DELT. ;
ENDDOT ;
LSQ (NOPRINT,STEP=BARDB) EQ72-EQ76 ;
```

In the above example, the series, equations, and parameters for a panel data model (5 years of data on each of several hundred units) can all be referred to by their LIST names to save the repetitive typing of each year's variables. Obviously, LISTs defining more than one variable cannot be used within equations, because they do not reduce to algebraic expressions, but it is useful in some applications to use EQSUB in a DOT loop to reproduce similar equations.

Suppose you do not know the number of variables in a LIST until runtime; you can use the options to construct a variable length list in this case:

```
BEGYR=72 ; ENDYR=76 ;
LIST (PREFIX=EQ,FIRST=BEGYR,LAST=ENDYR) EQS ;
```

creates a list called EQS consisting of EQ72, EQ73, EQ74, EQ75, and EQ76.

More simple examples:

```
LIST (FIRST=5,LAST=1) DECR;   ? 5 4 3 2 1
LIST (DROP) DECR 3 2;          ? 5 4 1
LIST (PREFIX=B) BS X Y Z(-1);  ? BX BY BZ(-1)
LIST (SUFFIX=5) X5 LW ED EX ;  ? X5 is LW5 ED5 EX5
LIST (SUFFIX=US) XLIST LW ED EX ;  ? XLIST is LWUS EDUS EXUS
LIST BSMID BS(2);              ? Y (Subscripted list)
PRINT BS(-1);                  ? X(-1) Y(-1) Z(-2) (Lagged list)
LIST (PRINT) BS;               ? prints the names X Y Z(-1), but not their values
LIST (DELETE) BS;              ? removes the list definition for BS
```
Programming trick: in fact, a list can be used to store any sequence of words or arguments. Here are some examples of lists which do not contain variable names, but which TSP will understand:

```
LIST STRING 'A title string';
TITLE STRING;          ? same as TITLE 'A title string';
LIST OPTIONS MEAN=5;
RANDOM (OPTIONS) X;    ? same as RANDOM(MEAN=5) X;
```
LMS computes least median of squares regression. This is a very robust procedure that is useful for outlier detection. It is the highest possible "breakdown" estimator, which means that up to 50% of the data can be replaced with bad numbers and it will still yield a consistent estimate. Proper standard errors (such as asymptotically normal) for LMS coefficients are not known at present.

\textit{LMS (ALL, LTS, MOST, PRINT, SILENT, SUBSETS=<value>, TERSE)<dependent variable> <list of independent variables> ;}

**Usage**

To estimate by least median of squares in TSP, use the LMS command just like the OLSQ command. For example,

\textit{LMS CONS,C,GNP ;}

estimates the consumption function. The PRINT option enables the printing of the outliers, or you can define a set of outliers for printing by screening on the residuals, which are stored in @RES.

**Output**

The usual regression output is printed and stored (see \texttt{OLSQ} for further discussion). The number of possible subsets and the best subset found are also printed. The following results are stored:

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@LHV</td>
<td>list</td>
<td>1</td>
<td>name of dependent variable</td>
</tr>
<tr>
<td>@RNMS</td>
<td>list</td>
<td>#vars</td>
<td>list of names of independent variables</td>
</tr>
<tr>
<td>@YMEAN</td>
<td>scalar</td>
<td>1</td>
<td>mean of dependent variable</td>
</tr>
<tr>
<td>@SDEV</td>
<td>scalar</td>
<td>1</td>
<td>standard deviation of dependent variable</td>
</tr>
<tr>
<td>@SSR</td>
<td>scalar</td>
<td>1</td>
<td>sum of squared residuals</td>
</tr>
<tr>
<td>@S2</td>
<td>scalar</td>
<td>1</td>
<td>standard error squared</td>
</tr>
<tr>
<td>@S</td>
<td>scalar</td>
<td>1</td>
<td>standard error of estimate</td>
</tr>
<tr>
<td>@RSQ</td>
<td>scalar</td>
<td>1</td>
<td>R-squared</td>
</tr>
<tr>
<td>@ARSQ</td>
<td>scalar</td>
<td>1</td>
<td>adjusted R-squared</td>
</tr>
</tbody>
</table>
Method

The LMS estimator minimizes the square (or the absolute value) of the median residual with respect to the coefficient vector $b$:

$$\min_b \text{median}\{y_i - X_i b\}$$

Clearly this ignores the sizes of the largest residuals in the sample (i.e. those whose absolute values are larger than the median), so it will be robust to the presence of any extreme data points (outliers).
If there are $K$ independent variables (excluding the constant term), LMS will consider many different subsets of $K$ observations each. An exact fit regression line is computed for each subset, and residuals are computed for the remaining observations using these coefficients. The residuals are essentially sorted to find the median, and a slight adjustment is made to allow for the constant term. If $K$ or the number of observations is large, the number of subsets could be very large (and sorting time could be lengthy), so random subsets will usually be considered in this case. This is controlled with the ALL, MOST, and SUBSETS= options.

The result is not necessarily the global Least Median of Squares optimum, but a feasible close approximation to it, with the same properties for outlier detection. Least Trimmed Squares (LTS) also has about the same properties. The LMS estimator occasionally produces a non-unique estimate of the coefficient vector $b$; TSP reports the number of non-unique subsets in this case. It uses the following steps to determine the preferred solution:

1. Try an arithmetic average of the parameter vectors. If this yields the same objective function value, this is the reported solution. (This is equivalent to using the average of the two middle values for the median, when there is an even number of observations).

2. Choose the parameter vector with minimum L1 norm (sum of absolute values of the coefficients). If this yields a tie, use rule 3.

3. Choose the parameter vector with minimum absolute value of the first coefficient. If this yields a tie, pick the first vector of those tied.

An extremely rough estimate of the variance-covariance of the estimated coefficients is computed with an OLS-type formula:

$$V(\hat{\beta}) = \sigma^2 (X'X)^{-1}$$

where the estimate of $\sigma$ squared is computed after deleting the largest residuals. This estimate is not asymptotically normal, and is likely to be an underestimate, so it should not be used for serious hypothesis testing. It all depends on how the outliers are generated by the underlying model.

The code used in TSP was adapted from Rousseeuw’s Progress program, obtainable from his web page, referenced below.

Options

ALL/NOALL uses all possible observation subsets (see Method), even if there are over one million of them.
LMS/NOLTS computes the Least Trimmed Squares estimates, which minimize the sum of squared residuals, from the smallest up to the median (instead of LMS, which minimizes just the squared median residual). Usually the LTS and LMS estimates are fairly close to each other.

MOST/NOMOST uses all possible subsets, unless the number of subsets is one million or more (in which case random subsets are used).

PRINT/NOPRINT prints better subsets (as progress towards a minimum is made), and the final outliers.

SILENT/NOSILENT suppresses all output.

SUBSETS= number of random subsets to use. The default is 500 to 3000. If the number of possible subsets is less than the number of random subsets, all possible subsets will be evaluated systematically.

TERSE/NOTERSE suppresses all printed output except the table of coefficient estimates and the value of the objective function.

Examples

LMS Y C Z1-Z6;  ? uses 3000 random subsets
LMS (MOST) Y C Z1-Z6 ;  ? probably all subsets
? 2000 subsets (but not the same set as the default)
LMS (SUBSET=2000) Y C Z1-Z6 ;

References


http://www.agoras.ua.ac.be (Rousseeuw / Progress)
LOAD

LOAD is a synonym for READ.

LOAD (BYOBS, BYVAR, FILE='filename string' or filename,
   FORMAT=BINARY or DATABANK or EXCEL or FREE or LOTUS
   or RB4 or RB8 or '(format text string)', FULL, NCOL=<number of
   columns>, NROW=<number of rows>, PRINT, SETSMPL,
   TYPE=CONSTANT or DIAG or GENERAL or SYMMETRI or
   TRIANG, UNIT=<I/O unit number>) <list of series or matrices> ;
LOCAL

Example

LOCAL specifies the variables which will be considered local to a TSP PROC. Their values will not be saved on exit from the procedure.

LOCAL <list of variables> ;

Usage

The LOCAL statement should be placed following the PROC statement to which it applies. Follow the word LOCAL with the names of variables which will be created during the execution of the PROC, but which will not be needed on exit. This can save storage space, or allow the use of duplicate names, which can be especially convenient if you wish to build a library of PROCs and don't want variable name conflicts.

Output

LOCAL produces no output.

Example

The procedure below computes moving averages of variable length LEN. The local variables LAST, which is the index of the last observation but one and LAG, the loop index, are not saved on return from PROC MA:

PROC MA X,LEN,XMA ;
   LOCAL LAST LAG ;
   XMA=X ;
   SET LAST=1-LEN ;
   DO LAG=LAST TO -1 ;
      XMA = XMA+X(LAG) ;
   ENDDO ;
   XMA=XMA/LEN ;
ENDPROC MA ;
LOGIT

LOGIT

Output  Options  Examples  References

LOGIT is used to estimate a conditional and/or multinomial logit model. The explanatory variables in the model may vary across alternatives (choices) for each observation or they may be characteristics of the observation, or both. There is no limit on the number of alternatives.

\[
\text{LOGIT (CASE=}<\text{series name}>\text{, COND, NCHOICE=<number>, NREC=}<\text{series name}>\text{, SUFFIX=}<\text{list of names}, \text{ nonlinear options}) <\text{dependent variable} > <\text{independent variables}> \ ;
\]

or

\[
\text{LOGIT (CASE=}<\text{series name}>\text{, COND, NCHOICE=<number>, NREC=}<\text{series name}>\text{, SUFFIX=}<\text{list of names}, \text{ nonlinear options}) <\text{dependent variable} > <\text{conditional variables} > | <\text{multinomial(alternative) variables}> \ ;
\]

Usage

There are three types of logit model: those where the regressors are the same across all choices for each observation, i.e., they are characteristics of the chooser, those where the regressors are characteristics of the specific choice, and mixed models, which have regressors of both kinds. In the first case (multinomial logit), a separate coefficient for each regressor is estimated for all but one of the choices. In the second case (conditional logit), the regressors change across the choices, and a single coefficient is estimated for each set of regressors.

1. Binary or multinomial logit -- like OLSQ or PROBIT:

\[
\text{LOGIT <dependent variable} > <\text{multinomial variables (chooser characteristics)}> ;
\]

\[
\text{LOGIT Y C X1 X2 ... XK;}
\]

\(Y\) can be 0/1 or 1/2 or any integral values. If \(Y\) takes on more than 2 values, the model is multinomial logit. The names of the coefficients are determined by appending the values of \(Y\) for each choice to the names of the explanatory variables. The coefficients are normalized by setting the coefficients for the lowest choice to 0. If \(Y\) is 0/1, the coefficients \(C1, X11, X21, \ldots, XK1\) would be estimated, with \(C0, X10, X20, \ldots, XK0\) normalized to zero. If \(Y\) is 1/2/3, the coefficients \(C2, X12, X22, \ldots, XK2\) and \(C3, X13, X23, \ldots, XK3\) would be estimated, with \(C1, X11, X21, \ldots, XK1\) normalized to zero.

\[
\text{LOGIT (NCHOICE=2) Y C X1 X2 ... XK;}
\]
Including the NCHOICE option causes TSP to check the range of \( Y \) to make sure there are only 2 choices. The model estimated has \( K+1 \) coefficients and \( K+1 \) variables.

The multinomial logit procedure checks for "univariate complete and quasi-complete separation", which prevents identification of the coefficients.

2. **Conditional logit:**

\[ \text{LOGIT (COND,NCHOICE=n) <dependent variable> <conditional variables (choice characteristics)>;} \]

For example,

\[ \text{LOGIT (COND,NCHOICE=2) Y X Z;} \]

looks for the variables \( X_1, X_2, Z_1, Z_2 \) corresponding to the 2 choices. The coefficients \( X \) and \( Z \) would be estimated. \( C \) is not allowed as a conditional variable (since it does not vary across choices, it is not identified). For a choice-specific set of dummies, use \( C \) as a multinomial variable in mixed logit. In this case, the example shown becomes

\[ \text{LOGIT (COND,NCHOICE=2) Y X Z | C;} \]

The **CASE** option allows you to use data organized with one choice per observation rather than with one case per observation. For example,

\[ \text{LOGIT (CASE=V) Y X Z ;} \]

where the variable \( V \) is a case ID which is equal for adjacent observations which belong to the same case. In this case, the variable names \( X \) and \( Z \) are used directly (not \( X_1, X_2, \) etc.). There need not be an equal number of observations per case. Only the first \( Y \) for each case is examined for a valid choice number.

For datasets with multiple observations per case, you can also use

\[ \text{LOGIT (NREC=W) Y X Z;} \]

\( W \) specifies the number of records per case and you do not need to supply an ID variable with the CASE option.

3. **Mixed logit:**

The general form of the command is now

\[ \text{LOGIT (COND,NCHOICE=n) <dependent variable> <conditional variables> | <multinomial variables> ;} \]
For example,

`LOGIT (COND,NCHOICE=3) Y ZA ZB / XA XB XC;`

`Y` takes on the values 1, 2 and 3. TSP looks for the conditional variables `ZA1, ZA2, ZA3, ZB1, ZB2, ZB3` corresponding to the 3 choices. `XA, XB, XC` are the multinomial variables. The coefficients `ZA, ZB, XA2, XB2, XC2, XA3, XB3, XC3` would be estimated, with `XA1, XB1, XC1` normalized to zero.

**Output**

The printed output of the LOGIT procedure is similar to that of the other nonlinear estimation procedures in TSP. A title is followed by a table of the frequency distribution of the choices. Then the starting values and iteration-by-iteration printout is printed; the amount is controlled by the PRINT and SILENT options. This is followed by a message indicating final convergence status, the value of the likelihood and a table of parameter estimates and their asymptotic standard errors and t-statistics. The variance-covariance matrix of the estimates is also printed if it has been unsuppressed. The @DPDX or @DPDZ matrices (described below) are printed unless they have been SUPRESed. The following are also stored in data storage:

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@LOGL</td>
<td>scalar</td>
<td>1</td>
<td>Log of likelihood function</td>
</tr>
<tr>
<td>@IFCONV</td>
<td>scalar</td>
<td>1</td>
<td>Convergence status (1 = success)</td>
</tr>
<tr>
<td>@LR</td>
<td>scalar</td>
<td>1</td>
<td>Likelihood ratio test for zero slopes</td>
</tr>
<tr>
<td>@SRSQ</td>
<td>scalar</td>
<td>1</td>
<td>Scaled R-squared for multinomial logit</td>
</tr>
<tr>
<td>@RSQ</td>
<td>scalar</td>
<td>1</td>
<td>Squared correlation between Y and @FIT for binary logit</td>
</tr>
<tr>
<td>@SSR</td>
<td>scalar</td>
<td>1</td>
<td>Sum of squared residuals (Y-@FIT) for binary logit</td>
</tr>
<tr>
<td>@RNMS</td>
<td>list</td>
<td>#params</td>
<td>List of parameter names</td>
</tr>
<tr>
<td>@GRAD</td>
<td>vector</td>
<td>#params</td>
<td>Gradient of likelihood function at maximum</td>
</tr>
<tr>
<td>@COEF</td>
<td>vector</td>
<td>#params</td>
<td>Estimated values of parameters</td>
</tr>
<tr>
<td>@SES</td>
<td>vector</td>
<td>#params</td>
<td>Standard errors of estimated parameters</td>
</tr>
<tr>
<td>@T</td>
<td>vector</td>
<td>#params</td>
<td>T-statistics</td>
</tr>
<tr>
<td>%T</td>
<td>vector</td>
<td>#params</td>
<td>p-values for T-statistics</td>
</tr>
<tr>
<td>@VCOV</td>
<td>vector</td>
<td>#params</td>
<td>Estimated variance-covariance of estimated parameters</td>
</tr>
</tbody>
</table>
| @DPDX    | matrix   | #vars* | Mean of probability derivatives for multinomial variables. This matrix is invariant to the set of coefficients which are normalized to zero (as are the

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differences between sets of coefficients).

@DPDZ matrix \#vars* \#choices
Mean of probability derivatives for conditional variables (#vars = \#condvars* \#choices). This matrix consists of NCHOICE submatrices of dimension (NCOND x NCHOICE) stacked vertically, and it is block-symmetric. Not calculated if NCHOICE varies by case.

@FIT matrix \#obs* \#choices
or series or \#obs
Matrix of fitted probabilities when NCHOICE>2 and there is one observation per case. Length NOB series when NCHOICE=2 or when there are multiple observations per case. If NCHOICE=2, @FIT will contain the probabilities for the highest choice only (just like binary PROBIT).

Method

If \( C(t) \) is the choice set for the \( t \)th observations, and observation \( t \) chooses the \( i \)th alternative out of \( C(t) \), then the expression for the choice probability is

\[
Pr(\text{\( i \) chosen from } C_t | Z) = \frac{\exp(Z_{yi} b)}{\sum_j \exp(Z_{yj} b)}
\]

The likelihood function is

\[
LogL = -\sum_{t=1}^{T} \log \left( \sum_j \exp((Z_{yj} - Z_{yi})b) \right)
\]

The coefficient vector to be estimated is \( b \). If some of the \( Z \)s do not vary across the choices, these equations would apply to an expanded \( Z \) vector formed by taking the Kronecker product of the fixed \( Z \)s with an identity matrix of order of the number of choices (less one for a normalization). The actual implementation does not expand the \( Z \)s, but treats the conditional and multinomial variables differently to conserve space.
Newton's method is used to maximize this likelihood function with respect to the parameter vector \( b \). The global concavity of the likelihood function makes estimates fairly straightforward to obtain with this method. Zero starting values are the default, unless @START is supplied. See the NONLINEAR section in this manual for more information about TSP's nonlinear optimization procedures in general.

The evaluation of the \( \exp() \) functions in the likelihood function and derivatives avoids floating overflows and zero divides. When these conditions occur, the appropriate limit is taken instead. This may result in some slight inaccuracy in the likelihood function, but it is certainly preferable to halting the estimation. Observations subject to these problems can be identified by exact 0 and 1 values in @FIT.

Before estimation, LOGIT checks for univariate complete and quasi-complete separation of the data and flags this condition. The model is not identified in this case, because one or more of the independent variables perfectly predict \( Y \) for some observations, and therefore their coefficients would slowly iterate to + or - infinity if estimation was allowed to proceed.

The scaled R-squared is a measure of goodness of fit relative to a model with just a constant term; it replaced the Kullback-Leibler R-squared beginning with TSP 4.5 since it has somewhat better properties for discrete dependent variable problems. See the Estrella (1998) article.

Options

The standard options for nonlinear estimation are available: see the NONLINEAR section in this manual. Note that HITER=N, HCOV=N are the defaults for the Hessian approximation and standard error computations. In addition, the following options are specifically for the LOGIT procedure:

**CASE**= case series for multiple observations per case. This variable holds a case identification which is equal for adjacent observations that belong to the same case. Note that any such variable may be used; it does not necessarily have to be the case identification.

**COND/NOCOND** for conditional or mixed models versus pure multinomial estimation (see the Usage section). If CASE= or NREC= is used, COND is assumed and need not be specified.

**NCHOICE**= number of choices can be supplied when the number is equal for all observations. The program then checks to make sure the data satisfy this constraint. This is not used with CASE= or NREC=, since then it is valid to take the first choice every time.
LOGIT

**NREC= choice count series** for multiple observations per case. This variable specifies the number of observations in each case. Usually the number is repeated in each observation, but only the count in the first observation for each case is used. You cannot say NREC=3, but you can say NCHOICE=3.

**SUFFIX=** a list of short names (suffixes) for the alternatives. These names are used in 4 places: in the initial table of frequencies for the dependent variable, as coefficient names for multinomial variables, as labels for the probability derivatives $dP/dZ$, and as the suffixes for conditional variable names (when there is one observation per case). Note that **SUFFIX** does not imply **COND**. See the examples below.

The SUFFIX names need to be in the proper order, relative to the values of the dependent variable. In the examples below, $Y=1$ is CAR, $Y=2$ is BUS, and $Y=3$ is RT. If some of the alternatives are never chosen, be sure to use **SUFFIX=** or **NCHOICE=** to ensure the full set of conditional variables is used (variables corresponding to all available alternatives).

**Examples**

```plaintext
LOGIT Y C X ;
```

```plaintext
LOGIT (COND, NCHOICE=3) Y XA XB ;
```

looks for conditional variables XA1 XA2 XA3 and XB1 XB2 XB3, whereas

```plaintext
LOGIT (COND,NCHOICE=3,SUFFIX=(CAR,BUS,RT)) Y XA XB ;
```

looks for conditional variables XACAR, XABUS, XART and XBCAR, XBBUS, XBRT. Note that the suffixes must be in the proper order (1,2,3) for correct interpretation of the output.

See the usage section for other examples of how to use this procedure.

**References**


LOGIT


LP

LP (Linear Programming) solves linear models with or without linear equality or inequality constraints.

\[ \text{LP (MAX/MIN, SILENT, NLW=<number of less than or equal constraints,} \]
\[ \text{NGE=<number of greater than or equal constraints, NEQ=<number of equality constraints, TOL=eps)} \]
\[ b_{\text{obj}} b_{\text{constraints}} \text{ rhs ;} \]

Usage

LP solves for the \( M \) by 1 vector \( X \) that minimizes the objective \( \phi = b_{\text{obj}}'X \), with constraints \( b_{\text{constraints}}'X (<= or >= or =) \text{ rhs} \) and \( X>=0 \). The order of the constraints must be the following: 1) \( \text{NLW} \) less than or equal; 2) \( \text{NGE} \) greater than or equal; 3) \( \text{NEQ} \) equality. \( b_{\text{obj}} \) is an \( M \) by 1 vector and \( b_{\text{constraints}} \) is a \( \text{NLW+NGE+NEQ} \) by 1 vector.

Output

LP stores some of the results in data storage for later use.

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@PHI</td>
<td>scalar</td>
<td>1</td>
<td>The objective function at the minimum</td>
</tr>
<tr>
<td>@IFCONV</td>
<td>scalar</td>
<td>1</td>
<td>=1 if convergence achieved, 0 otherwise</td>
</tr>
<tr>
<td>@COEF</td>
<td>vector</td>
<td>M</td>
<td>the solution for ( X )</td>
</tr>
</tbody>
</table>

Method

The Simplex method subroutine from \textit{Numerical Recipes} (Press et al. 1992) is used, which has a nice explanation of the method and some examples which were used for testing. The LP command was also tested on LAD estimation with the Stackloss dataset (Brownlee 1965; Dodge 1991).

Options

\textbf{MAX/MIN} specifies whether the objective function \( \phi \) is to be maximized or minimized (the default).
SILENT suppresses all output; results are still stored.

NLW = number of less than or equal constraints in b_constraint and rhs.

NGE = number of greater than or equal constraints in b_constraint and rhs.

NEQ = number of equality constraints in b_constraint and rhs.

TOL = convergence criterion.

Examples

Here is an example of using the LP command to solve the first example in the Numerical Recipes description:

? Maximize: 2*x2 - 4*x3
? Subject to: x1 + 6*x2 - x3 = 2
? -3*x2 + 4*x3 + x4 = 8
? The solution is: x2 = 1/3, x4 = 9 (x1=0, x3=0),
? with objective function = 2/3

read(nrow=1,ncol=4) objf1;
0 2 -4 0 ;
read(nrow=2,ncol=1) dep1;
2 8 ;
read(nrow=2,ncol=4) constr1;
1 6 -1 0
0 -3 4 1 ;
LP(NEQ=2,MAX) objf1 constr1 dep1;

References


http://www.nr.com
LSQ is used to obtain least squares or minimum distance estimates of one or more linear or nonlinear equations. These estimates may optionally be instrumental variables estimates. LSQ can compute nonlinear least squares, nonlinear two stage least squares or instrumental variables, nonlinear multivariate regression with cross-equation constraints, seemingly unrelated regression, and nonlinear three stage least squares. The equations for any of these estimators may be linear or nonlinear in the variables and parameters, and there may be arbitrary cross-equation constraints. LSQ can also be invoked with the SUR, 3SLS, THSLS, and GMM commands.

**Usage**

There are four basic estimators available in LSQ: single or multi-equation least squares and single or multi-equation instrumental variables. They are all iterative methods which minimize a distance function of the general form

\[ f(y, X, b)' [S^{-1} \otimes H (H' H)^{-1} H] f(y, X, b) \]

where \( f(y, X, b) \) is the (stacked) vector of residuals from the nonlinear model, \( S \) is the current estimate of the residual covariance matrix being used as a weighting matrix, and \( H \) is a matrix of instruments.

The form of \( f(y, X, b) \) is specified by the user as a FRML, which may be either unnormalized (in the form \( f(y, x, b) \) with no = sign), or normalized (the usual form of \( y = f(x, b) \)). The latter form will cause equation by equation statistics for the estimated model to be printed.

To obtain a particular estimator, various assumptions are made about the exact form of this distance function. These assumptions are described below.

**Nonlinear single equation least squares:** In this case, there are no instruments (\( H \) is identity) and \( S \) is assumed to be unity. This makes the objective function the sum of squared residuals of the model; minimizing this function is the same as obtaining maximum likelihood estimates of the parameters of the model under the assumption of normality of the disturbances.
The form of the LSQ statement for estimating this model is LSQ followed by options in parentheses, and then the name of the equation. Any of the standard NONLINEAR options can be used.

**Nonlinear two stage least squares:** for this estimator, $H$ is the matrix of instrumental variables formed from the variables in the INST option, and $S$ is again assumed to be unity. The estimator is described in Amemiya (1974). If the model is linear, conventional two stage least squares or instrumental variable estimates result.

**Nonlinear multivariate regression:** In this case, there are no instruments ($H$ is the identity matrix) and $S$ is either estimated or fixed. This estimator can be computed with two completely different objective functions. The default in TSP is to compute maximum likelihood estimates if the LSQ command is specified with no instruments and more than one equation. These estimates are obtained by concentrating variance parameters out of the multivariate likelihood and then maximizing the negative of the log determinant of the residual covariance matrix. They are efficient if the disturbances are multivariate normal and identically distributed.

Using the option MAXITW=0, it is possible to obtain minimum distance estimates of a nonlinear multivariate regression model. For these estimates, the objective function is the distance function given above with the instrument matrix $H$ equal to identity. The $S$ matrix is given by the WNAME option: it can be identity, which is similar to estimating each equation separately (except that cross-equation constraints will be enforced and the parameter standard errors will be wrong unless the true residual variances are unity), it can be supplied by you from a previous estimation, or it can be computed from the parameters (the WNAME=OWN option). The $S$ matrix is always a symmetric matrix of the order of the number of equations.

To obtain conventional seemingly unrelated regression estimates of a nonlinear multivariate regression model, use the SUR command, which is a special form of the LSQ command. This version of the procedure obtains single equation estimates of the parameters of the model, uses these to form a consistent estimate of the residual covariance matrix, and then minimizes the objective function shown above with respect to the parameters $b$. If the model is linear, this is a two stage procedure (only two iterations). The plain LSQ command will iterate simultaneously on the parameters and the residual covariance matrix. In this case, linear models may take more than one iteration to converge.
Nonlinear three stage least squares: this estimator uses the distance function as shown, with $S$ equal to a consistent estimate of the residual covariance (either supplied or computed), and $H$ equal to the Kronecker product of an identity matrix of the order of the number of equations and the matrix of instruments. This means that all the instruments are used for all the equations.

Three stage least squares estimates can be obtained in two ways: LSQ with the `WNAME` option, the `INST` option, and more than one equation name will give three stage least squares estimates using the $S$ matrix you specify. Alternatively, if you use the `3SLS` form of the LSQ command with the `INST` option, LSQ automatically computes consistent nonlinear two stage least squares estimates of the parameters, uses them to form an estimate of the residual covariance matrix $S$, and then computes three stage least squares estimates.

To use any of these estimators, first specify the equations to be estimated using FRML statements and name the parameters and supply starting values with `PARAM` statements (an alternative to this is the `FORM (PARAM)` command after a linear estimation procedure). Any parameters which appear in more than one equation are assumed to be the same parameter and the equality constraint is automatically imposed.

LSQ always determines the linearity or nonlinearity of the model; if the model is linear in the parameters, it prints a message to that effect, and uses just one iteration.

Output

LSQ stores its results in data storage. The estimated values of the parameters are stored under the parameter names. The fitted values and residuals will only be stored if the `RESID` option is on (the default). In addition, the following results are stored:

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@LOGL</td>
<td>scalar</td>
<td>1</td>
<td>Log of likelihood function (if valid).</td>
</tr>
<tr>
<td>@TR</td>
<td>scalar</td>
<td>1</td>
<td>Trace of COVT (if the minimum distance estimator is used).</td>
</tr>
<tr>
<td>@PHI</td>
<td>scalar</td>
<td>1</td>
<td>$E'PZ'E$, the objective function for instrumental variable estimation.</td>
</tr>
<tr>
<td>@FOVERID</td>
<td>scalar</td>
<td>1</td>
<td>test of overidentifying restrictions (for 2SLS)</td>
</tr>
<tr>
<td>@IFCONV</td>
<td>scalar</td>
<td>1</td>
<td>Convergence status (1 = success).</td>
</tr>
<tr>
<td>@RNMS</td>
<td>list</td>
<td>#params</td>
<td>list of parameter names</td>
</tr>
<tr>
<td>@GRAD</td>
<td>vector</td>
<td>#params</td>
<td>Gradient of objective function at the convergence</td>
</tr>
</tbody>
</table>
Normal LSQ output begins with a listing of the equations. The model is checked for linearity in the parameters (which simplifies the computations). A message is printed if linearity is found and LSQ does not iterate because it is unnecessary. The amount of working space used by LSQ is also printed - this number can be compared with the amount printed at the end of the run to see how much extra room you have if you wish to expand the model.
Next LSQ prints the values of constants and the starting conditions for the parameters, and then iteration-by-iteration output. If the print option is off, this output consists of only one line, showing the beginning value of the log likelihood, the ending value, the number of squeezes in the stepsize search (ISQZ), the final stepsize, and a criterion which should go to zero rapidly if the iterations are well-behaved. This criterion is the norm of the gradient in the metric of the Hessian approximation. It will be close to zero at convergence.

When the print option is on, LSQ also prints the value of the parameters at the beginning of the iteration and their direction vector. These are shown in a convenient table so that you can easily spot parameters with which you are having difficulty.

Finally LSQ prints the results of the estimation (whether or not it converged); these results are printed even if the NOPRINT or TERSE options are set. The names of the equations and endogenous variables are printed, the value of the objective function at the optimum, and the corresponding estimate of the covariance of the structural disturbances. If minimum distance estimation was used, the trace of the weighted residual covariance matrix is the objective function (the equation given above with $H$ equal to the identity matrix). Otherwise the objective function is the negative of the log of the likelihood function.

If instrumental variable estimation was used, the objective function is labelled $E'PZ^*E$ and stored as @PHI. This is analogous to the sum of squared residuals in OLSQ -- it can be used to construct a pseudo-F test of nested models. Note that it is zero for exactly identified models (if they have full rank). For two stage least squares, a test of overidentifying restrictions (@FOVERID) is also printed when the number of instruments is greater than the number of parameters. It is given by

\[ \frac{@PHI}{@S2^{(#inst-#params)}} \]

Following this is a table of parameter estimates and asymptotic standard errors, as well as their estimated variance-covariance (unless it has been suppressed). For each equation, LSQ prints a few goodness-of-fit statistics: the sum of squared residuals, standard error, mean and standard deviation of the dependent variable, and the Durbin-Watson statistic. The computation of these statistics is described in the regression output section of the User's Manual. If the equations are unnormalized, only the standard error, sum of squared residuals, and Durbin-Watson are printed.
Method

The method used by LSQ is a generalized Gauss-Newton method. The Gauss-Newton method is Newton's method applied to a sum of squares problem where advantage is taken of the fact that the squared residuals are very small near the minimum of the objective function. This enables the Hessian of the objective function to be well approximated by the outer product of the gradient of the equations of the model. "Generalized" refers to the fact that the objective function contains a fixed weighting matrix also, rather than being a simple sum of squares.

This implementation of the Gauss-Newton method in TSP uses analytic first derivatives of the model, which implies that the estimating equations must be differentiable in the parameters (TSP defines the derivatives of discontinuous functions like \textsc{SIGN}() to be zero, so this will always be true). The method is one of the simplest and fastest for well-behaved equations where the starting values of the parameters are reasonably good. When the equation is highly nonlinear, or the parameters are far away from the answers, this method often has numerical difficulties, since it is fundamentally based on the local properties of the function. These problems are usually indicated by numerical error messages from TSP; the program tries to continue executing for a while, but if things do not improve, the estimation will be terminated. When you encounter a problem like this, you can often get around it by estimating only a few parameters at a time to obtain better starting values. Use \textsc{CONST} to fix the others at reasonable values.

It is possible that a nonlinear model with cross-equation restrictions that estimates slowly using the default algorithm in LSQ will be estimated much faster if \textsc{FIML} with the \textsc{HITER=C} option is used instead. Iterated multivariate regression (LSQ without instruments, iterating on the covariance matrix) will produce answers identical to FIML; other LSQ methods such as \textsc{SUR} or \textsc{3SLS} produce estimates that will differ from FIML on the corresponding model. Asymptotically, however, they are equivalent.

For details on the estimation method, see the Berndt, Hall, Hall, and Hausman article.

Options

\texttt{COVU=} residual covariance matrix (same as the old \texttt{WNAME=} option below).

\texttt{DEBUG/NODEBUG} specifies whether detailed computations of the model and its derivatives are to be printed out at every iteration. This option produces extremely voluminous output and is not recommended for use except by systems programmers maintaining TSP.
FEI/NOFEI specifies that models with additive individual fixed effects are to be estimated. The panel structure must have been defined previously with the `FREQ (PANEL)` command. The equations specified must be linear in the parameters (this will be checked) and variables. Individual-specific means will be removed from both variables and instruments.

`INST` (list of instrumental variables). If this option is included, the LSQ estimator becomes nonlinear two stage least squares or nonlinear IV (if there is one equation) and nonlinear three stage least squares (if there is more than one equation). The list of instrumental variables supplied is used for all the equations. See the `INST` section of this manual and the references for further information on the choice of instruments.

ITERU/NOITERU specifies iteration on the COVU matrix; provides the same function as the old `MAXITW=` option.

`MAXITW=` the number of iterations to be performed on the parameters of the residual covariance matrix estimate. If `MAXITW` is zero the covariance matrix of the residuals is held fixed at the initial estimate (which is specified by `WNAME`). This option can be used to obtain estimates that are invariant to which equation is dropped in a shares model like translog.

`HCOMEGA=` `BLOCK/DIAGONAL` specifies whether standard errors robust to heteroskedasticity and serial correlation within group (panel-robust) are to be computed. The `BLOCK` option specifies that the variance-covariance matrix of the residuals has the block-diagonal structure (as opposed to a simple `DIAGONAL` covariance matrix of the residuals). When `FREQ (PANEL)` is in effect and the `ROBUST` or `HCOV=R` options are used in LSQ, the default option is `HCOMEGA=BLOCK`. This option is also available for `OLSQ`, `2SLS`, and `PANEL`.

`HETERO/NOHETERO` causes heteroskedastic-consistent standard errors to be used. See the `GMM (NMA=)` command for autocorrelation-consistent standard errors. Same as the old `ROBUST` option, or `HCOV=R`.

`WNAME=` the name of a matrix to be used as the starting value of the covariance matrix of the residuals.

`WNAME=OWN` specifies that the initial covariance matrix of the residuals is to be obtained from the residuals corresponding to the initial parameter values. If neither form of `WNAME=` is used, the initial covariance matrix is an identity matrix.

Nonlinear options control the iteration methods and printing. They are explained in the `NONLINEAR` section of this manual. Some of the common options are `MAXIT`, `MAXSQZ`, `PRINT/NOPRINT`, and `SILENT/NOSILENT`. 
The legal choices for \texttt{HITER=} are \texttt{G} (Gauss, the default) and \texttt{D} (Davidon-Fletcher-Powell). \texttt{HCOV=G} is the default method for calculating standard errors; \texttt{R} (Robust) and \texttt{D} are the only other valid options, although \texttt{D} is not recommended.

**Examples**

Assume that the following equations have been specified for the illustrative model of the U.S. economy:

\begin{verbatim}
FRML CONSEQ CONS = A+B*GNP ;
FRML INVEQ I = LAMBDA*I(-1) + ALPHA*GNP/(DELTA+R) ;
FRML INTRSTEQ R = D + F*(LOG(GNP)+LP-LM) ;
FRML PRICEQ LP = LP(-1) + PSI*(LP(-1)-LP(-2)) + PHI*LOG(GNP) +
    TREND*TIME+P0 ;
PARAM A B LAMBDA ALPHA D F PSI PHI TREND P0 ;
CONST DELTA 15 ;
\end{verbatim}

The model as specified has four equations: the parameters to be estimated are A, B, LAMBDA, ALPHA, D, F, PSI, PHI, TREND, and P0. There are 7 variables in the model, CONS, GNP, I, R, LP, LM, and TIME, and one additional instrument, G. To estimate the investment equation by nonlinear least squares, use the following command:

\texttt{LSQ (NOPRINT,TOL=.0001) INVEQ ;}

We can obtain multivariate regression estimates of the whole model with the following command, although these estimates are probably not consistent due to the simultaneity of the model (there are endogenous variables on the right hand side of the equations):

\texttt{LSQ (MAXIT=50) CONSEQ INVEQ PRICEQ INTRSTEQ ;}

The example below obtains three stage least squares estimates of the model, using a weighting matrix based on the starting values of the parameters (which are obtained by nonlinear two stage least squares):

\texttt{LSQ (INST=(C,LM,G,TIME)) CONSEQ ;}
\texttt{LSQ (INST=(C,LM,G,TIME)) INVEQ ;}
\texttt{LSQ (INST=(C,LM,G,TIME)) INTRSTEQ ;}
\texttt{LSQ (INST=(C,LM,G,TIME)) PRICEQ ;}
\texttt{LSQ (INST=(C,LM,G,TIME),WNAME=OWN) CONSEQ, INVEQ,}
\texttt{ INTRSTEQ, PRICEQ ;}

You can get the same three stage least squares estimates without the intermediate two stage least squares printout by using this command:

\begin{verbatim}
LSQ (INST=(C,LM,G,TIME)) CONSEQ
\end{verbatim}
3SLS (INST=(C,LM,G,TIME)) CONSEQ, INVEQ, INTRSTEQ, PRICEQ ;

See the description of the LIST command for an example of using cross-equation restrictions

References


MATRIX

Examples

MATRIX processes matrix algebra expressions. Operations on matrices are specified in matrix equations preceded by the word MAT; these equations are just like the variable transformations performed by GENR, except for two things: they do not operate under control of the current SMPL and the results are stored as a matrix. The MAT procedure checks the matrices for conformability of the operations and gives an error message if the operation specified is not possible. Often printing the matrices in question will reveal why the operation cannot be performed.

MATRIX <matrix name> = <matrix equation> ;

Usage

All the ordinary operators and functions used in TSP equations can also be used in the MAT command. They operate on an element-by-element basis (and hence require conforming matrices if they are binary operators). There is one important exception to this, the multiply operator *. For simplicity, this operator denotes the usual matrix multiplication, and element-by-element multiplication (the Hadamard product) is denoted by the operator %.

In the descriptions of the matrix operators that follow, we use the following symbols to denote the inputs and outputs of operations:

<table>
<thead>
<tr>
<th>Symbol</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>s</td>
<td>scalar or subscripted variable</td>
</tr>
<tr>
<td>i</td>
<td>integer scalar</td>
</tr>
<tr>
<td>m</td>
<td>any matrix (if scalar, treated as 1 by 1 matrix)</td>
</tr>
<tr>
<td>q</td>
<td>square matrix, N by N</td>
</tr>
<tr>
<td>sm</td>
<td>symmetric matrix, assumed positive semi-definite</td>
</tr>
<tr>
<td>dm</td>
<td>diagonal matrix, assumed positive semi-definite</td>
</tr>
<tr>
<td>tm</td>
<td>upper-triangular matrix, assumed positive semi-definite</td>
</tr>
<tr>
<td>v</td>
<td>column vector, N by 1</td>
</tr>
</tbody>
</table>

Here are the symbolic operators understood by the MAT command (in addition to the ordinary operators used also in GENR). Remember that the operands must be conformable for the operations that you request; TSP will check the dimensions for you and refuse to perform the computation if this condition is violated.

<table>
<thead>
<tr>
<th>Operator</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>m = m*m</td>
<td>matrix product</td>
</tr>
<tr>
<td>$m = m * s$ or $s * m$</td>
<td>scalar multiplication</td>
</tr>
<tr>
<td>------------------------</td>
<td>-----------------------</td>
</tr>
<tr>
<td>$m = m'$</td>
<td>matrix transpose</td>
</tr>
<tr>
<td>$m = m' m$</td>
<td>matrix transpose with implied matrix product</td>
</tr>
<tr>
<td>$m = q m''$</td>
<td>matrix inverse</td>
</tr>
<tr>
<td>$m = q m'' m$</td>
<td>matrix inverse with implied matrix product</td>
</tr>
<tr>
<td>$m = m # m$</td>
<td>Kronecker product</td>
</tr>
<tr>
<td>$m = m % m$</td>
<td>Hadamard product (element by element)</td>
</tr>
</tbody>
</table>

When TSP processes a MAT command, it recognizes several operations where great savings of computation time can be made by eliminating duplicate calculations. These situations include, but are not limited to, the cross-product operation (which generates a symmetric matrix) and the calculation of a quadratic form (the expression $A^T B A'$). This occurs even when the arguments to these expressions are complicated expressions themselves. Thus, you should be careful to express any such complex arguments the same way whenever they appear in the matrix expression.

The following functions take matrices as their input and produce scalars as output. They may be used anywhere in a MAT statement where scalars are allowed, keeping in mind that a scalar is also a 1 by 1 matrix.

<table>
<thead>
<tr>
<th>$s = DET(qm)$</th>
<th>determinant (truncated to zero when &lt; 1.E-37)</th>
</tr>
</thead>
<tbody>
<tr>
<td>$s = LOGDET(qm)$</td>
<td>log of (positive) determinant, no truncation</td>
</tr>
<tr>
<td>$s = TR(qm)$</td>
<td>trace (sum of diagonal elements)</td>
</tr>
<tr>
<td>$s = MIN(m)$</td>
<td>element with minimum value</td>
</tr>
<tr>
<td>$s = MAX(m)$</td>
<td>element with maximum value</td>
</tr>
<tr>
<td>$s = SUM(m)$</td>
<td>sum of elements</td>
</tr>
<tr>
<td>$i = NROW(m)$</td>
<td>number of rows</td>
</tr>
<tr>
<td>$i = NCOL(m)$</td>
<td>number of columns</td>
</tr>
<tr>
<td>$i = RANK(m)$</td>
<td>rank (number of linearly independent columns or rows)</td>
</tr>
</tbody>
</table>

The following functions are matrix-to-matrix; that is, they take a matrix, perform some computation on it, and produce another matrix as output. They can be used anywhere in a MAT equation.

<table>
<thead>
<tr>
<th>$tm = CHOL(sm)$</th>
<th>Choleski factorization (matrix square root)</th>
</tr>
</thead>
<tbody>
<tr>
<td>$sm = YINV(sm)$</td>
<td>Positive semi-definite inverse via CHOL()</td>
</tr>
<tr>
<td>$dm = IDENT(i)$</td>
<td>Creates an identity matrix of order $i$</td>
</tr>
<tr>
<td>$v = EIGVAL(qm)$</td>
<td>Computes the vector of eigenvalues of $qm$. If $qm$ is not symmetric positive semi-definite, the imaginary parts of the eigenvalues are stored as</td>
</tr>
</tbody>
</table>
### MATRIX

<table>
<thead>
<tr>
<th>Function</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>( qm )</td>
<td>Computes the matrix of eigenvectors (columns). If ( qm ) is not symmetric positive semi-definite, the imaginary parts of the eigenvectors are stored as ( @EIGVECI ).</td>
</tr>
<tr>
<td>( v = VEC(m) )</td>
<td>Creates a vector of all the elements of ( m ), column by column. ( qm ): ( N \times N ) elements, ( sm, tm ): ( N \times (N+1)/2 ) elements. ( dm ): ( N ) elements.</td>
</tr>
<tr>
<td>( v = VECH(m) )</td>
<td>Creates a vector of all the unique elements of ( m ), column by column. ( qm ): ( N \times N ) elements, ( sm, tm ): ( N \times (N+1)/2 ) elements. ( dm ): ( N ) elements.</td>
</tr>
<tr>
<td>( dm = DIAG(m) )</td>
<td>Creates a diagonal matrix from a matrix. ( qm, sm, tm ): take the diagonal from input matrix; ( v ): convert the vector to a diagonal matrix; ( s ): illegal, use ( s \times @IDENT(i) ) to create a diagonal matrix with ( s ) on the diagonal.</td>
</tr>
<tr>
<td>( sm = SYM(qm) )</td>
<td>Creates a symmetric matrix from a square matrix (the upper triangular elements are ignored).</td>
</tr>
<tr>
<td>( m = GEN(qm) )</td>
<td>Creates a general matrix from a symmetric or diagonal matrix.</td>
</tr>
<tr>
<td>( \text{series} = \text{SER}(v) )</td>
<td>Converts a vector to a series under control of ( \text{SMPL} ).</td>
</tr>
</tbody>
</table>

### Output

MATRIX produces no printed output. Typically, one matrix is stored in data storage.

### Examples

**MAT** \( B = (X'X)^{-1}X'Y; \)

produces OLS regression coefficients (not a very accurate way to do this).

The example below computes the Eicker-White estimate of the variance-covariance of the estimated coefficients after a regression:

**OLSQ Y C X ;**
**MAKE XMAT C X ;**
**MAT XXI = (XMAT ' XMAT) '' ;**
**MAT VCOV = (XMAT'XXI) ' DIAG(@RES''2) * (XMAT'XXI) ;**
MFORM

Options  Examples

MFORM forms or reforms matrices used to make a matrix from a series or vector or to change the dimensions or type of an existing matrix. The matrix may be transposed as it is formed. Matrices can be renamed or copied without reformattting with the RENAME or COPY commands.

MFORM (NROW=<# of rows in matrix>, NCOL=<# of columns in matrix>, TRANS, TYPE = GENERAL or SYMMETRIC or TRIANG or DIAG) <variable name> or <new matrix> = <old variable> or <old variable> new matrix> or <new matrix> = <scalar> ;

or
MFORM (BAND,NROW=nrows) <new matrix> = <band_vector> [<corner matrix>] ;

or
MFORM (BLOCK) <new matrix> = <list of matrices> ;

Usage

If there is only one argument, either an existing matrix or series is transformed in place, or a new matrix is initialized to zero. The options specify the characteristics of the new matrix: the number of rows and columns, the type and whether it is to be transposed as it is formed.

When there is more than one argument, the old variable may be a series, a matrix, a vector, or a scalar. The new variable will be a matrix of the type specified on the command. If no type is specified, a general matrix is created. If the input variable is a series, it is retrieved under control of SMPL and only those observations in the current sample are placed in the matrix. If the input series or matrix is longer than NROW*NCOL, it is truncated (except in the case of the DIAG type - see the examples). If it is shorter, an error message is given, unless it is a scalar, in which case it is duplicated throughout the new matrix.

If you specify a type such as triangular which is not consistent with the input matrix, MFORM will change the input so that it is, i.e., the elements below the diagonal will be zeroed. You can use this feature, for example, to select the diagonal elements of a matrix and form a new matrix from them (like a diag function). UNMAKE can be used to form a series from the diagonal of a diagonal matrix.

Output

MFORM produces no printed output. A single matrix is stored in data storage.
Options

BAND/NOBAND specifies that a (symmetric) band matrix is to be formed from a vector of band values, and optionally a symmetric corner matrix (for the upper left and lower right corners, to override the band values).

BLOCK/NOBLOCK specifies that a block diagonal matrix is to be formed from a list of symmetric or general matrices (by placing them along the diagonal, and putting zeros elsewhere). This is useful for composing VCOV matrices from several independent estimations for minimum distance estimation with LSQ (or for ANALYZ).

NROW= the number of rows in the matrix to be formed. This is required for a general matrix.

NCOL= the number of columns in the matrix to be formed. This is required for a general matrix.

Either NROW or NCOL must be specified for symmetric, triangular, or diagonal matrices, unless the input variable is a matrix and you wish the new matrix to have the same dimensions.

TRANS/NOTRANS specifies whether the input variable is to be transposed to produce a matrix of the type and dimensions specified. If the input variable is a matrix, the output matrix will have the number of rows equal to the number of columns of input and the number of columns equal to the number of rows of input. MFORM(TRANS) is the same as the MATRAN command.

TYPE=GENERAL or SYMMETRIC or TRIANG or DIAG specifies the type of the new matrix. GENERAL, the default, may be used for any rectangular or square matrix. SYMMETRIC implies that the matrix is equal to its transpose; only the lower triangle will be stored internally to save space. TRIANG implies that the matrix is upper triangular (has zeroes below the diagonal). DIAG means a matrix whose off-diagonal elements are zero. Only the diagonal is stored, and it is expanded before use. A series may be used as input to MFORM(DIAG).

Examples

Suppose that we have a nine observation series called X containing the values 10,20,30,40,50,60,70,80,90. Each of the following MFORM commands will yield a different result:

SMPL 1,9;
MFORM (TYPE=GENERAL,NROW=3,NCOL=3) X ;
yields $X = \begin{bmatrix} 10 & 40 & 70 \\ 20 & 50 & 80 \\ 30 & 60 & 90 \end{bmatrix}$

**SMPL 1.6;**

*MFORM* *(TYPE=GENERAL,NROW=2,NCOL=3)* \(X \) \(XMAT \);  

yields $XMAT = \begin{bmatrix} 10 & 30 & 50 \\ 20 & 40 & 60 \end{bmatrix}$

*MFORM*(TRANS) \(XMAT=XMATT\); 

yields $XMATT = \begin{bmatrix} 10 & 20 \\ 30 & 40 \\ 50 & 60 \end{bmatrix}$

*MFORM*(TYPE=GENERAL,NROW=4,NCOL=3) \(X \);  
gives an error message; the resulting matrix is too big for the amount of data available.

**SMPL 1.9;**

*MFORM* *(TYPE=SYM,NROW=3)* \(XSYM=X \);  

yields $XSYM = \begin{bmatrix} 10 & 20 & 30 \\ 20 & 50 & 60 \\ 30 & 60 & 90 \end{bmatrix}$

*MFORM* *(TYPE=TRIANG,NCOL=3)* \(X \);  

yields $X = \begin{bmatrix} 10 & 40 & 70 \\ 0 & 50 & 80 \\ 0 & 0 & 90 \end{bmatrix}$

*MFORM* *(TYPE=DIAG,NCOL=3)* \(X \);
MFORM

yields X =

\[
\begin{bmatrix}
10 & 0 & 0 \\
0 & 50 & 0 \\
0 & 0 & 90 \\
\end{bmatrix}
\]

\text{MFORM (TYPE=diag,NROW=9) X ;}
\text{or}
\text{MAT X = DIAG(X);} \\

yields X =

\[
\begin{bmatrix}
10 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\
0 & 20 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\
0 & 0 & 30 & 0 & 0 & 0 & 0 & 0 & 0 \\
0 & 0 & 0 & 40 & 0 & 0 & 0 & 0 & 0 \\
0 & 0 & 0 & 0 & 50 & 0 & 0 & 0 & 0 \\
0 & 0 & 0 & 0 & 0 & 60 & 0 & 0 & 0 \\
0 & 0 & 0 & 0 & 0 & 0 & 70 & 0 & 0 \\
0 & 0 & 0 & 0 & 0 & 0 & 0 & 80 & 0 \\
0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 90 \\
\end{bmatrix}
\]

The next example shows how you can make a diagonal matrix from a column vector - if the input series or matrix is too short to make a diagonal matrix by selecting diagonal elements, the whole vector becomes the diagonal.

\text{MFORM (TYPE=diag,NCOL=3) X = 2;}
\text{or}
\text{MAT X = 2*IDENT(3);} \\

yields X =

\[
\begin{bmatrix}
2 & 0 & 0 \\
0 & 2 & 0 \\
0 & 0 & 2 \\
\end{bmatrix}
\]

A band matrix:

\text{MMAKE BVEC 2 -1;}
\text{READ(NROW=2,TYPE=SYM) CORNER;}
11 21 22;
\text{MFORM(BAND,NROW=5) B5 = BVEC CORNER;}

yields B5 =

\text{262}
<table>
<thead>
<tr>
<th></th>
<th>11</th>
<th>21</th>
<th>0</th>
<th>0</th>
<th>0</th>
</tr>
</thead>
<tbody>
<tr>
<td>21</td>
<td>22</td>
<td>-1</td>
<td>0</td>
<td>0</td>
<td></td>
</tr>
<tr>
<td>0</td>
<td>-1</td>
<td>2</td>
<td>-1</td>
<td>0</td>
<td></td>
</tr>
<tr>
<td>0</td>
<td>0</td>
<td>-1</td>
<td>22</td>
<td>21</td>
<td></td>
</tr>
<tr>
<td>0</td>
<td>0</td>
<td>0</td>
<td>21</td>
<td>11</td>
<td></td>
</tr>
</tbody>
</table>
ML

Output Options Examples References

ML is a general purpose maximum likelihood estimation procedure. It can be used to estimate the parameters of any (identified) model for which you can write down the logarithm of the likelihood in a TSP equation (FRML), or evaluate the log likelihood in a procedure (PROC).

ML (nonlinear options) <log likelihood equation name> ;
or
ML (nonlinear options) <procedure name> <list of parameters> ;

Usage

FRML method. Usually the simplest approach is to write the log likelihood equation in a FRML with LOGL as the dependent variable. Note that this equation is for each observation in the current SMPL vector. If there are different equations, depending on different cases, write the equation as the sum of the individual equations, with dummy variables multiplying each equation to select the appropriate one for any given observation. Often the "case" will be determined by a dependent discrete choice variable, and observations are usually i.i.d., but the likelihood function could be made different for different parts of the SMPL by using more general time-dependent dummy variables. Of course, the log likelihood must be additively separable over the sample for this method to work (if it is not, use the PROC method described below).

Use a PARAM statement to specify which of the variables in the LOGL equation are to be estimated and supply their starting values if desired. Follow this by an ML command with any of the standard NONLINEAR options and the name of the equation which specifies the likelihood function. ML will maximize this function with respect to the parameters using a standard gradient method; the exact form of the Hessian approximation used as a weighting matrix depends on the HITER option. The default is to use the BHHH method, a method of scoring, but with the sample covariance of the gradient of the likelihood used in place of its expectation.

Good Applications for the FRML method:

1. Truncation models involving CNORM(), such as two-limit Tobit.
2. Nonlinear equations for PROBIT, TOBIT, LOGIT, etc. This includes parameter restrictions and holding parameters fixed.
3. Checking your own second derivatives when you are writing a
maximization procedure for any program that uses natural language
equations.

4. Robust models like $\text{LOGL} = \text{ABS}(Y - XB)$.

5. Minimization problems (just negate the equation).

6. General maximum likelihood problems, using any functions
recognized by TSP (including SQRT, POS, and the gamma
(factorial) function, which can be used for the gamma, chi-squared,
beta, t, and F densities). Even complicated likelihood functions on
large datasets may be estimable using ML. Even though more
computer time may be required using TSP instead of a custom
program, programming time is considerably reduced (and the
relative price of CPU time is usually small and shrinking).
Maximization with analytic derivatives is usually much faster than
with numeric derivatives (the method which used to be lowest in
programming cost). See Timing example below.

**PROC method.** Sometimes it is extremely difficult, or impossible to write
down the log likelihood in a single **FRML** (even with use of **EQSUB**). See the
list below for some examples. For this form of the **ML** command, write a
**PROC** which evaluates the log likelihood and stores it in $@\text{LOGL}$. Give the
name of this PROC as the first argument (after any options) of the **ML**
command, and follow it with a list of **PARAMs** which are to be estimated.
Write the **PROC** so that it starts by checking any constraints on the
**PARAMs. If any constraint is violated, set $@\text{LOGL}$ to $@\text{MISS}$ before exiting
from the **PROC.** **OPTIONS DOUBLE**; is advised if you want to use double
precision to form intermediate results such as residuals.

The main disadvantage of using the **PROC** instead of the **FRML** method is
that analytic derivatives are not available. However, numeric derivatives (the
default **HITER=F** and **GRAD=C2**) will often be quite adequate. A slight
disadvantage is that you have to explicitly list the **PARAMs** to be estimated in
the command line. **HCOV=U** (numeric second derivatives) is the default
method of computing the standard errors for **MLPROC.** For iteration,
**HITER=F** is the default, but **HITER=U** can be chosen as an option.

Good Applications for the **PROC** method:

1. Time series models like **ARMA** and **GARCH**, where the equations
are recursive (depend on residuals or variance from the previous
time period(s). Models which can be evaluated by the **KALMAN**
command also fit into this category (**ML** thus allows estimation of the
hyperparameters).
2. Multi-equation models like FIML. These involve Jacobians, matrix inverses, and determinants, which would have to be written into the log likelihood equation by hand (very difficult for more than about 4 equations unless the Jacobian is sparse).

3. Models which require several diverse commands to evaluate, such as multivariate normal integrals via simulation, or other functions that are not built in to TSP. Another example in this class is a concentrated log likelihood function (the FRML method can only handle the unconcentrated log likelihood, which is usually more nonlinear and often harder to write).

4. Models with complicated constraints. A good example would be ARCH models, where the conditional variance must be positive for every observation.

Bad Applications for either method:

1. Existing linear models in TSP (PROBIT, TOBIT, LOGIT, SAMPSEL). The regular TSP commands are more efficient, more resistant to numerical problems, often have better starting values, and provide model-specific statistics. See Timing example below.

To give an idea of how much this convenience costs in terms of CPU time, here is a timing example run on the VAX 11/780 of a Probit on 385 observations, 8 variables.

<table>
<thead>
<tr>
<th>CPU seconds</th>
<th>Method</th>
<th>Procedure</th>
</tr>
</thead>
<tbody>
<tr>
<td>5.24</td>
<td>canned Probit procedure</td>
<td>PROBIT command</td>
</tr>
<tr>
<td>65.65</td>
<td>BHHH algorithm (method of scoring)</td>
<td>ML (HITER=B, HCOV=N)</td>
</tr>
<tr>
<td>75.97</td>
<td>Newton's method (uses 2nd derivatives)</td>
<td>ML (HITER=N, HCOV=N)</td>
</tr>
</tbody>
</table>

The moral is that ML should not be used when you have a Fortran-coded alternative estimation program, but could be useful if you don't want to spend your time developing such a program. Also, in this case, the method of scoring was somewhat faster than Newton's method, although the latter is more powerful (it takes fewer iterations).

Tips:
1. Write the equation carefully to avoid things like Log (x < =0) or Exp(x >88). These are fatal errors if they happen in the first function evaluation (using the starting values). They are not fatal during iterations (the program automatically uses a smaller stepsize), but they can be inefficient. Often these problems can be avoided by reparametrizing the likelihood function. The standard example of this is estimating SIGMA (or SIGMA-inverse) instead of SIGMA-squared. If you are getting numerical errors and you can't rewrite the likelihood function, try using SELECT to remove the problem observations. After you get convergence, use the converged values as starting values and reestimate using the full sample.

2. Choose starting values carefully (see previous).

3. Use EQSUB for less work rewriting equations and more efficient code.

4. The "Working space =" message gives an indication of the length of the derivative code.

5. If the second derivative matrix is singular, you may have sign errors in the log likelihood function (the inversion routine assumes the second derivative matrix is negative definite).

6. If you are using derivatives, make sure the functions you are using are differentiable. Logical operations are not differentiable everywhere, although they are differentiable at all but a finite number of points. TSP will do the best it can with them, but if you end up on a kink (corner), it may stall.

Output

The following results are stored:

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@LOGL</td>
<td>scalar</td>
<td>1</td>
<td>Log of likelihood function</td>
</tr>
<tr>
<td>@IFCONV</td>
<td>scalar</td>
<td>1</td>
<td>= 1 if convergence achieved, 0 otherwise</td>
</tr>
<tr>
<td>@NCOEF</td>
<td>scalar</td>
<td>1</td>
<td>Number of parameters to be estimated</td>
</tr>
<tr>
<td>@NCID</td>
<td>scalar</td>
<td>1</td>
<td>Number of identified parameters</td>
</tr>
<tr>
<td>@RNMS</td>
<td>list</td>
<td>params</td>
<td>Names of right hand side variables</td>
</tr>
<tr>
<td>@COEF</td>
<td>vector</td>
<td>params</td>
<td>Coefficient estimates</td>
</tr>
<tr>
<td>@GRAD</td>
<td>vector</td>
<td>params</td>
<td>Gradient of the log likelihood at convergence</td>
</tr>
<tr>
<td>@SES</td>
<td>vector</td>
<td>params</td>
<td>Standard errors</td>
</tr>
</tbody>
</table>

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ML

@T vector #params T-statistics
@VCOV matrix #params* Variance-covariance of estimated
t- statistics #params coefficients.

See the NONLINEAR section for the alternative names of VCOV stored
when the HCOV option is used.

Method

The method used is a standard gradient method, explained in somewhat
more detail in Chapter 9 of the User's Manual. Briefly, at each iteration, a
new parameter vector is computed by moving in the direction specified by
the gradient of the likelihood (uphill), weighting this gradient by an
approximation to the matrix of second derivatives at that point (in order to
adjust for the curvature). Convergence is declared when the changes in the
parameters are all "small", where small is defined by the TOL= option.

The ML procedure normally uses analytic first (and second) derivatives (for
the FRML method). The function can also be maximized numerically
(HITER=F or HITER=D). See NONLINEAR for more information on the
options (HCOV=N gives standard errors based on analytic second
derivatives). MLPROC uses HCOV=U (numeric second derivatives) to
compute the standard errors.

Options

Standard nonlinear options (see NONLINEAR section). HITER=B, HCOV=B
is the default for the FRML method; HITER=F, HCOV=U, GRAD=C2 is the
default for the PROC method. Starting values are from the PARAM and SET
statements. Note that the CONST command allows fixing parameters during
an estimation (for the FRML method).

Examples

FRML method. For example, in the Probit model, the likelihood is CNORM(-
XB) for Y<=0, and (1-CNORM(-XB)) for Y>0. This could be written in the
following way (see the User's Guide for alternate coding and many more
examples):

GENR Y0 = Y<=0 ;
GENR Y1 = Y>0 ;
FRML EQ1 LOGL = LOG (Y0*CNORM(-XB) + Y1*(1-CNORM(-XB)));

The XB expressions can be filled in later with the EQSUB command. Note
that this allows for nonlinear equations, as in this example:

FRML NLXB XB = B0 + B1*X1 + (B2/B1)*X2;

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**PROC method.** Here is a simple concentrated log likelihood function, where we estimate the mean of a time trend, and concentrate out the variance parameter to reduce the nonlinearity of the function.

? make sure that residuals are stored in double precision
OPTIONS DOUBLE;
SMPL 1,9;
TREND T;
PARAM MT,2 ;
ML NRMLC MT ; ? PROC form of the ML command
PROC NRMLC;
    E = T - MT; ? residual
    MAT SIG2 = (E'E)/@NOB; ? sigma-squared (variance)
    SET PI = 4*ATAN(1); ? tan(pi/4) = 1
    SET @LOGL = -(@NOB/2)*( LOG(SIG2) + 1 + LOG(2*PI) );
ENDPROC;

References


MMAKE

MMAKE makes a new matrix by stacking a set of series or matrices, or a new vector from a set of scalars. In the series case, the new matrix normally has the number of rows equal to the number of observations and the number of columns equal to the number of series. In the matrix case, the new matrix has the number of rows equal to the common number of rows of the matrices and the number of columns equal to the total number of columns. The vector has the number of rows equal to the number of scalars.

MMAKE is the reverse of UNMAKE, which breaks a matrix into a set of series, or a vector into a set of scalars. To make a matrix from another matrix or vector by changing its type, dimensions, or transposing it, use the MFORM procedure.

**MMAKE (VERT) <matrix name> <list of series> ;**

or

**MMAKE <vector name> <list of scalars> ;**

or

**MMAKE (VERT) <matrix name> <list of matrices> ;**

Usage

MMAKE's first argument is the name to be given to the new matrix, followed by a list of series or matrices which will form the columns of the new matrix. The number of series is limited only by the maximum size of the argument list (usually 2000 or more) and the space available in data storage for the new matrix.

Only the observations in the series which are within the current sample will be used, so you can select data for the matrix very conveniently by changing the SMPL. If you use matrices instead of series, the SMPL is ignored.

The matrix made by MMAKE will always be a general matrix and (when using series) its dimensions are normally NROW = @NOB (the number of observations) by NCOL = the number of input series. To change its type, use MFORM.

If the second and following arguments to MMAKE are scalars (CONSTs, PARAMs, or numbers), a vector will be created instead of a matrix. This is useful for creating vectors like @START or @COEF from estimated parameters or scalars created by UNMAKE from previous @COEF vectors.

Output
MMAKE produces no printed output. A single matrix or vector is stored in data storage.

Options

VERT/NOVERT stacks the input series or matrices vertically instead of horizontally.

Examples

If the current sample is SMPL 1 4 ; and there are two data series X1 = (1,2,3,4) and X2 = (9 8 7 6), the following command:

\[ \text{MMAKE } X \ X1 \ X2 ; \]

results in the matrix \( X = \)

\[
\begin{array}{ll}
1 & 9 \\
2 & 8 \\
3 & 7 \\
4 & 6 \\
\end{array}
\]

whereas

\[ \text{MMAKE (VERT) XV X1 X2;} \]

would create the 8 x 1 vector \( XV : \)

\[
\begin{array}{ll}
1 \\
2 \\
3 \\
4 \\
9 \\
8 \\
7 \\
6 \\
\end{array}
\]

Here is an example of adding a coefficient to a LOGIT estimation while retaining the starting values for the other coefficients from the previous estimation:

\[ \text{LOGIT(NCHOIC=2) WORK C SCHOOL EXPER RACE;} \]
\[ \text{UNMAKE @COEF C1-C4;} \]
\[ \text{MMAKE @START C1-C4 0;} \]
\[ \text{LOGIT WORK C SCHOOL EXPER RACE MSTAT;} \]

This example makes a matrix of regression output to be written to disk:
The final example creates the partitioned matrix $A = \begin{bmatrix} D & E & F \\ G & H & I \end{bmatrix}$

where $D$, $E$, and $F$ are matrices with the same number of rows, $D$ and $G$ have the same number of columns, etc.

```plaintext
MMAKE DEF D E F ;
MMAKE GHI G H I ;
MMAKE (VERT) A DEF GHI ;
```
MODEL

**Output**

MODEL determines the order in which the equations of a model should be solved and saves this order under a collected model name. It must be used before a `SOLVE` command invokes the model simulation procedure.

**MODEL (DONGALLO, FILE=<filename>, PRINT, SILENT) <equation list> [<endogenous variable list>] <ordered model name>;**

**Usage**

MODEL takes as its arguments the name of a list of equations in the model, and produces a collected and ordered model which is stored under the name supplied by the user. Each of the endogenous variables in the list must appear on the left hand side of one and only one of the equations. For compatibility with older versions of TSP, you may supply the endogenous variable list, but for the current version this is optional.

**Output**

MODEL prints information about the ordering of the model, the number of simultaneous and recursive blocks, and whether or not the simultaneous blocks are linear in the variables for which they are to be solved. This information is also stored as part of the collected model. A table is printed which shows for each equation in the model the number of its block, whether the block is simultaneous (S) or recursive (R), and which endogenous variables appear in that equation.

See the example output in the *TSP User's Guide*.

**Options**

**DONGALLO/NODONGAL** specifies that each simultaneous block should be ordered for a near-minimal feedback set. It prints an F next to the feedback variables to identify them, and a summary of the blocks. This ordering is sometimes useful when the Gauss-Seidel method is used to `SOLVE` the model. See the Don and Gallo reference for further details.

**FILE=filename** writes a file containing input for the CAUSOR program. CAUSOR provides detailed information on model structure, such as essential feedback sets. When `FILE=` is used, the equations do not have to be uniquely normalized, as long as the endogenous variable list is supplied. CAUSOR may be obtained from Manfred Gilli, Departement d'Econometrie, Universite de Geneve.
PRINT/NOPRINT specifies whether the older (more voluminous) output format is to be used.

SILENT/NOSILENT suppresses printing completely.

Examples

This example shows how to set up the well-known Klein Model I for simulation:

\[
\text{INST } CX \ C \ W \ P(-1) \ \text{INVR } C \ P(-1) \ K(-1) \ E(-1) \ TM \ W2 \ G \ TX ; \\
\text{FORM CONS ;} \\
\text{INST } I \ C \ P \ P(-1) \ K(-1) \ \text{INVR } C \ P(-1) \ K(-1) \ E(-1) \ TM \ W2 \ G \ TX ; \\
\text{FORM INV ;} \\
\text{INST } W1 \ C \ E(-1) \ TM \ \text{INVR } C \ P(-1) \ K(-1) \ E(-1) \ TM \ W2 \ G \ TX ; \\
\text{FORM WAGES ;} \\
\text{IDENT WAGE } W = W1+W2 ; \\
\text{IDENT BALANCE } E=E+CX+I+(TX+W+P) ; \\
\text{IDENT PPROD } P = E-TX-W1 ; \\
\text{IDENT CAPSTK } K=K(-1)+I ; \\
\text{LIST KLEIN CONS WAGES BALANCE PPROD INV WAGE CAPSTK ;} \\
\text{MODEL KLEIN KLEINC ;} \\
\text{SOLVE (TAG=S,TOL=.0001,METHOD=FLPOW) KLEINC ;}
\]

This model solves for CX (consumption), I (investment), W1 (wages in the private sector), W (total wage bill), E (production of the private sector), P (profits), and K (capital stock) using TM (time), W2 (government wage bill), TX (taxes), and G (government expenditures) as exogenous variables. At the end of this simulation, the solved variables are stored under the names CXS, IS, etc.

References


Gilli, Manfred, "Graph-theory based tools in the practice of macroeconometric modeling," in Methods and Applications of Economic Dynamics, S. K. Kuipers, L. Schoonbeek, and E. Sterken (eds), North Holland, Amsterdam.

MSD

See Also  CORR/COVA

Output  Options  Examples  References

MSD produces a table of means, standard deviations, minima, maxima, sums, variances, skewness, and kurtosis for all the variables listed. Only observations in the current sample with no missing values are included. The variables may be weighted before the statistics are computed.

**MSD (ALL, BYVAR, CORR, COVA, MOMENT, PAIRWISE, PRINT, SILENT, TERSE, WEIGHT=<series name>) <list of series>**;

Usage

For univariate statistics on a set of variables, use the MSD command with no options. The CORR, MOMENT, and COVA options enable you to get several forms of descriptive statistics on the variables at once, saving on computation time. The option ALL allows you to obtain additional statistics such as the median. The TERSE option restricts the statistics computed in order save space and time.

Output

MSD stores the statistics which are requested as well as printing them.

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@NOBMSD</td>
<td>vector</td>
<td>#vars</td>
<td>Number of non-missing observations (for BYVAR).</td>
</tr>
<tr>
<td>@MEAN</td>
<td>vector</td>
<td>#vars</td>
<td>Means.</td>
</tr>
<tr>
<td>@STDDEV</td>
<td>vector</td>
<td>#vars</td>
<td>Standard Deviations.</td>
</tr>
<tr>
<td>@MIN</td>
<td>vectpr</td>
<td>#vars</td>
<td>Minimums.</td>
</tr>
<tr>
<td>@MAX</td>
<td>vector</td>
<td>#vars</td>
<td>Maximums.</td>
</tr>
<tr>
<td>@SUM</td>
<td>vector</td>
<td>#vars</td>
<td>Sums.</td>
</tr>
<tr>
<td>@VAR</td>
<td>vector</td>
<td>#vars</td>
<td>Variances.</td>
</tr>
<tr>
<td>@SKEW</td>
<td>vector</td>
<td>#vars</td>
<td>Skewness</td>
</tr>
<tr>
<td>@KURT</td>
<td>vector</td>
<td>#vars</td>
<td>Excess kurtosis</td>
</tr>
<tr>
<td>@MEDIAN</td>
<td>vector</td>
<td>#vars</td>
<td>Median (for ALL option).</td>
</tr>
<tr>
<td>@Q1</td>
<td>vector</td>
<td>#vars</td>
<td>1st quartile.</td>
</tr>
<tr>
<td>@Q3</td>
<td>vector</td>
<td>#vars</td>
<td>3rd quartile.</td>
</tr>
<tr>
<td>@IQR</td>
<td>vectotr</td>
<td>#vars</td>
<td>Inter-quartile range.</td>
</tr>
<tr>
<td>@CORR</td>
<td>matrix</td>
<td>#vars*#vars</td>
<td>Correlation matrix.</td>
</tr>
<tr>
<td>@COVA</td>
<td>matrix</td>
<td>#vars*#vars</td>
<td>Covariance matrix.</td>
</tr>
</tbody>
</table>
Method

The mean, minimum, maximum, variance, and standard deviation are computed in the usual way. The estimated variance and covariance are computed by small sample formulas (division by \( N-1 \) instead of \( N \)). The formulas for the skewness and kurtosis are the following:

\[
Skewness = \frac{N^2 M_3}{(N-1)(N-2)S^3}
\]

\[
Excess \ Kurtosis = \frac{N^2 (N+1)M_4 - 3(N-1)M_2^2}{(N-1)(N-2)(N-3)S^4}
\]

where \( M_3 \) and \( M_4 \) are the centered third and fourth moments and \( S \) is the estimated standard deviation. These statistics can be used to test for normality of the variables. The skewness multiplied by the square root of \( N/6 \) and the kurtosis multiplied by the square root of \( N/24 \) both have a normal(0,1) distribution under the null (when the mean and standard deviation have been estimated; see Davidson and MacKinnon for a derivation).

The median is the value of the series at the \( (N+1)/2 \) observation (after sorting from low to high). The first and third quartiles are the values of the series at the \( (N+1)/4 \) and \( (3N+3)/4 \) observations respectively and the interquartile range is the difference between these two values. If these observation numbers are not integers, the values are a weighted average of the bracketing observations.

Options

ALL/NOALL computes the median, first and third quartiles, and the interquartile range, in addition to the normal statistics. The median, etc. are computed using any weight that has been supplied.
BYVAR/NOBYVAR treats missing values for each series separately, so that the maximum possible number of observations for each series is used. NOBMSD will be stored in this case. Normally, if any series has missing values for any observation, that observation is dropped for all series.

CORR tells MSD to compute and print the correlation matrix of the variables.

COVA tells MSD to compute and print a covariance matrix.

MOMENT tells MSD to compute and print an uncentered moment matrix also. This matrix is divided by the number of observations with positive weights to scale it conveniently.

PAIRWISE/NOPAIRWISE treats missing values for each pair of series separately from other series. It applies to CORR, COVA and MOMENT matrices. NOBCOVA will be stored.

PRINT/NOPRINT specifies whether the results of the procedure are to be printed, or just stored in data storage.

SILENT/NOSILENT specifies that all printed output is to be suppressed.

TERSE/NOTERSE specifies that only the means, standard deviations, minima, and maxima are computed and printed. The sum, variance, skewness, and kurtosis are suppressed.

WEIGHT= the name of series which will be used to weight the observations. The data are multiplied by the square roots of the weighting series before the statistics are computed, so that the series should be proportional to the inverses of the variances of the variables. If the weight is zero for a particular observation, that observation is not included in the computations nor is it counted in determining degrees of freedom. The quartile estimates including the median are also weighted estimates.

Examples

LIST VARS PAT RND ASSETS DRND DPAT ;
MSD (CORR) VARS ;
MSD (CORR,COVA,WEIGHT=POP) INCOME PHONES NEWBUS ;

References


NAME

**Examples**

NAME, often the first statement in a TSP job, is used to supply a job or user name to be printed at the top of each page and, optionally, a title for the run.

**NAME <jobname> ['text string to be used as title'] ;**

**Usage**

The only required argument on the NAME statement is the jobname, which may be any descriptive name of up to 8 characters which you wish to give your job, or could be your name to distinguish your jobs from others if they are being run together.

The job title is optional, but recommended - if included, it will be printed at the top of each page of output until a TITLE statement is executed which replaces the title. The title is a string of up to 60 characters enclosed in quotes. There can be no quotes (" or ") imbedded in the title.

**Output**

NAME causes a jobname to be printed in the upper right hand corner of each page of TSP output. If a title is included on the command, the title is also printed at the top of every page in columns 21 through 80.

If the terminal (CRT) option is on, no paging of TSP output is done and no titles are printed unless requested by a PAGE command.

**Examples**

**NAME KARLMARX ;**
**NAME ILLUS44 'ILLUSTRATIVE MODEL FOR TSP VERSION 4.4' ;**
**NAME KLEINLSQ '3SLS ESTIMATES OF KLEIN MODEL I' ;**
NEGBIN

Output  Options  Example  References

NEGBIN obtains estimates of the Negative Binomial model, where the dependent variable takes on only nonnegative integer count values and its expectation is an exponential linear function of the independent variables. In the Negative Binomial model, the variance of the dependent variable is larger than the mean, in contrast to the Poisson model, where the variance equals the mean (see the POISSON procedure).

NEGBIN (MODEL=1 or 2, nonlinear options) <dependent variable> <list of independent variables> ;

Usage

The basic NEGBIN statement is like the OLSQ statement: first list the dependent variable and then the independent variables. If you wish to have an intercept term in the regression (usually recommended), include the special variable C or CONSTANT in your list of independent variables. You may have as many independent variables as you like subject to the overall limits on the number of arguments per statement and the amount of working space, as well as the number of data observations you have available.

The observations over which the regression is computed are determined by the current sample. If any of the observations have missing values within the current sample, NEGBIN will print a warning message and will drop those observations. NEGBIN also checks that the observations on the dependent variable are integers and are not negative.

The list of independent variables on the NEGBIN command may include variables with explicit lags and leads as well as PDL (Polynomial Distributed Lag) variables. These distributed lag variables are a way to reduce the number of free coefficients when entering a large number of lagged variables in a regression by imposing smoothness on the coefficients. See the PDL section for a description of how to specify such variables.

Output

The output of NEGBIN begins with an equation title and frequency counts for the lowest 10 values of the dependent variable. Starting values and diagnostic output from the iterations will be printed. Final convergence status is printed.
This is followed by the number of observations, mean and standard deviation of the dependent variable, sum of squared residuals, correlation type R-squared, likelihood ratio test for zero slopes, log likelihood, and a table of right hand side variable names, estimated coefficients, standard errors and associated t-statistics.

NEGBIN also stores some of these results in data storage for later use. The table below lists the results available after a NEGBIN command.

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@LHV</td>
<td>list</td>
<td>1</td>
<td>Name of dependent variable</td>
</tr>
<tr>
<td>@RNMS</td>
<td>list</td>
<td>#vars</td>
<td>List of names of right hand side variables</td>
</tr>
<tr>
<td>@IFCONV</td>
<td>scalar</td>
<td>1</td>
<td>=1 if convergence achieved, 0 otherwise</td>
</tr>
<tr>
<td>@YMEAN</td>
<td>scalar</td>
<td>1</td>
<td>Mean of the dependent variable</td>
</tr>
<tr>
<td>@SDEV</td>
<td>scalar</td>
<td>1</td>
<td>Standard deviation of the dependent variable</td>
</tr>
<tr>
<td>@NOB</td>
<td>scalar</td>
<td>1</td>
<td>Number of observations</td>
</tr>
<tr>
<td>@HIST</td>
<td>vector</td>
<td>#values</td>
<td>Frequency counts for each dependent variable value.</td>
</tr>
<tr>
<td>@HISTVAL</td>
<td>vector</td>
<td>#values</td>
<td>Corresponding dependent variable values</td>
</tr>
<tr>
<td>@SSR</td>
<td>scalar</td>
<td>1</td>
<td>Sum of squared residuals</td>
</tr>
<tr>
<td>@RSQ</td>
<td>scalar</td>
<td>1</td>
<td>correlation type R-squared</td>
</tr>
<tr>
<td>@LR</td>
<td>scalar</td>
<td>1</td>
<td>Likelihood ratio test for zero slope coefficients</td>
</tr>
<tr>
<td>%LR</td>
<td>scalar</td>
<td>1</td>
<td>P-value for likelihood ratio test</td>
</tr>
<tr>
<td>@LOGL</td>
<td>scalar</td>
<td>1</td>
<td>Log of likelihood function</td>
</tr>
<tr>
<td>@SBIC</td>
<td>scalar</td>
<td>1</td>
<td>Schwarz Bayesian Information Criterion</td>
</tr>
<tr>
<td>@NCOEF</td>
<td>scalar</td>
<td>1</td>
<td>Number of independent variables (#vars)</td>
</tr>
<tr>
<td>@NCID</td>
<td>scalar</td>
<td>1</td>
<td>Number of identified coefficients</td>
</tr>
<tr>
<td>@COEF</td>
<td>vector</td>
<td>#vars</td>
<td>Coefficient estimates</td>
</tr>
<tr>
<td>@SES</td>
<td>vector</td>
<td>#vars</td>
<td>Standard errors</td>
</tr>
<tr>
<td>@T</td>
<td>vector</td>
<td>#vars</td>
<td>T-statistics</td>
</tr>
<tr>
<td>%T</td>
<td>vector</td>
<td>#vars</td>
<td>p-values for T-statistics</td>
</tr>
<tr>
<td>@GRAD</td>
<td>vector</td>
<td>#vars</td>
<td>Gradient of log likelihood at convergence</td>
</tr>
<tr>
<td>@VCOV</td>
<td>matrix</td>
<td>#vars*</td>
<td>Variance-covariance of estimated coefficients</td>
</tr>
<tr>
<td>@FIT</td>
<td>series</td>
<td>#obs</td>
<td>Fitted values of dependent variable</td>
</tr>
<tr>
<td>@RES</td>
<td>series</td>
<td>#obs</td>
<td>Residuals = actual-fitted values of</td>
</tr>
</tbody>
</table>
dependent variable

If the regression includes a PDL variable, the following will also be stored:

@SLAG    scalar  1    Sum of the lag coefficients
@MLAG    scalar  1    Mean lag coefficient (number of time periods)
@LAGF    vector  #lags Estimated lag coefficients, after “unscrambling”

Method

NEGBIN uses analytic first and second derivatives to obtain maximum likelihood estimates via the Newton-Raphson algorithm. This algorithm usually converges fairly quickly. TSP uses zeros for starting parameter values, except for the constant term and alpha. @START can be used to provide different starting values (see NONLINEAR).

Multicollinearity of the independent variables is handled with generalized inverses, as in all the estimation procedures in TSP.

The exponential mean function is used in the NEGBIN model. That is, if $X$ are the independent variables and $B$ are their coefficients,

$$E(Y|X) = \exp(X^*B)$$

This guarantees that predicted values of $Y$ are never negative.

The ML command can also be used to estimate Negative Binomial models, including panel data models with fixed and random effects. See our web page for the panel examples.

Options

MODEL= type of variance function. For MODEL=1, the variance is proportional to the mean:

$$V(Y|X) = E(Y|X) * (1 + alpha)$$

For the default MODEL=2, the variance is a quadratic function of the mean:

$$V(Y|X) = E(Y|X) + alpha * E(Y|X)^2.$$  

In both cases, the parameter $alpha$ is restricted to be non-negative. $alpha = 0$ corresponds to the Poisson.

Nonlinear options - see the NONLINEAR entry.
Examples

Negative Binomial 2 regression of patents on lags of log(R&D), science sector dummy, and firm size:

```
NEGBIN PATENTS C LRND LRND(-1) LRND(-2) DSCI SIZE ;
```

Negative Binomial 1 regression for the same model:

```
NEGBIN (MODEL=1) PATENTS C LRND LRND(-1) LRND(-2) DSCI SIZE ;
```

References


Nonlinear options

Options  References

These options are common to all of TSP's nonlinear estimation and simulation procedures: ARCH, BJEST, FIML, LSQ, PROBIT, TOBIT, LOGIT, Sampsel, Siml, Solve, ML, etc. See Chapter 10 of the User's Guide for further information.

\texttt{DROPMISS, EPSMIN=<value>, GRADCHEC, GRADIENT=method,} 
\texttt{HCOV=method, HESSCHEC, HITER=method, MAXIT=<number of iterations>, MAXSQZ=<number of squeezes>, NHERMITE=<value>,} 
\texttt{PRINT, SILENT, STEP=<squeezing method>, STEPMEM=<number of iterations>, SQZTOL=<squeezing tolerance>, SYMMETRIC,} 
\texttt{TERSE, TOL=<parameter convergence tolerance>,} 
\texttt{TOLG=<gradient/CRIT convergence tolerance>,} 
\texttt{TOLS=<squeezed parameter convergence tolerance>,} 
\texttt{VERBOSE}

Usage

Include these options among any other special options which you supply within parentheses after the name of the command which invokes the estimation procedure:

\texttt{FIML (ENDOG=..., nonlinear options) list of eq names ;}

Method

The method used for nonlinear estimation is generally a standard gradient method, explained in more detail in Chapter 10 of the User's Guide. Briefly, at each iteration, a new parameter vector is computed by moving in the direction specified by the gradient of the likelihood (uphill), weighting this gradient by an approximation to the matrix of second derivatives at that point (in order to adjust for the curvature). Convergence is declared when the changes in the parameters are all "small", where "small" is defined by the TOL= option.

Options

\texttt{DROPMISS/NODROPMISS} specifies whether observations with missing values in any variables are to be dropped. This can be useful if the equation being estimated varies according to the presence of good data, but use with caution.
Nonlinear options

**EPSMIN** = minimum parameter change for numeric derivatives [default is .0001]. This is also used to control the numeric stepsize when computing HCOV=C and HCOV=U. If you have parameters smaller than .00001 in magnitude, it will be helpful to use an EPSMIN with a value somewhat smaller than your smallest parameter. Otherwise, too large a stepsize is used and the parameters will appear to have zero standard errors.

**GRADCHEC/NOGRADCHEC** evaluates and compares the analytic and numerical gradient for the current model at the starting values. No actual estimation takes place. Useful for checking derivatives of a new likelihood function. The numeric gradient is evaluated in a time-consuming but accurate way. See the GRAD=C4 option.

**GRADIENT** = ANALYTIC or C2 or C4 or FORWARD specifies the method of calculating numeric first derivatives.

- **GRAD=A** is the default when analytic first derivatives are available (as is usually the case).
- **GRAD=FORWARD** calculates the numeric derivatives for a given parameter B as

  \[ D = \frac{(F(B+\text{EPS}) - F(B))}{\text{EPS}} \]

  (1 function evaluation per parameter)

- **GRAD=C2** (CENTRAL2) uses

  \[ D = \frac{(F(B+\text{EPS}) - F(B-\text{EPS}))}{2\times\text{EPS}} \]

  (2 function evaluations per parameter)

- **GRAD=C4** uses

  \[ D = \frac{(F(B+2\times\text{EPS}) - 8F(B+\text{EPS}) + 8F(B-\text{EPS}) - F(B-2\times\text{EPS}))}{12\times\text{EPS}} \]

  (4 function evaluations per parameter)

In all cases, \( \text{EPS} = \max(\text{ABS}(.001\times B), \text{EPSMIN}) \)
Nonlinear options

HCOV=B or N or G or F or D or W or R or P or Q or U or C or BNW, etc., specifies the method for calculating the asymptotic covariance matrix of the parameter estimates (and standard errors). The default is usually N or B, depending on the procedure. Some procedures may not have N (and thus W) available. A label is printed below each table of standard errors and asymptotic t-statistics identifying the method of calculation used. More than one method may be specified for alternative VCOV matrices and standard errors. In this case, the first method is stored in @VCOV and @SES, and the VCOV for each method is stored under the name constructed by appending the letter to @VCOV. For example, HCOV=NB would store @VCOV, @VCOVN, and @VCOVB. Consult the table below to see which option is the default in a particular procedure.

The P and Q options are for panel data and are available for ML and PROBIT (REI or FEI). HCOV=P computes grouped BHHH standard errors from the gradient of the objective function using the formula

\[ V_p = \left( \sum_{i=1}^{NI} G_i G_i' \right)^{-1} \]

where \( G_i = \sum_{t=1}^{\tau_i} G_{it} \)

instead of the usual formula

\[ V_B = \left( N \sum_{i=1}^{NI} \sum_{t=1}^{\tau_i} G_{it} G_{it}' \right)^{-1} \]

where \( G_{it} \) is the gradient vector for individual \( i \) and period \( t \). Unlike the usual formula, this version of the estimate does not assume independence within individual across different time periods. HCOV=Q computes the robust version of this matrix

\[ NV_P^{-1} N \]

where \( N \) is the Newton (inverse second derivative) matrix. See Wooldridge, p. 407. For linear models, this matrix is exactly equivalent to that computed by OLSQ, 2SLS, LSQ, and PANEL using the HCOMEQA=BLOCK option.

Note that the rank of \( V_p \) is at most \( NI \), where \( NI \) is the number of individuals, so that when \( NI \) is less than the number of coefficients \( K \), the grouped panel estimates of the variance-covariance matrix will be singular and therefore probably inappropriate. However in most cases, \( NI >> K \), and this problem will not arise. Note also that for fixed effect estimation, the gradient is always zero for the fixed effects at the optimum, so the block of \( V \) corresponding to these effects is zero. For this reason, standard errors for the fixed effects are always computed using the Newton (second derivative) matrix.

HESSCHEC/NOHESSCH compares analytic and discrete Hessian (differenced analytic gradient)
Nonlinear options

HITER=B or N or G or F or D specifies the method of Hessian (second derivative matrix) approximation to be used during the parameter iterations. The options are the same as those described above for the estimate of the covariance matrix of the parameter estimates.

Table of HCOV and HITER options

<table>
<thead>
<tr>
<th>Option</th>
<th>Used to iterate?</th>
<th>Name and description</th>
<th>Procedures for which it is the default</th>
</tr>
</thead>
<tbody>
<tr>
<td>B</td>
<td>yes</td>
<td>BHHH (Berndt-Hall-Hall-Hausman) Covariance of the analytic gradient.</td>
<td>ML</td>
</tr>
<tr>
<td>N</td>
<td>yes</td>
<td>Newton (Analytic second derivatives)</td>
<td>ARCH iterations, AR1, PROBIT, TOBIT, LOGIT, SAMPSEL</td>
</tr>
<tr>
<td>G</td>
<td>yes</td>
<td>GAUSS (Gauss-Newton). Quadratic form of the analytic gradient and the residual covariance matrix.</td>
<td>LSQ, FIML iterations</td>
</tr>
<tr>
<td>F</td>
<td>yes</td>
<td>BFGS (Broyden-Fletcher-Goldfarb-Shanno). Analytic or numeric first derivatives, and rank 1 update approximation of the Hessian from iterations. Usually HITER=F is superior to HITER=D. The HCOV=F option is valid only if HITER=F.</td>
<td>None</td>
</tr>
<tr>
<td>D</td>
<td>yes</td>
<td>DFP (Davidon-Fletcher-Powell). Analytic or numeric first derivatives, and rank 1 update approximation of the Hessian from iterations. This option is valid only if HITER=D. For upward compatibility, it implies a default of</td>
<td>None</td>
</tr>
</tbody>
</table>
### Nonlinear options

<table>
<thead>
<tr>
<th>Option</th>
<th>Value</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>W</td>
<td>no</td>
<td>Eicker-White. A combination of analytic second derivatives and BHHH (see the White Reference).</td>
</tr>
<tr>
<td>R</td>
<td>no</td>
<td>Robust. Robust to heteroskedasticity. This is equivalent to W and used in LSQ only.</td>
</tr>
<tr>
<td>P</td>
<td>no</td>
<td>Panel grouped estimate (allows for free correlation within panel) - PROBIT (FEI; REI) and PANEL only</td>
</tr>
<tr>
<td>Q</td>
<td>no</td>
<td>Panel grouped estimate robust to heteroskedasticity across units (allows for free correlation within panel) - PROBIT (FEI; REI) and PANEL. For OLSQ, 2SLS, and LSQ, use HCOMEGA=BLOCK</td>
</tr>
<tr>
<td>C</td>
<td>yes</td>
<td>Discrete Hessian (numeric second derivatives based on analytic first derivatives)</td>
</tr>
<tr>
<td>U</td>
<td>yes</td>
<td>Numeric second derivatives</td>
</tr>
<tr>
<td>NBW</td>
<td>no</td>
<td>Print all three standard error estimates.</td>
</tr>
</tbody>
</table>

**MAXIT** = maximum number of iterations. The default is 20.

**MAXSQZ** = maximum number of "squeezes" in the stepsiz search. The default depends on **STEP**:

<table>
<thead>
<tr>
<th>STEP option</th>
<th>MAXSQZ default</th>
</tr>
</thead>
<tbody>
<tr>
<td>BARD</td>
<td>10</td>
</tr>
<tr>
<td>BARDB</td>
<td>10</td>
</tr>
<tr>
<td>CEA</td>
<td>10</td>
</tr>
</tbody>
</table>
Nonlinear options

<table>
<thead>
<tr>
<th>Option</th>
<th>Value</th>
</tr>
</thead>
<tbody>
<tr>
<td>CEAB</td>
<td>10</td>
</tr>
<tr>
<td>GOLDEN</td>
<td>20</td>
</tr>
</tbody>
</table>

Note that some routines (BJEST, LOGIT) reserve MAXSQZ=123 for special options.

**NHERMITE**= number of Hermite quadrature points for numeric integration, used for PROBIT (REI). [default is 20]

**PRINT/NOPRINT** produces short diagnostic output at each iteration, including a table of the current parameter estimates and their change vector. The value of the objective function for each squeeze on the change vector is also printed. **CRIT** is the norm of the gradient in the metric of the Hessian, which approaches zero at convergence.

**SILENT/NOSILENT** suppresses all printed output.

**STEP**= BARD or BARDB or CEA or CEAB or GOLDEN specifies the stepsize method for squeezing. The default depends on HITER and the procedure:

<table>
<thead>
<tr>
<th>HITER option</th>
<th>STEP default</th>
</tr>
</thead>
<tbody>
<tr>
<td>Newton</td>
<td>CEA</td>
</tr>
<tr>
<td>BHHH</td>
<td>CEA</td>
</tr>
<tr>
<td>Gauss</td>
<td>BARD (for LSQ); CEAB (for FIML)</td>
</tr>
<tr>
<td>DFP</td>
<td>GOLDEN</td>
</tr>
</tbody>
</table>

**STEPMEM**= n, the number of iterations at which to reset the stepsize. The initial stepsize lambda is remembered from the previous iteration, and reset to lambda=1 every n'th iteration. The default is STEPMEM=1, which resets lambda=1 every iteration. This option may be helpful if the natural stepsize for a model is different from 1 and tends to be about the same for successive iterations.

**SQZTOL**= tolerance of determining stepsize. Used for STEP=GOLDEN. The default is 0.1.

**SYMMETRIC/NOSYMMETRIC** is an old option which has been replaced with **GRADIENT=method**. SYMMETRIC is the same as GRAD=C4; NOSYM is equivalent to GRAD=FORWARD.

**TERSE/NOTERSE** produces brief output consisting of the objective function for the estimation procedure, and a table of coefficient estimates and standard errors.
Nonlinear options

**TOL** = tolerance of determining convergence of the parameters, using a unit stepsize. The default for most procedures is .001; for AR1 it is .000001.

**TOLG** = tolerance of determining convergence of the norm of the gradient (printed as CRIT in the output). The default is .001. CRIT = g'H-1g , which is usually many orders of magnitude smaller than .001.

**TOLS** = tolerance of determining convergence of the parameters, using the squeezed step. The default is 0, that is, ignore the squeezed change in parameters and use the regular TOL instead.

**VERBOSE/NOVERBOSE** produces lots of diagnostic output, including the gradient, Hessian, and inverse Hessian at each iteration, and the non-inverted Hessian for each output VCOV.

Starting values:

The default values depend on the procedure. For the standard TSP models which are (potentially) nonlinear in the parameters (LSQ,FIML,ML), the user provides them with PARAM and SET. PROBIT and LOGIT use zeros. TOBIT uses a regression and formulas from Greene (1981). SAMPSEL uses probit and a regression.

The default is overridden in the linear model procedures (ARCH, BJEST, PROBIT, TOBIT,LOGIT, and SAMPSEL) if the user supplies a matrix named @START. The length of @START must be equal to the number of parameters in the estimation (otherwise it is ignored). The easiest way to create @START is with a statement like:

```
MMAKE @START 12.3 4.56 33 44 55;
```

The order of the parameters in @START is obvious for most of the linear models, except for the following:

**TOBIT**: SIGMA comes last.

**SAMPSEL**: the probit equation is first, then the regression equation, and then SIGMA and RHO.

References


Nonlinear options


NOPLIT

Examples

NOPLIT turns off the PLOTS options so that actual and fitted values are not plotted after each regression. NOPLIT is the default.

NOPLIT ;

Usage

NOPLIT takes no arguments; it is needed only when a PLOTS statement has appeared earlier. PLOTS/NOPLIT are also available on the OPTIONS statement.

The PLOTS/NOPLIT options apply to any procedures which produce residual plots: these are the OLSQ, INST, AR1, LSQ, and ACTFIT.

Examples

This example suppresses the printing and plotting of residuals after a regression on large amounts of data, and then uses ACTFIT with a different sample to plot a portion of them:

SMPL 1 896 ;
NOPLIT ;
OLSQ LOGP C LOGR SCISECT NPLNT72 DPAT0 LOGR(-1) ;
SMPL 1 16 881 896 ;
PLOTS ;
ACTFIT @ACT @FIT ;
SMPL 1 896 ;
NOPLIT ;
Examples

NOPRINT turns off the printing of input data in the load section. It applies only to free format data input, since data read under fixed format is not printed in any case.

NOPRINT ;

Usage

Include the NOPRINT statement at any point in your data section where you wish to turn off the printing of the input. It remains in force until the end of the data section. Do not put the statement between a LOAD statement and the data which goes with it; these must be contiguous. The PRINT/NOPRINT option is also available on the LOAD statement itself.

Examples

The following example of a LOAD section shows the use of the NOPRINT command to suppress printing of the entire data section:

NOPRINT ;
FREQ A ; SMPL 20 41 ;
LOAD YEAR CX I G YT K1 P W1 W2 Y ;
1920 39.8 2.7 4.6 47.1 180.1 12.7 28.8 2.2 43.7
1921 41.9 -.2 6.6 48.3 182.8 12.4 25.5 2.7 40.6
....... more data input ....
END ;
**NOREPL**

**Examples**

NOREPL turns off the replacement mode option (REPL). REPL specifies that series are to be updated rather than completely replaced when the current sample under which they are being computed does not cover the complete series.

```
NOREPL;
```

**Usage**

The REPLace mode is the default; use NOREPL if you do not want previously existing series updated when they are modified. For example, use the REPL mode to create a series element by element with a DO loop and SET statements. Later on, if you want to recreate the same series with a slightly different sample, the REPL mode could cause the old observations to be mixed in with the new, so you might want to use NOREPL just for safety.

**Examples**

```
REPL;
SMPL 1 10; GENR D = 0;
SMPL 11 20; GENR D = 1;
NOREPL;
```

This creates a series named D which is zero for observations 1 through 10, and one for observations 11 through 20.
NORMAL

Examples

NORMAL normalizes a series so that a chosen observation has a predetermined value. It accomplishes this by dividing all the observations of the series by the ratio of the supplied value to the chosen observation's value.

NORMAL <series name> <obs. id> <value> [<series name> <obs. id> <value> .......];

Usage

After NORMAL, list the name of the series, the observation identifier of the base observation, and the value to be assigned to the base observation. The normalized series will replace the original series (for those observations in the current sample).

The observation identifier must include the period if the frequency is neither NONE nor ANNUAL. It is written in the form YYYY:PP or YY:PP where YYYY or YY is the year and PP is the period.

You may normalize as many series with the same statement as you wish: just include three arguments (series name, observation identifier, and value) for each one.

Output

NORMAL produces no printed output. One or more series are replaced in data storage.

Examples

This example normalizes the CPI to have the value 100 in 1975:

NORMAL CPI,75,100 ;

This is equivalent to the following statements:

SET BASE=CPI(75) ;
CPI = 100*CPI/BASE ;

This example normalizes a set of quarterly price series so they have the value 1 in the first quarter of 1972:

NORMAL P1,72:1,1 P2,72:1,1 P3,72:1,1 ;
NOSUPRES turns off the suppression of output for the selected results from procedures.

*NOSUPRES* <list of result names>; 

**Usage**

The arguments to NOSUPRES can be any of the output names beginning with @ described in this help system. The printing of the output associated with these names will be suppressed throughout the TSP program unless a NOSUPRES or REGOPT command with these codes is issued. The output results are still stored in memory and may be accessed.

See also SUPRES and REGOPT.
OLSQ is the basic regression procedure in TSP. It obtains ordinary least squares estimates of the coefficients of a regression of the dependent variable on a set of independent variables. Options allow you to obtain weighted least squares estimates to correct for heteroskedasticity, or to obtain standard errors which are robust in the presence of heteroskedasticity of the disturbances (see the GMM command for robustness to autocorrelation).

**OLSQ (HCOMEGA=BLOCK or DIAGONAL, HCTYPE=<<robust SE type>>, HI, NORM or UNNORM, ROBUSTSE, SILENT, TERSE, WEIGHT=<name of weighting variable>, WTYPE=<weight type>)**

<dependent variable> <list of independent variables> ;

**Usage**

In the basic OLSQ statement, you list the dependent variable and then the independent variables in the equation. To have an intercept term in the regression, include the special variable C or CONSTANT in the list of independent variables. The number of independent variables is limited by the overall limits on the number of arguments per statement and the amount of working space; obviously, it is also limited by the number of data observations available.

The observations over which the regression is computed are determined by the current sample. If any observations have missing values within the current sample, they are dropped from the sample, and a warning message is printed for each series with missing values. The number of observations remaining is printed with the regression output. @RES and @FIT will have missing values in this case, and the Durbin-Watson will be adjusted for the sample gaps.

The list of independent variables on the OLSQ command may include variables with explicit lags and leads as well as PDL (Polynomial Distributed Lag) variables. These PDL variables are a way to reduce the number of free coefficients when you are entering a large number of lagged variables in a regression by imposing smoothness on the coefficients. See the PDL section for a description of how to specify a PDL variable.

**Output**
The output of OLSQ begins with an equation title and the name of the dependent variable. This is followed by statistics on goodness-of-fit: the sum of squared residuals, the standard error of the regression, the R-squared, the Durbin-Watson statistic for auto correlation of the residuals, a Lagrange multiplier test for heteroskedasticity, the Jarque-Bera test for normality, and an F-statistic for the hypothesis that all coefficients in the regression except the constant are zero. If there is no constant in the regression and the mean was not removed from the dependent variable prior to the regression, the F-statistic may be meaningless. See the REGOPT command for a large variety of additional regression diagnostics.

A table of right hand side variable names, estimated coefficients, standard errors and associated t-statistics follows. The variance-covariance and correlation matrices are printed next if they have been selected with the REGOPT command.

If there are lagged dependent variables on the right hand side, the regular Durbin-Watson statistic is biased, so an alternative test for serial correlation is computed. The statistic is computed by including the lagged residual with the right hand side variables in an auxiliary regression (with the residual as the dependent variable), and testing the lagged residual's coefficient for significance. See the Durbin reference for details; this method is very similar to the method used for correcting the standard errors for AR1 regression coefficients in the same lagged dependent variables case. This statistic is more general than "Durbin's h" statistic since it applies in cases of several lagged dependent variables. It is not computed if there is a WEIGHT or gaps in the SMPL, and the presence of lagged dependent variables is not detected if they are computed with GENR (instead of being specified with an explicit lag like OLSQ Y C X Y(-1); or in a PDL).

If the PLOTS option is on, TSP prints and plots the actual and fitted values of the dependent variable and the residuals. OLSQ also stores most of these results in data storage for your later use. The table below lists the results available:

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@LHV</td>
<td>list</td>
<td>1</td>
<td>Name of the dependent variable</td>
</tr>
<tr>
<td>@RNMS</td>
<td>list</td>
<td>#vars</td>
<td>Names of right hand side variables</td>
</tr>
<tr>
<td>@SSR</td>
<td>scalar</td>
<td>1</td>
<td>Sum of squared residuals</td>
</tr>
<tr>
<td>@S</td>
<td>scalar</td>
<td>1</td>
<td>Standard error of the regression</td>
</tr>
<tr>
<td>@S2</td>
<td>scalar</td>
<td>1</td>
<td>Standard error squared</td>
</tr>
<tr>
<td>@YMEAN</td>
<td>scalar</td>
<td>1</td>
<td>Mean of the dependent variable</td>
</tr>
<tr>
<td>@SDEV</td>
<td>scalar</td>
<td>1</td>
<td>Standard deviation of the dependent variable</td>
</tr>
<tr>
<td>@NOB</td>
<td>scalar</td>
<td>1</td>
<td>Number of observations</td>
</tr>
</tbody>
</table>
OLSQ computes the matrix equation:

\[
\begin{align*}
\text{SSRO} &= \sum (y_{it} - \hat{y}_{it})^2 \\
\text{COEF} &= \hat{\beta}_1, \hat{\beta}_2, \ldots, \hat{\beta}_k \\
\text{SES} &= \hat{\sigma}_1, \hat{\sigma}_2, \ldots, \hat{\sigma}_k \\
\text{T} &= t_1, t_2, \ldots, t_k \\
\text{VCOV} &= \text{Var} (\hat{\beta}_1, \hat{\beta}_2, \ldots, \hat{\beta}_k) \\
\text{RES} &= y_{it} - \hat{y}_{it} \\
\text{FIT} &= \hat{y}_{it} \\
\text{HI} &= \text{Diagonal of } \text{"hat matrix" if the HI option is on} \\
\end{align*}
\]

If the regression includes a PDL or SDL variable, the following will also be stored:

\[
\begin{align*}
\text{SLAG} &= \sum \text{lag coefficients} \\
\text{MLAG} &= \text{Mean lag coefficient (number of time periods)} \\
\text{LAGF} &= \text{Estimated lag coefficients, after } \text{"unscrambling"} \\
\end{align*}
\]

**REGOPT (NOPRINT) LAGF;**

will turn off the lag plot for PDL variables.

**Method**

If the regression includes a PDL or SDL variable, the following will also be stored:
\[ b = (X'X)^{-1}X'y \]

where \( X \) is the matrix of independent variables and \( y \) is the vector of independent variables. The method used to compute this regression (and all the other regression-type estimates in TSP) is a very accurate one, which involves applying an orthonormalizing transformation to the \( X \) matrix before computation of the inner products and inverse, and then untransforming the result (see ORTHON in this manual). See OPTIONS FAST; to compute faster and slightly less accurate regressions (without orthonormalization).

OLSQ has been tested using the data of Longley on everything from an IBM 370 to a modern Pentium computer; it gives accurate results to six digits when the data is single precision. For the artificial problem suggested by Lauchli (see the Wampler article), OLSQ gives correct results for the coefficients to about five places, until epsilon becomes so small that the regression is not computable. Before this happens, OLSQ detects the fact that it cannot compute the regression accurately and drops one of the variables by setting its coefficient to zero and printing a warning. The results for these special regressions become more accurate when OPTIONS DOUBLE; is in effect, because these data have an unusually high number of significant digits. See the Benchmarks section of the TSP web page for more information on regression accuracy with Longley and other standard regression datasets.

Options

\textbf{HCOMEGA} = BLOCK or DIAGONAL specifies the form of the \( \Omega = E[uu'] \) matrix to use when computing ROBUST standard errors. Ordinarily, the default is diagonal, which yields the usual robust standard errors. When FREQ (PANEL) is in effect, the default is BLOCK, which allows for cross-time correlation of the disturbances within individuals. This feature can be used for any kind of grouped data, simply by ensuring that the relevant PANEL setup has been defined.

\textbf{HCTYPE} the type of heteroskedastic-consistent standard errors to compute (between 0 and 3, with a default of 2. This option implies ROBUSTSE. In general, the robust estimate of the variance-covariance matrix has the form:

\[ V = (X'X)^{-1} \left[ \sum_{i=1}^{n} \frac{e_i^2 X_i X_i'}{d_i} \right] (X'X)^{-1} \]

The option HCTYPE specifies the formula used for \( d(i) \):
the usual Eicker-White asymptotic formula

use finite sample degrees of freedom

unbiased if \( e \) is truly homoskedastic

jackknife approximation

where \( h(i) \) is the diagonal of the "hat matrix" (defined below). If \( 1-h(i) \) is zero for some \( i \) and \( e(i) \) is nonzero, HCTYPE=1 is used. Both HCTYPE=2 and HCTYPE=3 have good finite sample properties. See Davidson and MacKinnon, pp. 552-556 for details.

HI/NOHI specifies whether the diagonal of the "hat matrix" is stored in the series @HI. The hat matrix is defined as

\[
H = X (X'X)^{-1}X'
\]

This is useful for detecting "influential" observations (data errors, outliers, etc.). For example,

\[
SELECT @HI > 2*@NCOEF/@NOB;
\]

identifies the influential observations. (See the Belsley, Kuh, and Welsch or Krasker, Kuh, and Welsch references).

NORM/UNNORM tells whether the weights are to be normalized so that they sum to the number of observations. This has no effect on the coefficient estimates and most of the statistics, but it makes the magnitude of the unweighted and weighted data the same, on average, which may help in interpreting the results. The coefficient standard errors and t-statistics are affected. NORM has no effect if the WEIGHT option has not been specified.

ROBUSTSE/NOROBUST causes the variance of the coefficient estimates, the standard errors, and associated t-statistics to be computed using the formulas suggested by White, among others. These estimates of the variance are consistent even when the disturbances are not homoskedastic, and when their variances are correlated with the independent variables in the model. They are not consistent when the disturbances are not independent, however. See the Davidson and MacKinnon reference. See the HCTYPE= option for the exact formulas.

SILENT/NOSILENT suppresses all output. The results are stored.

TERSE/NOTERSE suppresses all regression output except for the log likelihood and the table of coefficients and standard errors.
**WEIGHT=** the name of a series used to weight the observations. The data are multiplied by the square roots of the normalized weighting series before the regression is computed (see NORM above). The series will be proportional to the inverses of the variances of the residuals. If the weight is zero for a particular observation, that observation is not included in the computations nor is it counted in determining degrees of freedom.

**WTYPE=** HET or REPEAT, the weight type. The default is REPEAT, where the weight is a repeat count (it multiplies the likelihood function directly). This is used for grouped data and is consistent with the UNNORM option. WTYPE=HET means the weight is for heteroskedasticity only (it enters the likelihood function only through the variance). The only difference between these two options in the regression output is the value of the log likelihood function (all the coefficients, standard errors, etc. are identical). With WTYPE=HET, the log likelihood includes the sum of the log weights; the default WTYPE=REPEAT does not include this.

**Examples**

This example estimates the consumption function for the illustrative model:

```
OLSQ CONS,C,GNP ;
```

Using population as weights, the next example regresses the fraction of young people living alone on other demographic characteristics across states. Since the regression is in terms of per capita figures, the variance of the disturbances is proportional to the inverse of population.

```
OLSQ (WEIGHT=POP) YOUNG,C,RSALE,URBAN,CATHOLIC ;
```

Other examples of the OLSQ command:

```
OLSQ (ROBUSTSE) LOGP C LOGP(-1) LOGR ;
OLSQ TBILL C RATE(4,12,FAR) ;
```

**References**


OPTIONS

Options   Examples

OPTIONS is used to set various options for the TSP run.

OPTIONS APPEND, ARGSUB, BASEYEAR=value, CHARID, CRT, DATE, DEBUG, DISPLAY=<monitor type>, DOUBLE, FAST, HARDCOPY, INDENT=<# of spaces>, LEFTMG=<left margin>, LIMCOL=<column width for input>, LIMERR=<maximum # of errors>, LIMNUM=<maximum # of numerical errors>, LIMPRN=<printer line width>, LIMWARN=<maximum # of warning messages printed>, LIMWMISS=<maximum # of missing value warning messages printed>, LIMWNUMC=<maximum # of numeric warning messages printed per command>, LINLIM=<lines per printer page>, MEMORY=<size of memory for TSP>, NWIDTH=<# of digits printed>, PLOTS, REPL, RESID, SECONDS=<# of seconds>, SIGNIF=<# of significant digits printed>, TOL=<tolerance for matrix inversion> ;

Usage

Usually OPTIONS is the first statement in a TSP run, before the NAME statement (if there is one); doing this sets the output format for the entire run, for example, the CRT option. However, an OPTIONS statement may be included anywhere in your TSP program (except the load section) to change certain global parameters.

If you use the same options repeatedly, you may want to place them in a login.tsp file. Every time TSP starts, it checks for a login.tsp file, and sets the options accordingly. Normally, TSP looks for login.tsp in your working directory. If it does not find one, it looks in the directory in which you installed TSP for DOS and Windows, in the folder in which you installed TSP for Macs, and in the home directory on Unix.

Many options on the OPTIONS statement can also be set using individual commands for compatibility with older versions of TSP. These commands include PLOTS/NOPLOT, REPL/NOREPL, DEBUG/NODBUG, MAXERR (same as LIMERR), and TOL.

Options

APPEND/NOAPPEND updates the .OUT file (in batch mode) at each nonlinear iteration. This is useful for monitoring the progress of a long estimation on a multitasking operating system.
ARGSUB/NOARGSUB controls substitution of actual arguments for formal arguments inside a PROC. The default is ARGSUB. NOARGSUB is useful if the PROC has LOCAL variables with the same names as global variables being passed as arguments to the PROC. It prevents the local variables from being used instead of the PROC arguments. The disadvantage is that the labels in the output will be the formal argument names rather than that actual argument names.

BASEYEAR = value used to make dates from 2 digit numbers. The default is 1900. For example, by default, 86:2 means 1986:2. If you set BASEYEAR=2000, 86:2 would mean 2086:2. BASEYEAR can also be set to zero, so that you can use dates in the first two centuries.

CHARID/NOCHARID treats the ID series as characters (instead of numbers) when printing observation labels. CHARID requires use of the DOUBLE option also. To use this option, read in your ID series (called ID) using an A8 format statement.

CRT/NOCRT sets several output format options to values suitable for viewing output on a 24 line by 80 character screen. These are LIMPRN=80, LINLIM=24, and LEFTMG=0. The page headings (date, time, and page number) are also suppressed in CRT mode.

DATE/NODATE specifies whether the date and time headings at the top of each page of TSP output are to be printed. A user title, and page number if present, is still printed. This option only applies when page headings are being printed (NOCRT).

DEBUG/NODBUG sets the debug option, causing intermediate results to be printed. This is described more fully in the DEBUG section and is of use primarily to TSP programmers.

DISPLAY = monitor type for PC/386 graphics (essentially obsolete).

DOUBLE/NODOUBLE causes all subsequent series to be stored in double precision (15-16 digits, vs. the default single precision).

FAST/NOFAST performs fast regression calculations (without orthonormalization). These are slightly less accurate, but usually yield no differences in the first 5 or so digits. Such calculations are also used in the iterations of LSQ, 3SLS, and GMM. Used to speed up runs with Monte Carlo loops, or more than 1000 observations.

HARDCOPY sets several output format options to values suitable for output routed to a printer. These are LIMPRN=120, LINLIM=60, LEFTMG=20, INDENT=10, and DATE and are the default options for printer output.
INDENT = number of spaces to indent printed output from the left margin. The default value is 5.

LEFTMG = left margin for printed output. The default is 20 (the first column in which output will be printed is 21).

LIMCOL = column width for input (number of columns read in each input line). The default value is 500 - this option is essentially obsolete.

LIMERR = maximum number of errors allowed in this TSP run. The default value is 25.

LIMNUM = maximum number of numerical warnings (divide by zero, log of zero or negative number, and exponentiation of too large a number) allowed in this run, before each subsequent one is treated as an ERROR instead of as a WARNING. The default value is 100000.

LIMPRN = printer line width, the maximum number of printing positions on the printer, including the left margin. The default is 132, which is correct for most high-speed printers. Occasionally these printers have only 120 positions, and your local installation may change the default accordingly.

LIMWARN = maximum number of warning messages to print. The default value is 100000.

LIMWMISS = maximum number of warning messages about missing values to print. The default is 10.

LIMWNUMC = maximum number of numeric warning messages to print in any particular command. The default value is 10. This means that each command will print at most 10 numeric warning messages, and then the remainder for that command will be suppressed.

LINLIM = number of lines per printer page. The default is 60, which is correct for most conventional printed output.

MEMORY = approximate memory used by TSP (in MB). This option only works if OPTIONS is the first command in the run, or the first command in the login.tsp file. The default is 4MB, and the minimum is 2.1MB. Calculate memory as 2MB plus 4MB per million words of working space desired. MEMORY=4 should be enough for most time series datasets and small cross sections. The memory actually used is printed at the end of the TSP run.

NWIDTH = maximum number of digits to be printed for numbers in tables. This is the number of columns allowed for each number and the default value is 13.
OPTIONS

PLOTS/NOPLOT tells whether residual plots are to be printed following each estimation. See the PLOTS command description for further information.

NOREPL/REPL tells whether replacement mode is to be used in updating series. See REPL for further information.

RESID/NORESID tells whether residuals and fitted values are to be computed and stored after the estimation procedures (LSQ, FIML, INST, and AR1).

SECONDS= number of seconds. All commands which take longer than this amount of time to execute display a message on the screen giving line number, command name, and elapsed time for execution. The default is 10 seconds. For more precise control of timing single or multiple commands, use the DATE variable; command. If you supply a fractional part to the argument, like OPTIONS SECONDS=2.1; within-command profile timings will be given, for regression commands, ML, GMM, and MATRIX. This is used to investigate which parts of commands are slow, for use in improving speed.

SIGNIF= number of significant digits to be printed in tables. In general, this is the number of digits printed to the right of the decimal point and the default value is 5.

TOL= tolerance for matrix inversion. This parameter is used to decide when a matrix is singular. The value of TOL is compared to the diagonals of the square root matrix of the matrix being inverted as it is formed, and if the diagonal is smaller than TOL, it is set to zero, effectively dropping that row and column from the matrix before inversion. The default value of TOL on IBM is 10.E-13, a conservative value; this value causes nearly singular matrices to fail, rather than letting through some exactly singular ones.

Examples

This example uses the dates 53 BC to 86 AD:

OPTIONS BASEYEAR=0 ;
FREQ A ;
SMPL -53 86 ;

Here are some other OPTIONS commands:

OPTIONS REPL,PLOTS,TOL=1.E-10 ;
OPTIONS NWIDTH=10,SIGNIF=3 ;
ORDPROB obtains estimates of the linear Ordered Probit model, where the dependent variable takes on only nonnegative integer ordered category values. The scaling of the category values does not matter (although they should be positive and integer for convenience); only information about their order is used in estimation.

**ORDPROB (nonlinear options)** `<dependent variable>` `<list of independent variables>` ;

**Usage**

The basic ORDPROB statement is like the OLSQ statement: first list the dependent variable and then the independent variables. If you wish to have an intercept term in the regression (usually recommended), include the special variable C or CONSTANT in your list of independent variables. You may have as many independent variables as you like subject to the overall limits on the number of arguments per statement and the amount of working space, as well as the number of data observations you have available.

The observations over which the regression is computed are determined by the current sample. If any of the observations have missing values within the current sample, ORDPROB will print a warning message and will drop those observations. ORDPROB also checks that the observations on the dependent variable are integers and are not negative.

The list of independent variables on the ORDPROB command may include variables with explicit lags and leads as well as PDL (Polynomial Distributed Lag) variables. These distributed lag variables are a way to reduce the number of free coefficients when entering a large number of lagged variables in a regression by imposing smoothness on the coefficients. See the PDL section for a description of how to specify such variables.

**Output**

The output of ORDPROB begins with an equation title and frequency counts for the lowest 10 values of the dependent variable. Starting values and diagnostic output from the iterations will be printed. Final convergence status is printed.
ORDPROB

This is followed by the number of observations, mean and standard deviation of the dependent variable, sum of squared residuals, scaled R-squared, likelihood ratio test for zero slopes, log likelihood, and a table of right hand side variable names, estimated coefficients, standard errors and associated t-statistics.

ORDPROB also stores some of these results in data storage for later use. The table below lists the results available after a ORDPROB command.

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@LHV</td>
<td>list</td>
<td>1</td>
<td>Name of dependent variable</td>
</tr>
<tr>
<td>@RNMS</td>
<td>list</td>
<td>#params</td>
<td>List of names of right hand side variables</td>
</tr>
<tr>
<td>@IFCONV</td>
<td>scalar</td>
<td>1</td>
<td>=1 if convergence achieved, 0 otherwise</td>
</tr>
<tr>
<td>@YMEAN</td>
<td>scalar</td>
<td>1</td>
<td>Mean of the dependent variable</td>
</tr>
<tr>
<td>@SDEV</td>
<td>scalar</td>
<td>1</td>
<td>Standard deviation of the dependent variable</td>
</tr>
<tr>
<td>@NOB</td>
<td>scalar</td>
<td>1</td>
<td>Number of observations</td>
</tr>
<tr>
<td>@HIST</td>
<td>vector</td>
<td>#values</td>
<td>Frequency counts for each dependent variable value.</td>
</tr>
<tr>
<td>@HISTVAL</td>
<td>vector</td>
<td>#values</td>
<td>Corresponding dependent variable values</td>
</tr>
<tr>
<td>@SSR</td>
<td>scalar</td>
<td>1</td>
<td>Sum of squared residuals</td>
</tr>
<tr>
<td>@RSQ</td>
<td>scalar</td>
<td>1</td>
<td>correlation type R-squared</td>
</tr>
<tr>
<td>@SRSQ</td>
<td>scalar</td>
<td>1</td>
<td>Scaled R-squared</td>
</tr>
<tr>
<td>@LR</td>
<td>scalar</td>
<td>1</td>
<td>Likelihood ratio test for zero slope coefficients</td>
</tr>
<tr>
<td>%LR</td>
<td>scalar</td>
<td>1</td>
<td>P-value for likelihood ratio test</td>
</tr>
<tr>
<td>@LOGL</td>
<td>scalar</td>
<td>1</td>
<td>Log of likelihood function</td>
</tr>
<tr>
<td>@SBIC</td>
<td>scalar</td>
<td>1</td>
<td>Schwarz Bayesian Information Criterion</td>
</tr>
<tr>
<td>@NCOEF</td>
<td>scalar</td>
<td>1</td>
<td>Number of parameters (#params)</td>
</tr>
<tr>
<td>@NCID</td>
<td>scalar</td>
<td>1</td>
<td>Number of identified coefficients</td>
</tr>
<tr>
<td>@COEF</td>
<td>vector</td>
<td>#params</td>
<td>Coefficient estimates</td>
</tr>
<tr>
<td>@SES</td>
<td>vector</td>
<td>#params</td>
<td>Standard errors</td>
</tr>
<tr>
<td>@T</td>
<td>vector</td>
<td>#params</td>
<td>T-statistics</td>
</tr>
<tr>
<td>%T</td>
<td>vector</td>
<td>#params</td>
<td>P-values for T-statistics</td>
</tr>
<tr>
<td>@GRAD</td>
<td>vector</td>
<td>#params</td>
<td>Gradient of log likelihood at convergence</td>
</tr>
<tr>
<td>@VCOV</td>
<td>matrix</td>
<td>#params*#params</td>
<td>Variance-covariance of estimated coefficients</td>
</tr>
</tbody>
</table>
ORDPROB

@FIT series #obs Fitted values of dependent variable

If the regression includes a PDL or SDL variable, the following will also be stored:

@SLAG scalar 1 Sum of the lag coefficients
@MLAG scalar 1 Mean lag coefficient (number of time periods)
@LAGF vector #lags Estimated lag coefficients, after "unscrambling"

Method

Like the binary Probit model, the Ordered Probit model is based on an unobserved continuous dependent variable \( y^* \). The model is

\[
y^* = XB + e.
\]

Instead of \( y^* \), we observe a category value \( Y \), where a larger category value implies a larger value of \( y^* \). In binary Probit, the category values are 0 for \( y^* < 0 \), and 1 for \( y^* > 0 \). In Ordered Probit, more than 2 category values are usually involved. The category values need not be consecutive, and the lowest category does not have to be 0. The boundary values between the different categories are estimated parameters (\( \mu \)s). The lowest effective boundary value (\( \mu_1 \)) is normalized to 0, just as in binary Probit.

For example, suppose there are 3 categories, with category values 0, 1, and 2:

\[
egin{align*}
Y &= 0 \text{ if } \mu_0 <= XB + e < \mu_1 \text{ (} \mu_0 = \text{-infinity, and } \mu_1 = 0) \\
Y &= 1 \text{ if } \mu_1 <= XB + e < \mu_2 \text{ (} \mu_2 \text{ is an estimated parameter)} \\
Y &= 2 \text{ if } \mu_2 <= XB + e < \mu_3 \text{ (Note: } \mu_3 = \text{infinity)}
\end{align*}
\]

The \( \mu \)s are always given names based on the category value for which they are the lower bound -- \( \mu_2 \) in the example above is the lower bound for category with value 2. \( X \) normally includes a constant term (C), which can be though of as a replacement for \( \mu_1 \); in this case, the other \( \mu \)s can be interpreted as being measured relative to the value of C. The estimated \( \mu \) values are constrained to follow a strict ordering (\( \mu_0 < \mu_1 < \mu_2 \), etc.). Negative and non-integer category values are not allowed. Just recode such values to integers (preserving the proper ordering).
ORDPROB

ORDPROB uses analytic first and second derivatives to obtain maximum likelihood estimates via the Newton-Raphson algorithm. This algorithm usually converges fairly quickly. TSP uses zeros for starting parameter values, except for the constant term and the MUs. @START can be used to provide different starting values (see NONLINEAR). Multicollinearity of the independent variables is handled with generalized inverses, as in the other linear and nonlinear regression procedures in TSP.

If you wish to estimate a nonstandard ordered probit model (e.g. adjusted for heteroskedasticity or with a nonlinear regression function), use the ML command. See our website for an example.

Before estimation, ORDPROB checks for univariate complete and quasi-complete separation of the data and flags this condition, because the model is not identified in this case. Without this check, one or more RHS variables perfectly predict the dependent variable for some observations, and their coefficients would slowly iterate to plus or minus infinity.

The Scaled R-squared is a measure of goodness of fit relative to a model with just a constant term; it is a nonlinear transformation of the Likelihood Ratio test for zero slopes. See Estrella (1998). Although the paper is concerned with dichotomous dependent variables, the scaled R-squared applies to any model with a fixed number of categories, such as Ordered Probit and Multinomial Logit.

Options

See the NONLINEAR section of this manual for the usual nonlinear options.

Example

Ordered Probit regression of patents on lags of log(R&D), science sector dummy, and firm size:

```
ORDPROB PATENTS C LRND LRND(-1) LRND(-2) DSCI SIZE;
```

References


ORTHON

Example

ORTHON orthonormalizes an arbitrary matrix and saves the orthonormalizing transformation. The columns of the resulting matrix span the same space as the columns of the original matrix, but are orthonormal (orthogonal and scaled so that their Euclidean norm is one).

ORTHON <input matrix> <triangular matrix> <orthonormalized matrix>;

Usage

The input matrix $X$ is a general NROW by NCOL matrix. ORTHON obtains a triangular matrix $S$ of order NROW such that $X'X = S'S$. It uses the inverse of $S$ to transform $X$ by postmultiplying it. $S$-inverse and the orthonormalized $X$ are returned in the second and third arguments to the procedure.

ORTHON transforms a data matrix $X$ as it is done in TSP's regression calculation to obtain more accurate results. Even if $S$ is not determined very accurately due to inaccuracy in forming the cross product matrix $X'X$, a regression run on the transformed $X$s and then untransformed will produce extremely accurate results, since the actual matrix inversion is performed on an $X'X$ matrix from which most collinearity has been removed.

Method

ORTHON forms $X'X$ from the $X$ matrix, factors it using the Choleski factorization algorithm, inverts the result using the method described in MATRIX, and postmultiplies $X$ by the resulting upper triangular matrix.

Example

The following example shows how to use ORTHON in programming ordinary least squares explicitly in TSP:

```
MMAKE X C X1 X2 ;
ORTHON X S XTILDA ;
MAT XTXINV = (XTILDA'XTILDA)" ;
MAT BETA = S'XTXINV'XTILDA'Y ;
MAT XXINV = S'XTXINV'S" ;
```

The resulting BETA and XXINV are estimates of the untransformed coefficients and the inverse of the $X'X$ matrix.
OUT (Databank)

Examples

OUT specifies a list of external files on which all TSP variables created or modified will be stored.

\[ OUT \text{<list of filenames> or 'filename strings'} ; \]

Usage

Follow the word OUT with the names of the TSP databank(s) on which you wish to store your variables. On most computers, these are binary .TLB files. Up to 8 databanks may be active for output at one time.

After the OUT statement in your program, TSP marks any of the variables you modify or create so they will be stored on the databank files at the end of the run. Variables created before the OUT statement was executed and not modified later will not be stored. OUT remains in effect until another OUT statement is encountered. To stop writing data to any files, include an OUT statement with no arguments to cancel the previous statement; this will also cause the variables to be stored on the previous OUT file.

When time series are stored with an OUT statement, the whole series is stored, rather than just the observations in the current sample. The frequency of the run where you use the series later should be the same as the frequency of the run when the series was stored.

Since all variables to be saved on databanks are actually saved only upon execution of a new OUT statement, or at the end of your TSP run, the variables marked by the last OUT statement will not be stored if the run later aborts for any reason.

Output

OUT produces no printed output, except a message when a new databank is created.

Examples

\[ OUT \text{FOO} ; \]
\[ \text{? creates } \text{EXP.TLB in the } \text{C:\CONSUME directory on a PC} \]
\[ OUT \text{'C:\CONSUME\EXP'} ; \]

Also see the examples under the KEEP command.
OUTPUT (Interactive)

Examples

OUTPUT sends all subsequent output to a specified external file rather than to the terminal.

OUTPUT [<filename> or 'filename string'] ;

Usage

You may use OUTPUT to save the results of your entire terminal session, or to select portions for subsequent printing or review (plots, graphs, regression results). This command will stay in effect until you restore the output stream to the terminal with a TERMINAL command. It is not possible to send results to the screen and output file simultaneously, but warning and error messages will be displayed in both places as they occur.

OUTPUT will take only one filename as an argument, and if it is not in quotes, this filename must conform to restrictions placed on TSP variable names, i.e. it must be limited to eight characters, and the filename extension must be omitted. If the filename is provided on the command line, the extension .OUT will be assumed. If the filename is absent, you will be prompted for it -- in this case you may specify a directory other than the current as well as an extension or disk unit, the only limit is that the whole name must be 32 characters or less. Again, if the extension is omitted, .OUT will be assumed.

You may switch back and forth between your output file and TERMINAL, or between any number of output files as much as you like. If a file is found to exist already when you open it with the OUTPUT command, subsequent output will be appended to it rather than creating a new file.

You may view any output you've sent to a disk file by using the SYSTEM command. This feature enables you to give operating system commands, so you can EDIT the contents of your file, or call your favorite editor to display it for you.

Typing CONTINUE will return you to your interactive TSP session without any loss of continuity.

Note: In order to see the contents of your currently open output file, you must close it before giving the SYSTEM command. Otherwise your output will appear to be missing from the file. The TERMINAL command, or opening a new OUTPUT file will close the current file; you may reopen the original file when you return from SYSTEM and output will continue to be appended to it.
Examples

13? ? Sending regression output to a file
14? OLSQ Y C X Z ;
15? OUTPUT YXZ ;
16? EXEC 14
17? TERM

In addition to any output files you have stored results in, you may wish to
document your session before you quit:

74? OUTPUT AUG2385
75? ?
75? ? Interactive TSP session on Aug 23, 1985 -- RSS
75? ?
75? ? comments about results, files used, etc....
75? ?
75? ? REVIEW ? photo of session
76? ?
76? ? display of symbols created during session sorted
76? ? into classes
76? ?
76? ? SHOW SERIES,EQUATION,MATRIX,PROC
77? EXIT

These commands and comments will be written to the disk file as well as the
resulting output. You may choose to document just significant points by
REVIEWing specified ranges, or EXECing important results. The comment
delimiter "?" may be used freely to make output files more readable. Of
course, TSP automatically provides the file BKUP.TSP as an undocumented
session photo.
PAGE

PAGE is used to force the paging of the printed output of TSP. It only operates when the option HARDCOPY is in effect.

PAGE ;

Usage

Include a PAGE statement any place in your TSP program where you wish to force the printed output to start on a new page. This might be at the beginning of a series of regressions, or when doing a large simulation, or just to force a new title to be printed.

Normally TSP pages the output as well as it can so that the major procedures start on a new page, but it is not always possible to format exactly as the user would want without wasting a large amount of blank paper; the PAGE statement lets you control the printing to a certain extent.

Output

PAGE produces no printed output itself but causes paging to the next page to occur and the title line to be printed.
PANEL obtains estimates of linear regression models for panel data (several observations or time periods for each individual). Total, between groups, within groups, and variance components may be obtained. In addition one and two-way random effects models may be estimated by maximum likelihood. The data may be unbalanced (different number of observations per individual). PANEL can also compute means by group and perform F tests between groups.

**PANEL** *(ALL, BETWEEN, BYID, FEPRINT, HCOMEGA=BLOCK or DIAGONAL, HCTYPE=0 or 1, ID=<id series>, MEAN, PRINT, REG, RE, REIT, ROBUST, SILENT, T=<number of time periods>, TERSE, TIME=<time series>, TOTAL, VARCOMP, VBET=<between variance>, VSMALL, VWITH=<within variance>, WITHIN, Nonlinear options) <dependent variable> <list of independent variables> ;

**Usage**

The basic PANEL statement is like the **OLSQ** statement: first list the dependent variable and then the independent variables. C is optional; an intercept term is central to these models and will be added if it is not present. You may have as many independent variables as you like subject to the overall limits on the number of arguments per statement and the amount of working space, as well as the number of data observations you have available. The observations over which the models are computed are determined by the current sample. PANEL treats missing values, lags, and leads correctly. That is, lags and leads are applied only within an individual.

Your data must be set up with all the time periods for each individual together. Additionally, you must specify when the observations for one individual end and data for the next individual begins. The default method is to provide a series named @ID which takes on different values for each individual. If your data are balanced (the same number of time periods for every individual), the **T=** option can be used. If the data are not in this order, the **SORT** command can be used to reorder them; you could also sort the data by year and then individual if you wish to do variance components in the time dimension. Usually it is best to use the **FREQ** (PANEL) command at the top of your run to specify such ID variables, internal frequency and starting date, etc. Then these options will be used for all PANEL, **AR1**, **GENR**, etc. commands within the run.
The models you wish to estimate are specified in the options list. The default is to estimate the total, between, within, and variance components models. For the VARCOMP (random effects) model, there are additional options that specify how to compute the variance components. Small- or large sample formulas may be used, or the user can supply the values directly. If negative variances are computed using the small sample method, the method switches over to the large sample formulas, which always result in positive values. PANEL also computes a Hausman test for correlated effects by comparing the WITHIN (fixed effects) and VARCOMP (random effects) estimators.

The REI and REIT options are used to obtain maximum likelihood estimates of the one and two-way random effects models.

Output

The output begins with a title and a summary of the panel structure: number of individuals (NI), number of time periods (T), and total number of observations (NOB). If the data are unbalanced, TMIN and TMAX will be printed. For each estimator, a table of regression coefficients and their standard errors is printed, along with name of the dependent variable, the sum of squared residuals, standard error of the regression, mean and standard deviation of the dependent variable, R-squared, and adjusted R-squared.

Other output varies by estimator. If the data are unbalanced, the Ahrens-Pincus measure of the degree of unbalancedness is also printed; this measure is one for balanced data; values less than one provide an indication of how far the data is from balanced. See the method section for the definition of this statistic and the reference for details on its interpretation.

MEAN prints a table of means for each individual. @MEAN (#obs*#vars) is stored, and excludes any constant term.

BYID prints an F test vs. TOTAL (labelled F-stat for A,B= Ai,Bi), and an F test vs. WITHIN (labelled F-stat for Ai,B= Ai,Bi), in the output of the respective estimators. Only @COEFI (the individual coefficient estimates), @LOGLI, and @SSRI (the individual sum or squared residuals) are stored. Use the PRINT option to print @COEFI.

WITHIN prints an F test vs. TOTAL (labelled F-stat for A,B= Ai,B), and stores @FIXED effects vector.

VARCOMP prints the actual variance components, the method used to compute them, and the implied differencing factor (THETA). A Hausman specification test comparing VARCOMP (null hypothesis) and WITHIN is computed.
PANEL

PANEL stores the standard regression results in data storage for later use using @names, but with B, T, V, W, REI, and REIT appended to distinguish between the different estimators. For example, @COEFW is the within coefficients, @RESW are the within residuals, and @SSRV is the sum of squared residuals from VARCOMP. @RESB is a matrix.

In the table below, #vars is equal to the number of right hand side variables plus one (for the constant) for the T, B, W, and V estimators. For the REI estimator, #vars includes the estimate of RHO_I (the within group correlation) and SIGMA2 (the total standard error). For the REIT estimator, #vars includes the estimate of RHO_I, the estimate of RHO_T, the within time correlation, and SIGMA2 (the total standard error).

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@LHV</td>
<td>list</td>
<td>1</td>
<td>Name of the dependent variable</td>
</tr>
<tr>
<td>@SSRT/I/B/W/V/REI/REIT</td>
<td>scalar</td>
<td>1</td>
<td>Sum of squared residuals (@SSRI=BYID, etc.)</td>
</tr>
<tr>
<td>@S2T/B/W/V/REI/REIT</td>
<td>scalar</td>
<td>1</td>
<td>Variance of residuals (@S2B=BETWEEN, etc.)</td>
</tr>
<tr>
<td>@ST/B/W/V/REI/REIT</td>
<td>scalar</td>
<td>1</td>
<td>Standard error of the regression</td>
</tr>
<tr>
<td>@YMEANT/B/W/V/REI/REIT</td>
<td>scalar</td>
<td>1</td>
<td>Mean of the dependent variable</td>
</tr>
<tr>
<td>@SDEVT/B/W/V/REI/REIT</td>
<td>scalar</td>
<td>1</td>
<td>Standard deviation of the dependent variable</td>
</tr>
<tr>
<td>@NOB</td>
<td>scalar</td>
<td>1</td>
<td>Number of observations</td>
</tr>
<tr>
<td>@APUI</td>
<td>scalar</td>
<td>1</td>
<td>Ahrens-Pincus unbalancedness in i</td>
</tr>
<tr>
<td>@SPUT</td>
<td>scalar</td>
<td>1</td>
<td>Ahrens-Pincus unbalancedness in t</td>
</tr>
<tr>
<td>@RSQT/B/W/V</td>
<td>scalar</td>
<td>1</td>
<td>R-squared</td>
</tr>
<tr>
<td>@ARSQ/B/W/V</td>
<td>scalar</td>
<td>1</td>
<td>Adjusted R-squared</td>
</tr>
<tr>
<td>@NCOEFT/B/W/V/REI/REIT</td>
<td>scalar</td>
<td>1</td>
<td>Number of coefficients</td>
</tr>
<tr>
<td>@NCIDT/B/W/V/REI/REIT</td>
<td>scalar</td>
<td>1</td>
<td>Number of identified coefficients (number with non-zero standard errors)</td>
</tr>
<tr>
<td>@LMHETT/W/V</td>
<td>scalar</td>
<td>1</td>
<td>LM heteroskedasticity</td>
</tr>
</tbody>
</table>
%LMHETT/W/V scalar 1 test P-value of LM heteroskedasticity test
@DWT/W/V scalar 1 Durbin-Watson autocorrelation test
%DWUT/W/V scalar 1 Upper bound on P-value of DW
%DWLT/W/V scalar 1 Lower bound on P-value of DW
@LOGLT/I/W/REI/REIT scalar 1 value of the log likelihood
@SBICT/W/REI/REIT scalar 1 Schwarz-Bayes information criterion
@AICT/W/REI/REIT scalar 1 Akaike information criterion
@HAUS scalar 1 Hausman test value
%HHAUS scalar 1 Hausman test p-value
@HAUSDF scalar 1 Hausman test degrees of freedom
@RNMST/B/W/V/REI/REIT list #vars List of names of right hand side variables
@COEFT/I/B/W/V/REI/REIT vector #vars Coefficient estimates
@SEST/B/W/V/REI/REIT vector #vars Standard errors
@TT/B/W/V/REI/REIT vector #vars T-statistics
@COEFAI vector #individuals Estimated fixed effects
@SESAI vector #individuals Standard errors on fixed effects
@TAI vector #individuals t-statistics on fixed effects
%TAI vector #individuals p-values associated with @TAI
@AI series #obs Fixed effect estimates as a series
@VCOVT/B/W/V/REI/REIT matrix #vars*#vars Variance-covariance of estimated coefficients
@REST/I/B/W/V/REI/REIT series #obs Residuals = actual - fitted values of the dependent variable.
Method

The model estimated is

\[ y_{it} = X_{it} \beta + a_i + u_{it} \]

PANEL computes means for each variable by individual. These are used directly in the BETWEEN regression. WITHIN subtracts the individual means from each variable and runs a regression on this transformed data (any variables which are constant over time for every individual are not identified).

VARCOMP does a transformation similar to WITHIN. \((1 - \sqrt{\theta})\) times the mean is subtracted from each variable (including the constant term), where \(\theta\) is given by

\[ \theta = \frac{\text{Var}(u)}{\text{Var}(u) + T \text{Var}(a)} \]

\(T\) does not have to be the same for each individual. The small and large sample formulas used for the variance components are:

<table>
<thead>
<tr>
<th>variance</th>
<th>small sample</th>
<th>large sample</th>
</tr>
</thead>
<tbody>
<tr>
<td>within</td>
<td>@SSRW/(NOB-NX-NI)</td>
<td>@SSRW/NOB</td>
</tr>
<tr>
<td>total</td>
<td>@SSRT/(NOB-NX-1)</td>
<td>(not used)</td>
</tr>
<tr>
<td>between</td>
<td>VTOT-VWITH</td>
<td>(@SSRT-@SSRW)/NOB</td>
</tr>
</tbody>
</table>

If the small sample formula produces a non-positive variance, PANEL switches over to the large sample formulas automatically. The large sample formulas are asymptotically correct if \(T\) is (becomes) large relative to \(NI\) (not usually the case); otherwise they will be biased. Note that if \(\theta=1\), this corresponds to a zero between variance and VARCOMP will produce the same estimates as TOTAL. If \(\theta=0\), this corresponds to a zero within variance, and VARCOMP will be the same as WITHIN.

For each F test (described under Output), a P-value and an alternative critical value are printed. The critical value has a size which becomes smaller as the number of observations grows -- this is an alternative to the conventional testing procedure, which is certain to reject all point null hypotheses when sample sizes become large. It is based on a Bayesian flat prior, and computed from the formula in the Leamer reference:

\[ F_{\text{crit}} = \left( \frac{T - k}{p} \right) \left( T^{\theta/7} - 1 \right) \]
Where \( T \) = total number of observations, \( k \) = number of estimated parameters in the unrestricted model, and \( p \) = the number of restrictions.

All regressions are computed with the standard orthonormalized data matrices to insure accurate coefficients and variance estimates under possible multicollinearity (methods using moment matrices are less accurate).

The Durbin-Watson test and bounds on its P-values are computed following the Bhargava et al reference, extended to the unbalanced data case. The P-values are computed using the Farebrother-Imhof method, since there can be multiple equal eigenvalues.

The REI estimates are obtained with a grid search over RHO_I in order to avoid the problem of multiple local optima. Estimates are then refined to choose the global optimum and multiple optima are reported. RHO_I is bounded between \(-1/(\text{Max}(T)-1)\) and 1, where Max(T) is the maximum number of observations per individual. See Maddala and Nerlove (1971). The REIT estimates are obtained using the method of Davis (2002). The Ahrens-Pincus measure of unbalancedness in dimension \( i \) is defined as follows:

\[
\text{AP}_i = \frac{N}{\bar{T} \sum_{i=1}^{N} 1/T_i} \quad \text{where} \quad \bar{T} = \frac{1}{N} \sum_{i=1}^{N} T_i
\]

This can be interpreted as the ratio of the harmonic and arithmetic means of the \( T(i) \) over the sample of individuals. Note that \( AP \) is always less than or equal to 1 and that it equals one only when \( T(i)=T \) for all \( i \).

**Options**

**ALL/NOALL** turns all regressions on or off (equivalent to the combination of TOTAL, BETWEEN, WITHIN, VARCOMP, REI, REIT).

**BETWEEN/NOBETWEEN** selects the "between" estimator -- a regression on the means for each individual.

**BYID/NOBYID** does a separate regression for each individual, and computes F tests for equality with the TOTAL and WITHIN estimators.

**FEPRINT/NOFEPRINT** specifies that the fixed effect estimates are to be printed as well as stored.
**HC** OMEGA = BLOCK or DIAGONAL specifies the form of the $\Omega = E[uu']$ matrix to use when computing ROBUST standard errors. Ordinarily, the default is BLOCK for PANEL, which allows for cross-time correlation of the disturbances within individuals. This feature can be used for any kind of grouped data, simply by ensuring that the relevant PANEL setup has been defined.

**HCTYPE** = 0 or 1 specifies whether to apply a degrees of freedom correction to the robust s.e.s (0 is no and 1 is yes).

**ID**= the name of a series which takes on a different value for each individual. The default is @ID; alternatives are the T= and TIME= options.

**MEAN/NOMEAN** causes the means for each individual to be printed in a table. This can be used in conjunction with the NOREG option to print means only (to suppress all the default regression models). These individual means are stored in the NI x (1+NX) matrix @MEAN, where the first column is the dependent variable.

**PRINT/NOPRINT** prints @COEFI in conjunction with BYID, and prints @FIXED for within.

**REG/NOREG** is used with the MEAN option above. To suppress some regression models, but print others, use the individual options -- NOBETW to suppress the BETWEEN output, etc.

**REI/NOREI** specifies that ML estimates of the one-way random effects model are to be obtained. @START may be used to supply starting values.

**REIT/NOREIT** specifies that ML estimates of the two-way random effects model are to be obtained. This requires the TIME= option for unbalanced data in FREQ(PANEL). @START may be used to supply starting values.

**ROBUST/NOROBUST** calculates heteroskedasticity-robust standard errors (HCTYPE=1; see OLSQ) for the WITHIN coefficients. If this option is used, the Hausman test comparing WITHIN and VARCOMP is not computed.

**SILENT/NOSILENT** can be used to turn off all the regression output.

**T**= the number of time periods for each individual (for balanced data only). For unbalanced data, use the ID= option.

**TERSE/NOTERSE** can be used to turn off most of the regression output, except the coefficients and standard errors.
**PANEL**

**TIME** = the name of a time period series which increases in value for each individual and decreases between individuals. Alternatives are the **ID=** and **T=** options. Example: **TIME=YEAR**. This is not considered sufficient for identifying individuals, since the last time period for one individual may be less than the first time period of the next individual.

**TOTAL**/**NOTOTAL** selects the "total" or "pooled" estimator -- a plain OLS regression on the whole sample.

**VARCOMP**/**NOVARCOMP** selects the "variance components" or "random effects" estimator. The method of selecting the variance components is controlled with the **VBET**, **VSMALL**, and **VWITH** options described below. Unbalanced data are not a problem. For variance components in the time dimension, use the **REIT** option, or sort your data by time period and use time as the ID.

**VBET=** specifies the value of the "between" variance for **VARCOMP**.

**VSMALL**/**NOVSMALL** selects the small sample variance components formulas for **VARCOMP** (as opposed to the large sample formulas). Small sample formulas are unbiased but can result in negative variances, while large sample formulas are biased but always yield positive variances. To supply your own variance values, use **VBET=** and **VWITH=**.

**VWITH=** specifies the value of the "within" variance for **VARCOMP**.

**WITHIN**/**NOWITHIN** selects the "within" or "fixed effects" estimator (different intercepts for each individual).

Nonlinear options may be used for the **REI** and **REIT** estimators. See **NONLINEAR**.

**Examples**

Global **FREQ** (PANEL) command, with ID variable to identify individuals:

```
FREQ (PANEL,ID=FIRM); DY=Y-Y(-1);
PANEL DY C X X(-1);
```

Estimate all models (7 years per individual, balanced data), and print individual means:

```
PANEL (T=7,MEAN,BYID) LRNDL5 C PATENTS LRNDL4;
```

Print **VARCOMP** output only, using @ID or **FREQ**(PANEL) to distinguish individuals:

```
PANEL (NOTOT,NOBET,NOWITH) LRNDL5 C PATENTS LRNDL4;
```
Estimate all models except BYID, use large sample formulas for VARCOMP:

**PANEL (NOVSMALL) LRNDL5 C PATENTS LRNDL4;**

Print individual means only:

**PANEL (MEAN, NOREG) LRNDL5 C PATENTS LRNDL4;**

**References**


StatLib, [http://lib.stat.cmu.edu/apstat/](http://lib.stat.cmu.edu/apstat/)
PARAM

Example

PARAM is used to define parameters for the nonlinear estimation procedures and to assign starting values to them. To supply parameter starting values to PROBIT, TOBIT, SAMPSEL, and LOGIT, use the @START vector; see those procedures for further information.

PARAM <parameter name> [<value> <parameter name> <value> ....] ;

Usage

PARAM may be followed by as many argument pairs as desired (limited only by TSP's argument limit). Each pair is the name of the parameter followed by the value it is to be given. The parameter names may be that of new or previously defined variables. The value may be omitted, in which case the variable is given the value zero if it is new, or left unchanged if it has already been defined. Note: the keywords C and CONSTANT may not be used as parameters.

Procedures which estimate values for parameters defined by the PARAM statement are LSQ for nonlinear single and multi-equation least squares (including minimum distance estimators) and FIML. All other procedures treat parameters like constants (scalar variables) which have the arithmetic value they have been assigned, either by a PARAM statement or by later estimation. FORM (PARAM) can also be used to create parameters.

PARAM ignores any () or *** in the command. This is useful for pasting back in starting values of the parameters from a previous estimation.

Output

PARAM produces no printed output; it stores the variables named in data storage with a type equal to parameter.

Example

A common problem in nonlinear estimation is that one or more parameters may enter the model in a highly nonlinear fashion, making it difficult to estimate unless you have good starting values. In this example, we estimate a subset of the parameters of a model conditional on the value of another parameter, DELTA, and then reestimate with all the parameters free:

FRML INVEQ I = LAMBDA^1(-1) + ALPHA*GNP/(DELTA+R) ;
PARAM LAMBDA ALPHA ;
CONST DELTA 15 ;
When the second LSQ is done, the starting values for LAMBDA and ALPHA will be those determined by the first estimation, while the starting value for DELTA will be 15, which it was assigned by the `CONST` statement.
A PDL (polynomial distributed lag) variable specification may be used for a right hand side variable in any linear estimation procedure (OLSQ, INST, LIML, AR1, PROBIT and TOBIT). It constrains the coefficients on the lags of that variable to lie on a polynomial of the degree specified by the user. The Shiller lag, available with OLSQ only, uses an additional argument to relax this constraint somewhat.

PDL lag variables have the following form:

\[ \text{varname}(<\text{degree}>, <\# \text{lags}>, \text{NONE or FAR or NEAR or BOTH}) \]

Shiller lag variables add the \textit{XIPRIOR} argument:

\[ \text{varname}(<\text{degree}>, <\# \text{lags}>, \text{NONE or FAR or NEAR or BOTH, XIPRIOR}) \]

### Usage

You may include a PDL (polynomial distributed lag) specification anywhere in the list of right hand side variables in a linear estimation procedure. There is no explicit limit on the number of right hand side variables which may contain a PDL specification. The form of a PDL variable is the name of the variable whose lags you want in the model, followed by parentheses containing three items: the number of terms in the polynomial (the degree plus one), the number of lags of the variable to be included (including the zeroth lag), and what kind of end-point constraint you place on the polynomial.

The polynomial distributed lag is a method for including a large number of lagged variables in a model, while reducing the number of coefficients which have to be estimated by requiring the coefficients to lie on a smooth polynomial in the lag. The purpose of the constraints is to force the lag coefficients at either end of the lags over which you are estimating to go to zero, that is, the NEAR constraint forces the coefficient of the first lead to zero, while the FAR constraint forces the coefficient of the lag one past the last included lag to zero. The BOTH and NONE constraints are self-explanatory.
The number of coefficients which are estimated for a PDL variable are the number of terms in the polynomial less the number of constraints. This must be positive and less than or equal to the number of lags in the unconstrained model. For a long, totally unconstrained distributed lag, an implicit list (-) is best. For example, the following three statements are equivalent:

\[ \text{OLSQ CONS72 C GNP72 GNP72(-1)-GNP72(-15);} \]
\[ \text{OLSQ CONS72 C GNP72(16,16,NONE);} \]
\[ \text{OLSQ CONS72 C GNP72(-1) GNP72(-2) GNP72(-3) GNP72(-4)} \]
\[ \text{ GNP72(-5) GNP72(-6) GNP72(-7) GNP72(-8) GNP72(-9) GNP72(-10)} \]
\[ \text{ GNP72(-11) GNP72(-12) GNP72(-13) GNP72(-14) GNP72(-15);} \]

For Shiller lags, the additional argument specifies a prior for the variance of the differenced coefficients - smaller values imply coefficients that are "smoother." A value for the prior equal to zero is equivalent to simply using PDL. A very large value will yield unconstrained lag coefficients.

**Output**

The coefficients of PDL variables are estimated by forming linear combinations of the underlying lagged variables and including these variables in the regression. The estimates of these coefficients are not easily interpreted, and so TSP "unscrambles" these results for each PDL variable after the regression and presents the estimates in terms of the original variable and its lags.

The results begin with a title 'Distributed Lag Interpretation for: variable name,' followed by the estimated mean lag and its standard error, computed as the average lag weighted by the lag coefficients. If any of the lag coefficients are less than zero, this quantity does not have very much meaning. The estimated sum of the lag coefficients and its standard errors are also shown. Both these standard errors are computed by taking into account the covariance of the estimates of the lag coefficients.

Following these summary statistics a table and a plot of the individual lag coefficients is printed. This table also shows the standard errors of the estimates and plots standard error bands around the lag coefficients. These coefficients are also stored in data storage with the following names:

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@SLAG</td>
<td>scalar</td>
<td>1</td>
<td>Sum of the lag coefficients</td>
</tr>
<tr>
<td>@MLAG</td>
<td>scalar</td>
<td>1</td>
<td>Mean lag</td>
</tr>
<tr>
<td>@LAGF</td>
<td>vector</td>
<td>#lags</td>
<td>Estimated lag coefficients after &quot;unscrambling&quot;</td>
</tr>
<tr>
<td>@PDL1,2,etc</td>
<td>series</td>
<td>#obs</td>
<td>Scrambled right hand side variables</td>
</tr>
</tbody>
</table>
Method

PDL estimates are obtained by forming linear combinations of the underlying variable and its lags, with weights determined by the order of the lag polynomial and the value of the constraint options. These "scrambled" variables are used as regressors, and then "unscrambled" after estimation to obtain the implied lag coefficients. Further details are given in the TSP User's Guide and Almon (1965). The method TSP uses is described in Cooper (1972). It uses Lagrangian interpolation polynomials that are orthogonal, in order to minimize multicollinearity problems.


Examples

**OLSQ I,C,GNP(4,16,FAR) ;**  
specifies a distributed lag on GNP which covers 16 periods where the lag coefficients are constrained to lie on a third degree polynomial and go to zero at the 16th lag.

**OLSQ I C GNP(4,16,FAR) R(4,24,NEAR) ;**  
adds another distributed lag in R which covers 24 periods and is constrained to go to zero at the first lead. PDL variables can also be used with INST and AR1:

**INST I C GNP(3,5,NONE) INVR C LM LM(-1) LM(-2) ;**  
**AR1 (METHOD=CORC) CONS C GNP (3,5,NONE) ;**

The instrumental variable estimation specifies just enough instruments for the number of independent variables which will appear in the estimation: the constant and 3 weighted combinations of GNP and its lags.

References


PLOT

Output  Options  Examples

PLOT produces a plot of one or more series versus the observation number (usually in units of time). The series are plotted on the horizontal axis and time on the vertical axis. The user has a good deal of freedom in formatting this plot with options. For DOS, Windows, unix, or MAC PC with the graphics version of TSP, see the entry for PLOT (graphics version).

PLOT (BAND=STANDARD or <series name>, BMEAN, BMID, BOX, HEADER, ID, INTEGER, LINES=(list of values), MAX=<y-axis maximum>, MEAN, MIN=<y-axis minimum>, ORIGIN, RESTORE, VALUES) <series name> <plotting character> [<series name> <plotting character>.........] ;

Usage

PLOT is followed by a series name, the character to use in plotting the series, possibly a second series name and a second character, and so on. Up to nine series may be plotted. The characters may be anything except $ ; ."":

Parameters that control the appearance of the plot may be specified in an options list in parentheses following the word PLOT. These parameters all have default values, so you do not need to specify them if you just want a simple plot.

Any observations with missing data are excluded from the plot.

Output

PLOT prints a title, followed by the names of all the series being plotted and the characters used to plot them. If there are lines drawn on the plot, a message giving the locations of the lines is printed.

The plot itself is labelled at its four corners with the horizontal minima and maxima; the axes are labelled at several points if the HEADER option was specified, and the mean is marked with an M if a line at the mean was requested. The ID series labels the left hand side of the vertical axis and the values of the first series are on the right hand side if the VALUES option was specified.

If more than one series is being plotted, any points which are superimposed are plotted with the number of series which have that value instead of the plotting character. The plotting characters of the duplicate series are shown on the right hand side of the plot.
PLOT

PLOT uses the LIMPRN option to decide how wide to make the plot, so you have some control over the format by use of the OPTIONS command.

Options

**BAND** = STANDARD or `seriesname` specifies the name of a series which is used as the width of a band to be printed around the observations of the first series to be plotted (usually this series is a set of computed standard errors). The keyword STANDARD will cause the standard deviation of the series to be used as the band. The default is not to plot a band.

**BMEAN/NOBMEAN** causes the band to be printed about the series mean.

**BMID/NOBMID** causes the band to be printed about the midpoint of the plot.

**BOX/NOBBOX** draws a box around the plot.

**HEADER/NOHEADER** causes the horizontal axis to be labelled at equispaced intervals.

**ID/NOID** causes a vertical ("time") axis to be labelled on the left hand side with the ID series.

**INTEGER/NOINTEGER** causes the numeric labels on the horizontal axis to be rounded to the nearest integer value; this improves readability of the plot.

**LINES** = *(list of up to 9 numeric values)* - specifies points along the horizontal axis at which vertical lines will be drawn.

**MAX** = the maximum value on the horizontal axis. If not specified, the maximum value of all the series to be plotted is used.

**MEAN/NOMEAN** draws a vertical line from the mean of the series on the horizontal axis.

**MIN** = the minimum value on the horizontal axis. If not specified, the minimum value of all the series to be plotted is used.

**ORIGIN/NOORIGIN** causes a vertical line to be drawn starting at zero on the horizontal axis.

**RESTORE/NORESTORE** causes the options to be set to their default values.

**VALUES/NOVALUES** causes the value of each observation of the first series to be printed on the right hand side of the plot.
The list of options is obviously extensive and, to make things easier for the user, a set of default options has been chosen which produce a plot of attractive appearance. These options are

\textit{PLOT(LINES=none, BAND=none, NOORIGIN, BOX, NOMEAN, ID, NOINTEGER, VALUES, NOHEADER)}

For convenience, the options of PLOT which are set by you are retained in the next PLOT(s) until they are overridden either explicitly or by including the option RESTORE in the list. RESTORE causes the options to be reset at their default values.

\textbf{Examples}

\texttt{PLOT GNP,*,CONS,X;}
\texttt{PLOT(MIN=500,MAX=1500,LINES=(1000)) GNP G GNP S H CONS C CONSS D ;}
\texttt{PLOT(MIN=-25, MAX=25, BMEAN, HEADER, VALUES, BAND=STANDARD, INTEGER) RESID * ;}
PLOT (graphics version)

Output  Options  Examples

PLOT produces a plot of one or more series versus the observation number (usually in units of time). The series are plotted on the vertical axis and time on the horizontal axis. The plot may be printed as well as displayed if a hardcopy device such as a Laserjet or dot matrix printer is available. This section describes the graphics version of PLOT, which is available only for TSP/Oxmetrics, DOS/Win TSP, unix, and MAC TSP. For other versions, see the non-graphics PLOT command.

\[ \text{PLOT (A4, DASH, DEVICE=\{name of printer\}, FILE=\{name of file\}, HEIGHT=\{height of characters\}, HIRES, LANDSCAP or PORTRAIT, MIN=\{y axis minimum\}, MAX=\{y axis maximum\}, ORIGIN, PREVIEW, SYMBOL, TITLE= 'text string to be used as title', WIDTH) } <\text{list of series names}>; \]

Usage

PLOT is followed by a single series name, or a list of series names. On the plot, the series will be differentiated by colors or the style of the lines used to plot them. Parameters to control the appearance and printing of the plot may be included in parentheses following the word PLOT.

The graph will be displayed on the screen; if a DEVICE= is specified, a prompt is also displayed which instructs you to type "P" if you wish to print the graph. If you type anything else, the graph will not be printed; this is useful if you decide you do not like its appearance after you have seen the screen.

If there are observations with missing data or if the FREQ (PANEL) option is set, there will be breaks in the plotted lines.

Output

A high resolution plot is produced on the screen, with time (the observation index) on the horizontal axis and the series on the vertical access. If there is more than one series, they are differentiated by means of colors (see below for other options).

Options

General  DOS/Win only  MAC only
PLOT (graphics version)

For convenience, the DEVICE=, FILE= AND HEIGHT= options of PLOT that you set are retained in the next PLOT(s) or GRAPH(s) until they are overridden explicitly. They may also be set in a LOGIN.TSP file by not specifying any series to plot.

DASH/NODASH specifies whether the lines for different series on the screen are to be distinguished by using different dash patterns (of which there are seven). The default is no dashes (just color) on the screen and dashes on printed output.

MIN= minimum value for the y-axis. This value must be less than or equal to the minimum value in the data.

MAX= maximum value for the y-axis. This value must be greater than or equal to the maximum value in the data.

ORIGIN/NOORIGIN causes a horizontal line to be drawn starting at zero on the vertical axis.

PREVIEW/NOPREVIE specifies whether the graph is to be shown on the screen before printing or saving. The default is PREVIEW for interactive use and NOPREVIE for batch use. This option is not used when working inside the Oxmetrics shell.

SYMBOL/NOSYMBOL specifies that symbols are to be used for plotting.

TITLE= 'a string which will be printed across the top of the graph'.

DOS/Win only

A4/NOA4 specifies A4 paper size. Available for DEVICE=LJ3 or POSTSCRIPT only.

DEVICE= CHAR or EPSON or LJ2 or LJ3 or LJET or LJPLUS or LJR75 or LJR100 or LJR150 or LJR300 or POSTSCRI or PS specifies the hardcopy device to be used for printer output. LJ means HP LaserJet or compatible, EPSON is EPSON dot matrix or compatible, POSTSCRI and PS are Postscript output, and CHAR is the old line printer output (characters instead of graphics). The LJ suffixes specify models of the printer and the LJR suffixes specify the resolution of the LaserJet printer directly, rather than giving the printer type. The default resolutions for the LJET, LJPLUS, and LJ2 printers are 100, 150, and 300 respectively. Note that larger resolutions imply larger file sizes and printing times.

FILE= the name of a file to which the graphics image is to be written. This file can be printed later; for example, if you are running under DOS and your printer device is LPT1, print the plot with the command
PLOT (graphics version)

copy/b file LPT1:

HEIGHT= letter height in inches. The default is .25. Values in the range (0,1] are valid.

HIRES/NOHIRES controls how graphs are printed in batch mode (when PREVIEW is not being used). Normally (NOHIRES), graphs are printed in character mode to the batch output file. When the HIRES option is used, the patched DEVICE= and FILE= will be used; usually this will send a page to LPT1 for each graph.

LANDSCAP/PORTRAIT specifies the orientation of the plot. On the Mac, specify this option in the dialog box.

MAC only

WIDTH/NOWIDTH specifies whether varying width sizes are to be used to distinguish the lines corresponding to different series on the graph.

Examples

The following example plots the graph shown in the User's Guide:

FREQ Q;
SMPL 53:1 67:4;
LOAD expend approp; ? Original Almon data from Maddala, p. 370
2072 1660 2077 1926 2078 2181 2043 1897 2062 1695 2067 1705 1964
1731 1981 2151
...5715 5412 5637 5465 5383 5550 5467 5465;
PLOT (PORT, PREV, DEV=LJ3, FILE='ALMON.PLT', TITLE='ALMON DATA') EXPEND APPROP;

Here is an example of setting the plot options without actually plotting anything:

PLOT (DEV=LJ3, HEIGHT=.2);
PLOTS

Options  Examples  References

PLOTS turns on the option which produces plots of actual and fitted values and residuals following estimation. The default is not to produce plots. It can also be used to turn on plots of CUSUM and CUSUMQ, which are used to look for "structural change" in a regression.

**PLOTS (PREVIEW) [ALL CUSUM CUSUMSQ] ;**

**Usage**

Include a PLOTS statement in your program before the first regression (OLSQ, AR1, INST, or LSQ) for which you wish to see residual plots. PLOTS remains in force until a NOPLOT statement is encountered.

Note that even though residual plots are not printed, residuals and fitted values will still be stored in data storage. To suppress this feature also, use the OPTIONS NORESID ; statement.

The regression diagnostics are based on the maximum values of the CUSUMs relative to their bounds or mean. They provide a compact alternative to the plot; for example, if a P-value is < .05, the CUSUM crosses a bound.

\[ @CSMAX = \text{max} \left[ .9479 \times \frac{\text{CUSUM}(t)}{\text{CSUB5}(t)} \right] \]
\[ @CSQMAX = \text{max} \left[ \text{CUSUMSQ}(t) - \text{CSQMEAN}(t) \right] \]

The P-value %CSMAX is a function of @CSMAX given in Brown et al (1975). %CSQMAX is a function of @CSQMAX and the degrees of freedom, described in Durbin (1969). This P-value and the critical value for the CUSUMSQ plot are computed from the algorithms given in Edgerton and Wells (1994). The critical values are based on the asymptotic approximation when the number of observations is greater than 60, or on iterating to obtain the exact P-value calculation when the number of observations is less than 60.

All these results are based on recursive residuals, and they will not be calculated if the first \( K \) observations are not of full rank (where \( K \) = number of right hand side variables in the regression). Recursive residuals can also be computed by going backwards through the sample, and although this is not done at present in TSP, it may be useful if the plots are being used to locate points of structural change. Harvey (1990) contains some examples of interpreting the CUSUM and CUSUMSQ plots.

**Options**
PLOTS

PREVIEW/NOPREVIEW turns graphics plots off or on for the following subsequent commands:

- **OLSQ INST, 2SLS** - actual and fitted values, residuals
- **OLSQ** - CUSUM/CUSUMSQ

The default is PREVIEW in TSP/Oxmetrics, and NOPREVIEW in other versions.

Examples

> ? Residual plots are shown only for the second (AR) regression.
> NOPLOT;
> OLSQ CONS C GNP;
> PLOTS; AR1 CONS C GNP;

> ? turn on the CUSUM plots for each regression until a
> NOPLOTS; is given).
> PLOTS CUSUM CUSUMSQ;

> ? turn on all plots (both CUSUMs plus residuals and fitted values).
> PLOTS ALL;

> ? turn on the regression diagnostics.
> REGOPT (PVPRINT) CSMAX CSQMAX;

> ? also include the diagnostics (and P-values).
> REGOPT (PVPRINT) AUTO;

> ? include both the diagnostics and all plots.
> REGOPT (PVPRINT) ALL;

In all versions except TSP/Oxmetrics, the CUSUM plots are done in low resolution with characters, so they can be included in standard output files and implemented on all platforms. The CUSUM plot stores: @CUSUM, @CSUB5%, and @CSLB5%. The CUSUMSQ plot stores: @CUSUMSQ, @CSQMEAN, @CSQUB5%, and @CSQLB5%. These variables can be stored without displaying the low resolution plot if you want to make only high-resolution plots. To do this, use:

`REGOPT(CALC,NOPRINT) CUSUM CUSUMSQ ;`

or

`PLOTS CUSUM CUSUMSQ; SUPRES CUSUM CUSUMSQ ;`

To then make high-resolution plots on the 386/486 and Mac versions of TSP, use:

`PLOT (ORIGIN,PREVIEW) @CUSUM @CSUB5% @CSLB5% ;`
`PLOT (PREVIEW) @CSQMEAN @CUSUMSQ @CSQUB5% @CSQLB5% ;`
TSP only calculates the bounds automatically for the 5% two-tailed tests. Bounds for other size/one-tailed tests can be calculated manually with simple transformations of the bounds which are stored, using the appropriate critical values from a table.

References


POISSON obtains estimates of the Poisson model, where the dependent variable takes on nonnegative integer count values and its expectation is an exponential linear function of the independent variables. In the Poisson model, the variance of the dependent variable equals its mean, which is rarely the case in practice. More general models, where the variance is larger than the mean, are the Negative Binomial types 1 and 2. See the NEGBIN command. The Poisson command is

**POISSON (nonlinear options)** `<dependent variable>` `<list of independent variables>` ;

**Usage**

The basic POISSON statement is like the OLSQ statement: first list the dependent variable and then the independent variables. If you wish to have an intercept term in the regression (usually recommended), include the special variable C or CONSTANT in your list of independent variables. You may have as many independent variables as you like subject to the overall limits on the number of arguments per statement and the amount of working space, as well as the number of data observations you have available.

The observations over which the regression is computed are determined by the current sample. If any of the observations have missing values within the current sample, POISSON will print a warning message and will drop those observations. POISSON also checks that the observations on the dependent variable are integers and are not negative.

The list of independent variables on the POISSON command may include variables with explicit lags and leads as well as PDL (Polynomial Distributed Lag) variables. These distributed lag variables are a way to reduce the number of free coefficients when entering a large number of lagged variables in a regression by imposing smoothness on the coefficients.

**Output**

The output of POISSON begins with an equation title and frequency counts for the lowest 10 values of the dependent variable. Starting values and diagnostic output from the iterations will be printed. Final convergence status is printed.
This is followed by the number of observations, mean and standard deviation of the dependent variable, sum of squared residuals, correlation type R-squared, a test for overdispersion, likelihood ratio test for zero slopes, log likelihood, and a table of right hand side variable names, estimated coefficients, standard errors and associated t-statistics. The default standard errors are the robust/QMLE Eicker-White estimates. These are consistent even for a model whose variance is not equal to the mean, as long as the mean is correctly specified. For most economic data, the overdispersion test rejects the Poisson model, and you may wish to use the Negative Binomial model instead (although as a member of the linear exponential class, the Poisson model with Eicker-White standard errors may be more robust against misspecification even when the data are overdispersed - see Cameron and Trivedi for further information on this point).

POISSON also stores some of these results in data storage for later use. The table below lists the results available after a POISSON command.

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@LHV</td>
<td>list</td>
<td>1</td>
<td>Name of dependent variable</td>
</tr>
<tr>
<td>@RNMS</td>
<td>list</td>
<td>#vars</td>
<td>List of names of right hand side variables</td>
</tr>
<tr>
<td>@IFCONV</td>
<td>scalar</td>
<td>1</td>
<td>=1 if convergence achieved, 0 otherwise</td>
</tr>
<tr>
<td>@YMEAN</td>
<td>scalar</td>
<td>1</td>
<td>Mean of the dependent variable</td>
</tr>
<tr>
<td>@SDEV</td>
<td>scalar</td>
<td>1</td>
<td>Standard deviation of the dependent variable</td>
</tr>
<tr>
<td>@NOB</td>
<td>scalar</td>
<td>1</td>
<td>Number of observations</td>
</tr>
<tr>
<td>@HIST</td>
<td>vector</td>
<td>#values</td>
<td>Frequency counts for each dependent variable value.</td>
</tr>
<tr>
<td>@HISTVAL</td>
<td>vector</td>
<td>#values</td>
<td>Corresponding dependent variable values</td>
</tr>
<tr>
<td>@SSR</td>
<td>scalar</td>
<td>1</td>
<td>Sum of squared residuals</td>
</tr>
<tr>
<td>@RSQ</td>
<td>scalar</td>
<td>1</td>
<td>correlation type R-squared</td>
</tr>
<tr>
<td>@OVERDIS</td>
<td>scalar</td>
<td>1</td>
<td>Overdispersion test</td>
</tr>
<tr>
<td>%OVERDIS</td>
<td>scalar</td>
<td>1</td>
<td>p-value for overdispersion test</td>
</tr>
<tr>
<td>@LR</td>
<td>scalar</td>
<td>1</td>
<td>Likelihood ratio test for zero slope coefficients</td>
</tr>
<tr>
<td>%LR</td>
<td>scalar</td>
<td>1</td>
<td>P-value for likelihood ratio test</td>
</tr>
<tr>
<td>@LOGL</td>
<td>scalar</td>
<td>1</td>
<td>Log of likelihood function</td>
</tr>
<tr>
<td>@SBIC</td>
<td>scalar</td>
<td>1</td>
<td>Schwarz Bayesian Information Criterion</td>
</tr>
</tbody>
</table>
POISSON

@NCOEF scalar 1 Number of independent variables (#vars)
@NCID scalar 1 Number of identified coefficients
@COEF vector #vars Coefficient estimates
@SES vector #vars Standard errors
@T vector #vars T-statistics
%T vector #vars p-values for T-statistics
@GRAD vector #vars Gradient of log likelihood at convergence
@VCOV matrix #vars* Variance-covariance of estimated coefficients
@FIT series #obs Fitted values of dependent variable
@RES series #obs Residuals = actual-fitted values of dependent variable

If the regression includes a PDL variable, the following will also be stored:

@SLAG scalar 1 Sum of the lag coefficients
@MLAG scalar 1 Mean lag coefficient (number of time periods)
@LAGF vector #lags Estimated lag coefficients, after "unscrambling"

Method

POISSON uses analytic first and second derivatives to obtain maximum likelihood estimates via the Newton-Raphson algorithm. This algorithm usually converges fairly quickly. Zeros are used for starting parameter values, except for the constant term. @START can be used to provide different starting values (see NONLINEAR in this help system). As in other regression procedures in TSP, estimation is done using a generalized inverse in the case of multicollinearity of the independent variables.

The overdispersion test is a Lagrange multiplier test based on regressing the difference between the estimated variance and the dependent variable on the fitted value. The statistic is T (the number of observations) times the R-squared from the following regression:

\[(e_i^2 - y_i) = \alpha + \beta \hat{y}_i + u_i\]


The exponential mean function is used in the Poisson model. That is, if \( X \) are the independent variables and \( B \) are their coefficients,

\[E(Y|X) = \exp(X'B)\]
This guarantees that predicted values of $Y$ are never negative. Because the Poisson model implies that the variance of the dependent variable is equal to the mean, the effect of Poisson estimation is to downweight the large $Y$ observations relative to ordinary regression. If you use LSQ to run an unweighted nonlinear regression with the same exponential mean function, you will get a better fit to large $Y$ values than with the Poisson model.

The ML command can also be used to estimate Poisson models, including panel data models with fixed and random effects. See our web page for the panel examples.

**Options**

The usual nonlinear estimation options can be used. See the [NONLINEAR](#) entry.

**Examples**

Poisson regression of patents on lags of log(R&D), a scientific sector dummy, and firm size:

```
POISSON PATENTS C LRND LRND(-1) LRND(-2) DSCI SIZE;
```

**References**


PRIN obtains the principal components of a group of series. The number of such components obtained may be a fixed number, or it may be determined by the amount of variance in the original series explained by the principal components.

Principal components are a set of orthogonal vectors with the same number of observations as the original set of series which explain as much variance as possible of the original series. Users of this procedure should be familiar with the method and uses of principal components, which are described in many standard texts such as Harman (1976) or Theil (1971).

PRIN (FRAC=<fraction of variance>, NAME=<name of components>, NCOM=<number of components>, PRINT) <list of series> ;

Usage

To obtain principal components in TSP give the word PRIN followed by a list of series whose principal components you want. The options determine how many principal components will be found. The resulting principal components are also series and are stored in data storage under the names created from the NAME= option.

Output

If PRINT is on, the output of the principal components procedure begins with a title, the list of input series, the number of observations, and the correlation matrix of the input series. This is followed by a table for the components, showing the corresponding characteristic root, and the fraction of the variance of the original series which was explained by all the components up to and including this one.

Finally a table of factor loadings is printed: this table shows the weights applied to each component in expressing each input series as a function of the components.

PRIN stores the correlation matrix of the input variables under the name @CORR and stores the components themselves under the names P1, P2, P3, etc. as time series. If you supply a different prefix for the names from P, PRIN will use that when making the names.

Method
TSP standardizes the variables (subtracts their means and divides by their standard deviations) before computing the principal components. The resulting components have the following properties:

1. They have mean zero, standard deviation unity, and are orthogonal.

2. The correlation coefficients between a principal component vector and the set of original variables are identical to that component's loading factor.

3. The sum of squared loading factors equals the characteristic root. (In some other principal component packages, the sum of squared factor loadings equals unity; this is a matter of arbitrary scaling.) In calculating the principal components, the factor loadings are divided by the characteristic root to obtain a principal component with standard deviation of unity. Other programs treat the scaling differently.

4. The fraction of the variance of the original variables explained by a principal component is its characteristic root divided by the number of variables.

The TSP commands below obtain the same results as the PRIN X Y Z; command:

```
CORR X Y Z;
MAT EVEC = EIGVEC(@CORR);
MAT FACTLD = EVEC*DIAG(SQRT(@EIGVAL));
MFORM (TYPE=TRI,NROW=3) ONE=1;
MAT FRACVAR = ONE'@EIGVAL/3;
PRINT @EIGVAL FRACVAR FACTLD;
```

```
Assumes X Y Z are already standardized
MAT PCOM = XM*(FACTLD*(DIAG(@EIGVAL))
```

Options

FRAC= the fraction of the variance of the input variables which you wish to explain with the principal components.

NAME= the prefix to be given to the names of the principal components: the components will be called prefix1, prefix2, and so forth. You may use any legal TSP name as the name for the principal components, but the names generated by adding the numbers must also be legal TSP names (i.e., of the appropriate length).
PRIN

NCOM= the maximum number of components to be determined. The actual number will be the minimum of the number requested, the number of variables, and the number needed to explain FRAC of the variance.

PRINT/NOPRINT tells whether the results of PRIN are to be printed or just stored in data storage.

The default values of the PRIN options are NAME=P, NCOM=number of variables, FRAC=1. This set of options is non-limiting, that is, the maximum number of components possible will always be constructed.

Example

PRIN (NAME=PC,NCOM=3,FRAC=.95) I TIME CONS GOVEXP EXPORTS

specifies that three principal components are to be found for five variables I, TIME, GOVEXP, and EXPORTS. If 95% of the variance of the five variables can be explained by fewer than three components, the program will stop there. The principal components found will be stored under the names PC1, PC2, and PC3, for further use in the program.

References

Harman, Harry H., Modern Factor Analysis, University of Chicago Press, First Edition (1960), Sec. 9.3 or Third Edition (1976), Sec. 8.3.


PRINT

PRINT is a synonym for WRITE.

PRINT <list of variables> ;
PROBIT obtains estimates of the linear probit model, where the dependent variable takes on only two values. Options allow you to obtain and save the inverse Mills ratio as a series so that the sample selection correction due to Heckman can be estimated (also see the SAMPSEL command).

PROBIT (FEI, FPRINT, MILLS=<name for output inverse Mills ratio>, NHERMITE=<number of points for hermite quadrature>, REI, nonlinear options) <dependent variable> <list of independent variables> ;

Usage

The basic PROBIT statement is like the OLSQ statement: first list the dependent variable and then the independent variables. If you wish to have an intercept term in the regression (usually recommended), include the special variable C or CONSTANT in your list of independent variables. You may have as many independent variables as you like subject to the overall limits on the number of arguments per statement and the amount of working space, as well as the number of data observations you have available.

The observations over which the regression is computed are determined by the current sample. If any of the observations have missing values within the current sample, PROBIT will print a warning message and will drop those observations. PROBIT also checks for complete or quasi-complete sample separation by one of the right hand side variables; such models are not identified.

The list of independent variables on the PROBIT command may include variables with explicit lags and leads as well as PDL (Polynomial Distributed Lag) variables. These distributed lag variables are a way to reduce the number of free coefficients when entering a large number of lagged variables in a regression by imposing smoothness on the coefficients. See the PDL section for a description of how to specify such variables.

The dependent variable need not be a strictly zero/one variable. Positive values are treated as one and zero or negative values are treated as zero.
The FEI and REI options compute estimates for models with fixed and random effects for individuals respectively. FREQ (PANEL) must be in effect. For fixed effects, a very efficient algorithm is used, so large unbalanced panels can easily be handled. The FEPRINT option prints a table of the effects, their standard errors, and t-statistics. Individuals that have dependent variable values that are all zero or all one are allowed, although their data are not informative for the slopes. The fixed effects for such individuals will be either a very large negative number (in the case of zero) or a very large positive number (in the case of one). These values yield the correct probability for these observations (zero or one). Note that this estimator has a finite-T bias, so the number of time periods per individual should not be too small. The random effects model is estimated by maximum likelihood; see the method section below for details.

Output

The output of PROBIT begins with an equation title and the name of the dependent variable. Starting values and diagnostic output from the iterations will be printed. Final convergence status is printed.

This is followed by the mean of the dependent variable, number of positive observations, sum of squared residuals, R-squared, and a table of right hand side variable names, estimated coefficients, standard errors and associated t-statistics.

PROBIT also stores some of these results in data storage for later use. The table below lists the results available after a PROBIT command.

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@LOGL</td>
<td>scalar</td>
<td>1</td>
<td>Log of likelihood function</td>
</tr>
<tr>
<td>@IFCONV</td>
<td>scalar</td>
<td>1</td>
<td>Convergence status (1 = success)</td>
</tr>
<tr>
<td>@NOB</td>
<td>scalar</td>
<td>1</td>
<td>Number of observations</td>
</tr>
<tr>
<td>@NPOS</td>
<td>scalar</td>
<td>1</td>
<td>Number of positive observations</td>
</tr>
<tr>
<td>@SRSQ</td>
<td>scalar</td>
<td>1</td>
<td>Scaled R-squared for binary probit</td>
</tr>
<tr>
<td>@RSQ</td>
<td>scalar</td>
<td>1</td>
<td>Squared correlation between Y and @FIT</td>
</tr>
<tr>
<td>@SSR</td>
<td>scalar</td>
<td>1</td>
<td>Sum of squared residuals</td>
</tr>
<tr>
<td>@RNMS</td>
<td>list</td>
<td>#params</td>
<td>List of parameter names</td>
</tr>
<tr>
<td>@GRAD</td>
<td>vector</td>
<td>#params</td>
<td>Gradient of likelihood function at maximum</td>
</tr>
<tr>
<td>@COEF</td>
<td>vector</td>
<td>#params</td>
<td>Estimated values of parameters</td>
</tr>
<tr>
<td>@SES</td>
<td>vector</td>
<td>#params</td>
<td>Standard errors of estimated parameters</td>
</tr>
<tr>
<td>@T</td>
<td>vector</td>
<td>#params</td>
<td>T-statistics</td>
</tr>
<tr>
<td>%T</td>
<td>vector</td>
<td>#params</td>
<td>p-values for T-statistics</td>
</tr>
</tbody>
</table>
PROBIT

@VCOV vector #par*#par Estimated variance-covariance of estimated parameters
@DPDX matrix #vars*2 Matrix of mean probability derivatives for the two values of the dependent variable
@MILLS series #obs Inverse Mills ratios
@FIT series #obs Fitted probabilities
@NCOEFAI scalar 1 Number of fixed effects
@NCIDAI scalar 1 Number of identified fixed effects
@AI series #obs estimated fixed effects stored as a series (for FEI)
@COEFAI vector #individuals estimated fixed effects (for FEI)
@SESAI vector #individuals standard errors for fixed effects (for FEI)
@TAI vector #individuals T-statistics for fixed effects (for FEI)
%TAI vector #individuals p-values corresponding to T-statistics for fixed effects (for FEI)

If the regression includes a PDL variable, the following will also be stored:

@SLAG scalar 1 Sum of the lag coefficients
@MLAG scalar 1 Mean lag coefficient (number of time periods)
@LAGF vector #lags Estimated lag coefficients, after "unscrembling"

Method

PROBIT uses analytic first and second derivatives to obtain maximum likelihood estimates via the Newton-Raphson algorithm. This algorithm usually converges fairly quickly. TSP uses zeros for starting parameter values, unless @START is used to override this (see the NONLINEAR entry). As in other regression procedures in TSP, estimation is done using a generalized inverse in the case of multicollinearity of the independent variables.

The numerical implementation involves evaluating the normal density and cumulative normal distribution functions. The cumulative normal distribution function is computed from an asymptotic expansion, since it has no closed form. See the reference under the CDF command for the actual method used to evaluate CNORM(). The ratio of the density to the distribution function is also known as the inverse Mills ratio. This is used in the derivatives and with the MILLS= option.

@MILLS is actually the expectation of the structural residual, where the model is given by

350
$y_i = x_i \beta + \varepsilon_i \quad \varepsilon_i \sim N(0,1)$
$D_i = I(y_i > 0)$

@MILLS is the value of the following two expressions, depending on whether $D=0$ or $1$:

$$E(\varepsilon | X, D = 1) = \frac{\text{NORM}(-Xb)}{1 - \text{CNORM}(-Xb)}$$
$$= \frac{\text{NORM}(Xb)}{\text{CNORM}(Xb)} = \text{DLCNORM}(Xb)$$

$$E(\varepsilon | X, D = 0) = -\frac{\text{NORM}(-Xb)}{\text{CNORM}(-Xb)}$$
$$= -\text{DLCNORM}(-Xb)$$

where NORM is the normal density, CNORM is the cumulative normal and DLCNORM is the derivative of the log cumulative normal with respect to its argument. Before estimation, PROBIT checks for univariate complete and quasi-complete separation of the data and flags this condition. The model is not identified in this case, because one or more of the independent variables perfectly predict the dependent variable for some of the observations, and therefore their coefficients would slowly iterate to plus or minus infinity if estimation was allowed to proceed.

The scaled R-squared is a measure of goodness of fit relative to a model with just a constant term; it replaced the Kullback-Leibler R-squared beginning with TSP 4.5 since it has somewhat better properties for discrete dependent variable problems. See the Estrella (1998) article.

The Probit random effects model estimated is the following:

$$y_{it} = I(X_{it} \beta + \alpha_i + \varepsilon_{it} > 0)$$
$$\alpha_i \sim N(0, \sigma^2_{\alpha}) \quad \text{and} \quad \varepsilon_{it} \sim N(0, \sigma^2)$$
$$\sigma^2_\alpha + \sigma^2 = 1$$
$$\rho = \frac{\sigma^2}{\sigma^2_\alpha + \sigma^2}$$
This normalization means that the slope estimates are normalized the same way as the results from the usual Probit command. The parameter \textit{RHO} is estimated and corresponds to the share of the variance that is within individual. The likelihood function involves computing a multivariate integral and this is done with Hermite quadrature, using a default 20 points; when \textit{RHO} is high, it may be necessary to increase this using the NHERMITE option.

\section*{Options}

\texttt{FEI/NOFEI} specifies that the fixed effects Probit model should be computed. \texttt{FREQ} (PANEL) must be in effect.

\texttt{FEPRINT/NOFEPRIN} specifies whether the estimated effects and their standard errors should be printed.

\texttt{MILLS=} the name of a series used to store the inverse Mills ratio series evaluated at the estimated parameters. The default is \texttt{@MILLS}.

\texttt{NHERMITE=} number of points for the Hermite quadrature in computing the integral for the random effects Probit model. The default is 20. The value set is retained throughout the TSP run.

\texttt{REI/NOREI} specifies that the random effects Probit model should be computed. \texttt{FREQ} (PANEL) must be in effect.

The usual nonlinear estimation options can be used. See the \texttt{NONLINEAR} entry.

\section*{Examples}

Standard probit model:

\begin{verbatim}
PROBIT MOVE C WAGE1 WAGE2 COST1 COST2;
\end{verbatim}

Heckman sample selection model (see the \texttt{SAMPSEL} command for ML estimation of this model):

\begin{verbatim}
PROBIT (MILLS=RMILL) WORK C OCC1 OCC2 TENURE MSTAT AGE;
SELECT WORK;
OLSQ LWAGE C SCHOOL EXPER IQ UNION OCC1 OCC2 RMILL;
\end{verbatim}

Computing fitted probabilities and inverse Mills ratios explicitly:

\begin{verbatim}
PROBIT MOVE C WAGE1 WAGE2 COST1 COST2; FORCST XB;
MOVEP = CNORM(XB);
MILLSR = MOVE * DLCNORM(XB) + (1-MOVE) * (-DLCNORM(-XB));
\end{verbatim}
References


PROC

Examples

PROC defines a TSP user procedure. It is always the first statement in a procedure and must be matched by a corresponding ENDPROC statement.

PROC <procedure name> [<list of arguments>] ;

Usage

PROC gives the procedure name (any legal TSP name) and the list of "dummy" arguments for the procedure. When the procedure is called by specifying its name somewhere else in the TSP run, any dummy arguments which are used in the procedure are replaced by the actual arguments on the statement which invokes the procedure.

Always include an ENDPROC statement at the end of your procedure.

A user procedure can use any of the TSP variables which are at a higher level, that is, any variables in the procedure(s) which call it, or in the main TSP program. Variables which are created in a lower level procedure are not available after leaving that procedure unless they are the names of dummy arguments. In programming terminology, they are local variables.

When you enter a procedure, the last SMPL processed is in force. When you leave be careful to restore the SMPL if you have changed it during the procedure, or you may get unpredictable results if you call the PROC from different parts of the program.

If the name of a LIST is used as an argument to a PROC, the list is not expanded until it is used inside the PROC. This allows the number of items in the list to vary between PROC uses. The number of items in the list for a particular call can be determined by the LENGTH command.

Output

PROC produces no output, unless your procedure generates output.

Examples

This simple example creates a dummy variable over a sample specified by START and STOP, which is one every SKIPth observation and zero elsewhere. This is an inefficient way to create a seasonal dummy, or year dummies for panel data (it is better to use a repeating TREND and the DUMMY command). Note how the SMPL and FREQ are saved and restored on exit from the PROC.
The next example is a procedure to compute the prediction error for the classical linear regression model. The formula for the error variance is

\[ s^2 \left[ 1 + X_0 (X'X)^{-1} X_0 \right] \]

where \( s \)-squared is the estimate of the residual variance, \( X'X \) is the moment matrix for the data in the regression, and \( X(0) \) is the matrix of data for the prediction interval. This PROC is invoked by the statements

\[ \text{LIST VARLIST VAR1 VAR2 VAR3 ; ? Example with 3 variables} \]

\[ \text{VARFORC ERROR VARLIST ;} \]

The sample over which you wish the forecast and error must be specified before invoking VARFORC. VARFORC expects that the estimation of the model for which you are doing the forecast has just been executed and that \@S2 and \@VCOV contain the estimated variances of the residuals and coefficients. The series ERROR contains the prediction error for each observation on return. This example also shows how to pass a list of variable names as an argument.

\[ \text{PROC VARFORC PREDERR XLIST ;} \]

\[ \text{MMAKE X XLIST ;} \]

\[ \text{MAT PREDERR = SER(SQRT(@S2 + VECH(DIAG(X*@VCOV*X'))));} \]

\[ \text{ENDPROC ;} \]

Here is a user procedure to compute Theil's inequality coefficient (U) as a normalized measure of forecast error:

\[ \text{PROC THEILU ACT PRED U ;} \]

\[ \text{GENR RESID = ACT-PRED ;} \]

\[ \text{MAT U = SQRT(RESID^2/RESID^2)/(ACT*ACT)} ; \]

\[ \text{ENDPROC ;} \]
QUIT (Interactive)

QUIT exits from interactive TSP without saving the backup file. Otherwise, it is equivalent to the END, EXIT, or STOP command.

> QUIT
RANDOM

Options   Examples   References

RANDOM creates pseudo-random variables. It can create random variables which follow the normal, uniform, Poisson, Negative Binomial, Laplace, t, Cauchy, exponential, gamma, or an empirical distribution. The user may specify (optionally) parameters of the distribution.

**RANDOM (CAUCHY, DF=<scalar>, DRAW=<series> or <matrix>,
EDF=<series>, EXPON, GAMMA, GEN=1 or 2,
MEDIAN=<scalar>, LAPLACE, MEAN=<scalar> or <series>,
NEGBIN, POISSON, REPEAT, SEEDIN=<scalar>,
SEEDOUT=<scalar>, STDEV=<scalar> or <series>, T, UNIFORM,
VAR=<scalar> or <series>, VCOV=<covariance matrix>,
VMEAN=<mean vector>) <series> or <matrix> or <list of series> ;

Usage

RANDOM with no options causes a normal random variable with mean zero and variance one to be generated and stored as a series (under control of the current SMPL). If you want a non-standardized random variable, include the MEAN and STDEV options.

Other options cause the random variable generated to follow the Poisson, negative binomial, uniform, t, Cauchy, exponential, gamma, Laplace (double exponential), or general empirical distributions. See the Examples Section to learn how to obtain random variables from other distributions.

TSP "randomizes" the seed to start every run based on the current time, so you need to specify a fixed seed if you want to reproduce results from run to run. To change the starting seed for any given run or to save it for future use, use the SEEDIN and SEEDOUT options.

If multivariate normal random deviates are desired, the VCOV option is required. The dimension of this (symmetric or diagonal) matrix is the number of series to create. If only one argument is supplied before the semicolon, the series are stored in a matrix with this name. The VMEAN option should be used if a non-zero mean vector is desired.
To create a random variable from an empirical distribution function, a series (usually a set of residuals) generated by the distribution function must be supplied. This feature is useful for computing bootstrap standard errors. A set of residuals generated by the model is used as input to the DRAW option and a new series with the same distribution as the old one is obtained by drawing observations from a discrete distribution with probability mass equal to one divided by the number of observations placed on each observed value of the residuals. This new sample of residuals may then be used in further computations to obtain estimates of functions of these random variables. To draw without replacement, use the NOREPLACE option.

Method

The method used by RANDOM is the multiplicative congruential method. The new uniform generator (GEN=2) has period $2^{31}9$, and is a combination of 2 multiple recursive generators with 8 seeds. See L'Ecuyer (1999), generator MRG32k5a. The old uniform generator (GEN=1) has period $2^{31}-1$ and multiplier 41358; this choice is described in L'Ecuyer(1990) (it has optimal "randomness" in its class). Both are implemented in integer math for speed, and to insure a full period (no repeats in $2^{31}$ or $2^{31}$ draws).

The multiplicative congruential method produces random numbers which are uniform on the (0,1) interval. Normal and Poisson random variables are created from uniform random variables with ACM Algorithms #488 and #369, respectively. Gamma random variables use ACM Algorithm #599. Negative binomial random variables are computed by drawing Gamma random variables to determine $Y$, and then drawing Poisson random variables with mean $Y$. All other random variables are derived from the uniform random variables using the inverse distribution function, which usually involves an asymptotic expansion (see the CDF references).

Options

CAUCHY/NOCAUCHY specifies that the random number generated is to follow the Cauchy distribution:

$$f(x) = \frac{1}{\pi(1+x^2)}$$

DF = the degrees of freedom for student's t distribution (see the t option).
**RANDOM**

**DRAW** = the name of a series or matrix which will be sampled with probability one divided by the number of observations to generate the random numbers. This series does not have to be sorted in any order. If DRAW= a matrix, a multivariate set of random numbers is drawn. The number of random variables is equal to the number of columns in the matrix. Note that the matrix from which you are drawing does not have to have the number of rows equal to the number of observations. This is very useful for simulation or bootstrap standard errors.

**EDF** = Empirical Distribution Function. Same thing as DRAW=.

**EXPON/NOEXPON** specifies that the random number generated is to follow the exponential distribution:

\[ f(x) = \lambda \exp(-\lambda x) \]
\[ E(x) = \lambda^{-1} \quad V(x) = \lambda^{-2} \]

Use the LAMBDA= option to specify the parameter lambda.

**GAMMA/NOGAMMA** specifies that the random number generated is to follow the gamma distribution:

\[ f(x) = \frac{\gamma \alpha}{\Gamma(\alpha)} x^{\alpha-1} \exp(-\gamma x) \]
\[ E(x) = \alpha / \gamma \quad V(x) = \alpha / \gamma^2 \]

Use the MEAN= and STDEV= options to specify the parameters, which must be non-negative.

**GENERATOR** = 1 or 2. Type of uniform random number generator. Use GEN=1 to reproduce results from older versions of TSP (to 4.4); this generator has period equal to 2**31-1. The new (default) generator has period 2**319.

**LAMBDA** = the exponential or double exponential parameter. (See the EXPON and LAPLACE options).

**LAPLACE/NOLAPLACE** specifies that the random number generated is to follow the Laplace (double exponential) distribution:

\[ f(x) = \lambda \exp(-2|\lambda x|) \]
\[ E(x) = 0 \quad V(x) = 1/(2\lambda^2) \]
**RANDOM**

**MEAN** = the expected value of the random variable or a series containing expected values. Applies only to the normal, gamma, Poisson, and negative binomial random variables. The default value is zero (you will want to change this for the gamma, Poisson, and negative binomial distributions). When a series is supplied, each random number drawn will come from a distribution with a different mean.

**NEGBIN/NONEGBIN** specifies that the random number generated is to follow the negative binomial\((N, p)\) distribution, which is excess waiting time to obtain \(N\) successes (the number of trials minus \(N\)) with success probability \(p\) for each trial. \(N\) does not have to be an integer. For this model, the mean and variance of the data are given by

\[
E(y) = \frac{N(1-p)}{p}, \quad V(y) = \frac{N(1-p)}{p^2} = \frac{E(y)}{p}
\]

An alternative widely used specification is the following:

\[
E(y) = \frac{\alpha}{\delta}, \quad V(y) = \frac{\alpha(1+\delta)}{\delta^2} = E(y)\left(1+\frac{\delta}{\delta}\right)
\]

The two specifications are equivalent and imply the following identities:

\[
N = \alpha, \quad p = \frac{\delta}{1+\delta}, \quad \text{or} \quad \delta = \frac{p}{1-p}, \quad \delta \geq 0
\]

Use the **MEAN=** and **STDEV=** options to specify the parameters. The MEAN must be non-negative, and STDEV must be larger than or equal to the square root of the mean.

**POISSON/NOPoisson** specifies that the random number generated is to follow the Poisson distribution:

This distribution has one parameter, alpha, which is both the expected value and the variance. Supply this parameter using the **MEAN=** option. It must be a non-negative number.

**REPLACE/NOREPLACE** specifies that DRAWing from the empirical distribution function is to be done with replacement (the default) or no replacement.
SEEDIN = value of random seed to start random generator (replaces the current value of the random seed). This must be an integer in the range \([1,2.1\text{ billion}]\) (otherwise it is moved into this range). Note that scalar values are stored in double precision, which allows for 16 significant digits, so don't make the seed too large if you are trying to reproduce results. When used with the new default uniform generator, all 8 seeds are set to the SEEDIN value.

SEEDOUT = random seed of the random generator (the current random seed, before any random variables are created by this command). To print the seed, use OPTIONS NWIDTH=20; to provide enough digits. SEEDOUT is not useful with the new uniform generator, because it uses 8 seeds.

STDEV = the standard deviation of the random variable or a series containing standard deviations. This option applies only to normal, gamma, and negative binomial random variables. The default value is one. When a series is supplied, each random number drawn will come from a distribution with a different standard deviation.

T/NOT specifies that the random number generated is to follow student's t distribution with degrees of freedom given by the DF= option.

UNIFORM/NOUNIFORM specifies that the random number generated is to follow the uniform distribution between zero and one.

VAR= the variance of the random variable or a series containing variances. This option applies only to normal, gamma, and negative binomial random variables. The default value is one. When a series is supplied, each random number drawn will come from a distribution with a different variance.

VCOV = (symmetric) variance-covariance matrix for multivariate normal random variables. You can also supply the (triangular) square root of this matrix, which saves a step.

VMEAN = mean vector for multivariate normal random variables (the default is a vector of zeroes) or for creating several series at the same time, each with a different mean.

Examples

These examples each generate a thousand random numbers and store them under the series name given.

SMPL 1 1000 ;
RANDOM STDNORM ; ? Std normal random variable

RANDOM (UNIFORM) FLAT ; ? Uniform (0,1) r. v.
FLATAB = FLAT*(B-A) + A ;  ? Uniform on the interval (A,B)
CDF (CHISQ,DF=3,INV) FLAT CHI3;  ? Chi-square(3) from uniform
T1EV = -LOG(-LOG(FLAT));  ? Type I Extreme Value from uniform

RANDOM (CAUCHY) FAT:
RANDOM (EXPON, LAMBDA=2) Z;
RANDOM (LAPLACE, LAMBDA=0.5) DE;
RANDOM (T, DF=5) FATAIL;
RANDOM (GAMMA, MEAN=10, STDEV=2) GAMMA10;
RANDOM (NEGBIN, MEAN=10, STDEV=5) NEGBIN10;
RANDOM (MEAN=10, STDEV=3.1623) NORM10;
RANDOM (MEAN=10, POISSON) POISS10;

The last two examples produce random numbers with the same mean and variance, but different distributions. A normal variable with mean 10 and standard deviation 3.1623 could also have been generated by the following:

NORM10 = 10 + STDNORM*3.1623;

The next example generates a bivariate normal random vector with correlation 0.5:

LOAD (TYPE=SYM, NROW=2) COVMAT; 1 .5 1;
RANDOM (VCOV=COVMAT) NU1 NU2;

Now we use the estimated residuals from a regression to generate 10 samples with the same empirical distribution function:

LIST RES RES1-RES10;
RANDOM (EDF=@RES) RES;

When we are done, the list RES consists of ten series which have the same distribution as the original residual series @RES.

Example with inverse distribution functions:

RANDOM (SEED=94298, UNIFORM) U;
EV = -LOG(-LOG(U));  ? Type I Extreme Value
CDF(INV, CHISQ, DF=n) U CHIV;  ? Chi-square (n)

An example using the matrix version of DRAW to draw data from an empirical distribution function in order to investigate the potential rate of convergence of a particular estimator:

? original data set has 100 observations, compute residuals
SMPL 1 100;
OLSQ Y C X;
UNMAKE @COEF A B;
MMAKE EDF @RES X;
? estimation using 1000 obs drawn from EDF
SMPL 1 1000;
RANDOM (DRAW=EDF) E X;
Y = A+B*X+E;
OLSQ Y C X;
? estimation using 10,000 obs drawn from same EDF
SMPL 1 10000;
RANDOM (DRAW=EDF) E X;
Y = A+B*X+E;
OLSQ Y C X;

Draw 5 cards from a deck of 52 without replacement:

SMPL 1 52;
TREND OBS; SUITE = 1+INT((OBS-1)/13);
TREND (PER=13) NUMBER;
MMAKE CARDS SUITE NUMBER;
SMPL 1 5;
RANDOM (DRAW=CARDS,NOREPL) SUITE NUMBER;

Permute a series of residuals:

SMPL 1 100;
OLSQ Y C X1 X2;
RANDOM (DRAW=@RES,NOREPL) U;

Verify that the new uniform generator is properly implemented (check sum of first 10,000,000 r.v.s for seed 12345):

OPTIONS DOUBLE MEMORY=5;
SMPL 1,100000;
RANDOM(GEN=2,SEEDIN=12345);
SET TOTAL=0;
DO I=1,100;
RANDOM(UNIFORM) X;
MAT TOTAL = TOTAL + SUM(X);
ENDDO;
SET CORRECT = 5000494.15;
PRINT TOTAL, CORRECT;

References


READ

Options   Examples

READ is used to read series, matrices, or scalars into data storage. Normally, data will be read from an external file, but small quantities of data can be included in a "data section" at the bottom of your program file (when running TSP in batch mode). READ can be also be used to read spreadsheets or stata™ data files. If you plan to repeatedly use a very large dataset, the fastest way to access it is as a TSP databank - see the OUT and IN commands for further details.

```
READ (BYOBS, BYVAR, FILE='filename string' or <filename>,
       FORMAT=BINARY or DATABANK or EXCEL or FREE or LOTUS
       or RB8 or STATA or 'format string', FULL, NCOL=<number of
       columns>, NROW=<number of rows>, PRINT, SETSMPL,
       SHEET='sheetname string', TYPE=CONSTANT or DIAG or
       GENERAL or SYMMETRI or TRIANG, UNIT=<I/O unit number>)
       <list of series or matrices or constants> ;
```

or

```
READ ;
```

Usage

If TSP encounters a simple READ statement with no arguments, it transfers to the data section and begins reading data until it reaches an END statement (or end of file), at which point it returns to the line in the TSP program following the READ statement. A NOPRINT command in the data section will stop the data values from echoing to the output.

A READ statement followed by a list of series names is the easiest way to read small quantities of data. If no options are specified, the data is assumed to follow the READ statement directly in free format, each number separated from the others by one or more blanks. Each group of data may be terminated by a semicolon (;), although this is not required. If there is more than one series to be read in, the order of the data is the first observation of each of the series, followed by the second observation of each of the series, and so forth.

READ can also be used to read several variables from an external file in free format, such as:

```
READ (FILE='FOO.DAT') X Y Z ;
```
There are two special features for free-format numbers. The first is the use of the dot (.) to specify a missing value (and is similar to the SAS convention). However, note that in other places in TSP (such as in formulas), dot (.) is treated as a DOT variable. Use @MISS or @NA to represent a missing value in a formula (see FRML). The other special feature is a repeated number, specified by an integer repeat count, a star (*), and the value to be repeated. For example, 3*0 is equivalent to 0 0 0. This is most often used for repeated zeroes in special matrices (like band matrices) and resembles the repeat count feature in the FORTRAN free-format (*) READ.

When the data is read in free format, the number of items read must be equal to the number of series times the number of observations in the current SMPL. TSP checks for this and prints a message when the check fails. TSP will determine the length of the SMPL itself if the SETSMPL option is on.

To read matrices use a READ statement with options to define the matrix, and the matrix's name for storage. Follow the statement with the numbers which compose the matrix in free format, a row at a time. That is, a 3 (# rows) by 2 (# of columns) matrix is read in the following order: (1,1), (1,2), (2,1), (2,2), (3,1), (3,2).

A matrix of any type may be read by specifying all its elements, but there are special forms for reading symmetric, triangular, and diagonal matrices. If a matrix is symmetric, only the lower triangle needs to be read, i.e., elements (1,1), (2,1), (2,2), (3,1), (3,2), (3,3), and so forth. The FULL option specifies whether the full matrix is being specified or only the lower triangle. If the matrix is triangular, you need only specify the transpose of the upper triangular portion: (1,1), (1,2), (2,2), (1,3), (2,3), (3,3), and so forth. If the matrix is diagonal, only the diagonal needs to be given: (1,1), (2,2), (3,3), .... It will be filled out with zeroes when used.

Stata™ files

TSP reads stata version 2-10 .dta files. The variables have the same names as they have in stata, although string (text) variables are generally not supported.

Spreadsheet files

TSP can read series and matrices directly to or from spreadsheet files. The following files are supported by TSP:

<table>
<thead>
<tr>
<th>spreadsheet version</th>
<th>filename extensions</th>
<th>TSP support*</th>
</tr>
</thead>
<tbody>
<tr>
<td>Lotus 123</td>
<td>.wks</td>
<td>Read and</td>
</tr>
<tr>
<td>Lotus 123 3, Symphony 1,2</td>
<td>.wk3 .wk1 .wrk .wr1</td>
<td>Read</td>
</tr>
<tr>
<td>---------------------------</td>
<td>----------------------</td>
<td>------</td>
</tr>
<tr>
<td>Lotus 123/J (Japanese) 1,2</td>
<td>.wj1 .wj2 .wk2 .wt2</td>
<td>Read</td>
</tr>
<tr>
<td>Microsoft Excel 2</td>
<td>.xls</td>
<td>Read and write</td>
</tr>
<tr>
<td>Microsoft Excel 3, 4</td>
<td>.xls</td>
<td>Read</td>
</tr>
<tr>
<td>Microsoft Excel 5, 7, 8, 97, 98, 2000, 2002, 2007</td>
<td>.xls .xlw</td>
<td>Read mult. sheets</td>
</tr>
<tr>
<td>Microsoft Excel 2007</td>
<td>.xlsx .xslb</td>
<td>Neither (yet)</td>
</tr>
<tr>
<td>Quattro Pro</td>
<td>.wq1</td>
<td>Read</td>
</tr>
</tbody>
</table>

* Note that TSP generally writes the oldest file formats, which are always readable by more recent spreadsheet releases.

Spreadsheet files should be in the format of the following example, SML.XLS, which consists of quarterly data on two series, CJMTL and PMTL, for 1948:1 to 1949:1):

<table>
<thead>
<tr>
<th>A</th>
<th>B</th>
<th>C</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>Date</td>
<td>CJMTL</td>
</tr>
<tr>
<td>2</td>
<td>Mar-48</td>
<td>183.4</td>
</tr>
<tr>
<td>3</td>
<td>Jun-48</td>
<td>185.2</td>
</tr>
<tr>
<td>4</td>
<td>Sep-48</td>
<td>192.1</td>
</tr>
<tr>
<td>5</td>
<td>Dec-48</td>
<td>193.3</td>
</tr>
<tr>
<td>6</td>
<td>Mar-49</td>
<td>206.9</td>
</tr>
</tbody>
</table>

The above file could be read with the following command:

```
READ (FILE='SML.XLS') ;
```

Series are read from individual columns in the file. Series names are optionally supplied in the first row of the file (aligned) above the data columns. Dates may be given in the first column. Many different file configurations are possible, and it is possible to read in some series while ignoring others. Following are some simple guidelines for creating a spreadsheet file for TSP:
1. Put the column names in the first row. They should be valid series names in TSP (lowercase is fine - it will be converted to uppercase, but imbedded blanks and special characters are not allowed). If the file has no names, you can supply them when you read the file in TSP, but this can be inconvenient. If the file contains invalid names, the data can be read by TSP as a matrix, ignoring the current names, and you can supply your own names inside of TSP. TSP will not recognize names in lower rows or in sheets other than the first; they will be treated as missing values and numbers below them will be read as data for the original column names.

2. The second row must contain data.

3. If you are reading time series, the first column should contain dates. This will ensure the series are read with the proper frequency and starting date, regardless of the current \texttt{FREQ} and \texttt{SMPL} in TSP. If you have dates in other columns, they will be read as numbers. If you are reading a matrix, the date column will be ignored (i.e. it will not be read into the matrix).

Dates can be strings such as 48:1 or numbers formatted as dates (Mar-48, 3/31/48, etc.). You only have to supply enough dates so that TSP can detect the frequency (5 is enough to distinguish between quarterly and monthly). TSP ignores any dates after these, assuming that the data is contiguous (no missing periods/years, or SMPL gaps in TSP terminology). If you have missing periods/years for all series, leave the corresponding rows blank. Below is a table of examples showing recommended ways of defining the starting date and frequency with a dates column:

<table>
<thead>
<tr>
<th>first date</th>
<th>second date</th>
<th>resulting frequency</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>2</td>
<td>A if current FREQ is A; otherwise N</td>
</tr>
<tr>
<td>48.</td>
<td>any</td>
<td>A</td>
</tr>
<tr>
<td>48:5</td>
<td>any</td>
<td>M</td>
</tr>
<tr>
<td>48:4</td>
<td>49:1</td>
<td>Q</td>
</tr>
<tr>
<td>1948:1</td>
<td>1949:1</td>
<td>A</td>
</tr>
<tr>
<td>a2:3</td>
<td>any</td>
<td>invalid</td>
</tr>
</tbody>
</table>

**string dates - first character is ' ^ or ":**

**numeric dates:**

<table>
<thead>
<tr>
<th>first date</th>
<th>second date</th>
<th>resulting frequency</th>
</tr>
</thead>
<tbody>
<tr>
<td>12/31/48</td>
<td>12/31/49</td>
<td>A (any dates 365/366 days apart)</td>
</tr>
<tr>
<td>12/31/48</td>
<td>1/31/49</td>
<td>M (any dates 28-31 days apart)</td>
</tr>
<tr>
<td>12/31/48</td>
<td>3/31/49</td>
<td>Q (any dates 90-92 days apart)</td>
</tr>
<tr>
<td>12/31/48</td>
<td>1/1/49</td>
<td>N (any other date range)</td>
</tr>
</tbody>
</table>
4. Missing values can be represented by blank cells or by formulas which evaluate to NA, @NA, #N/A, etc.

To read in a spreadsheet file, use the FILE='filename' option. FORMAT=LOTUS or EXCEL is optional. If the filename contains one of the extensions listed earlier (.WKS, .WK3, .XLS, etc.), TSP checks the first few bytes of the file to confirm that it is one of the spreadsheet versions listed above. Conversely, if FORMAT=LOTUS or EXCEL is specified, but the filename does not contain a extension, then .WKS or .XLS is appended to the filename. To read a matrix (bypassing column names and dates), use the TYPE=GEN option. TYPE=CONSTANT is not supported for spreadsheet files; series will be defined instead. The NCOL=, NROW=, IFULL, UNIT=, etc. options are ignored, but SETSMPL is supported.

If no series names are supplied on the READ command, TSP looks for column names in the file and creates series with those names. If you supply series names, TSP attempts to match them to column names in the file. If the file does not have column names, you must supply a READ argument for each data column. If you are unsure of the file's contents, check it with your spreadsheet or read it as a matrix. If you think the file has column names, but you don't know what they are, try supplying a dummy name which won't be matched. TSP will print an error message listing the column names in the file. If you are reading a matrix (using TYPE=GEN as mentioned above), TSP will create a matrix named @LOTMAT unless you supply an argument (matrix name) to READ.

If for some reason your series are in rows instead of columns, you can read the file as a matrix, transpose it, and UNMAKE the matrix into series.

TSP checks the first row in the file for string names (cells beginning with the characters ' ^ or "). The names are truncated to 8 characters (if necessary), and are translated to uppercase. They must be aligned above their corresponding data columns. If you have dates in the first column, no name is required for the date column. Any names with imbedded blanks will be ignored.

Output

READ prints the data as it is read when the free format option is used and the PRINT options is on, otherwise READ produces no output, and stores all the data read in data storage under the series names specified. In some cases, a few special variables may be stored in data storage:

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@NOB</td>
<td>scalar</td>
<td>1</td>
<td>Number of observations read (when SETSMPL is in effect).</td>
</tr>
</tbody>
</table>
READ

@RNMS list #vars Variable names read (for spreadsheets, when no names are given).

Options

BYOBS/NOBYOBS specifies that the data is organized by observation -- the first observation for all series, then the second observation for all series, etc. This is the default.

BYVAR/NOBYVAR specifies that the data is organized by series: all observations for the first series, then all observations for the second series, etc.

FILE=’filename string’ or filename specifies the actual name of the file where your data is stored. If the filename string is 8 characters or less and does not include non-alphabetic or lowercase characters, it does not need to be enclosed in quotes.

FORMAT=Binary or DATABANK or EXCEL or FREE or LOTUS or RB4 or RB8 or STATA or ’(format text string)’ specifies the format in which your data is to be read. The default is free format, which means the fields (numbers) are separated by blanks or tabs and may be of varying length. Each of the format options is described in more detail below, and under the FORMAT entry.

FORMAT=Binary specifies that the data is in binary single precision (REAL*4) format. To read data in this format, it must be on an external file, since binary data cannot be intermixed in a TSP program input file. This method of reading data is quite fast - about the same as a TSP databank, but not quite as easy to use. This is the same as FORMAT=RB4. FORMAT=RB8 is for double precision binary.

FORMAT=DATABANK specifies that the data are to be read from a TSP databank.

FORMAT=EXCEL reads an Excel spreadsheet file (similar to FORMAT=LOTUS). If the filename ends with .XLS, this is the default. It handles version 5, 7/95, 97/98/2000/2002 Excel files. Excel 97 and later files can have up to 65536 rows of data.

FORMAT=FREE is the default. If the number of values read does not match the expected number of observations times the number of variables, an error message is printed, and TSP tries to make its best guess as to what was meant.

FORMAT=LOTUS reads Lotus 123, Excel, or Quattro Pro worksheet files (most files with the extension .WKx, where x is any character)


**FORMAT=RB4** is the same as **FORMAT=BINARY** (single precision binary).

**FORMAT=RB8** is used for double precision binary.

**FORMAT=STATA** reads Stata Version 10 or earlier files.

**FORMAT=** a format string enclosed in quotes '(format string)' specifies the format with which the data are to be read. The quotes are required and should surround a Fortran FORMAT statement, including the parentheses but excluding the word FORMAT. If you are unfamiliar with the construction of a Fortran FORMAT statement, see the FORMAT entry.

**FULL/NOFULL** applies only to reading diagonal, symmetric, or triangular matrices. It specifies whether the complete matrix is to be read, or only the upper triangle (in the case of triangular), the lower triangle (symmetric), or the diagonal (diagonal).

**NCOL=** the number of columns in the matrix. This is required for a general matrix.

**NROW=** the number of rows in the matrix. This is required for a general matrix.

Either NROW or NCOL must be specified for symmetric, triangular, or diagonal matrices. These options only apply to matrices.

**PRINT/NOPRINT** specifies whether or not the data is to be printed as it is read. This option applies only to free format reading. It may be set globally for the READ section by use of the NOPRINT statement.

**SETSMPL/NOSETSMP** specifies whether the SMPL is to be determined from the number of data items read. The default is on (SETSMPL) if no SMPL has been specified yet in the program and off otherwise. This option does not apply to matrix reading.

**SHEET= "sheet name"** reads a particular worksheet in an Excel 5 or higher file that has multiple worksheets. The default is to read the first sheet. If you request a sheet that is not in the file, TSP will list the available sheet names in the file.
TYPE=GENERAL or SYMETRIC or TRIANG or DIAG or CONSTANT specifies the type of the matrix which is to be read. GENERAL, the default, may be used for any rectangular or square matrix. SYMETRIC implies that the matrix is equal to its transpose; only the lower triangle will be stored internally to save space. TRIANG implies that the matrix is triangular (has zeroes above the diagonal). Although a lower triangular matrix is read, its transpose is stored since the TSP matrix procedures expect upper triangular matrices. DIAG means a matrix all of whose off-diagonal elements are zero. Only the diagonal is stored, and it is expanded before use. CONSTANT means a scalar or scalars are to be read and none of the matrix options will apply. If no type is specified, a warning is printed and the matrix is assumed to be general.

UNIT= an integer number (usually between 1 and 4, or 8 and 99) which is the Fortran input/output unit number of an external file from which the variables listed will be read. Usually, just FILE= is used, but UNIT= could be used to avoid typing in a long filename for several READ commands from the same file.

Examples

Reading a single series in free format:

```
SMPL 1 9 ;
READ IMPT ; 100 106 107 120 110 116 123 133 137 ;
```

This example reads formatted data from the TSP input file:

```
SMPL 1 50 ;
READ (FILE='STATES.DAT', FORMAT =
    (F2.0,F4.0,3F4.1,F6.0/F7.0,5F4.1,F4.0)' ) STATE X1-X12 ;
```

where STATE.DAT contains:

```
01284952.326.4 1.9 4120
  1905553.320.813.7 8.9 5.46677
024592 0.021.211.0 403
  303168.176.819.815.5 8.29047
.....and so forth....
```

After this data set has been read, the series STATE and X1 through X12 have the following values:

```
STATE: 1,2,.....
X1: 2849,4592,.....
X2: 52.3,0.0,.....
.....
```
X11: 8.9,15.5,.....
X12: 5.4,8.2,.....
X12: 6677,9047,.....

Reading a Stata (.dta) file, printing documentation for variables, and checking the min/max/mean of the variables:

? read all variables from file; names are stored in @RNMS
READ (FILE="acq95.dta") ;
? show documentation (Stata labels) for all variables
SHOW (DOC) @RNMS;
? check min/max/mean for all variables
MSD (TERSE) @RNMS;

Examples of reading matrices:

READ (NROW=4,NCOL=3) COEFMAT ;
  0.32 0.5 1.3
  0.30 0.4 1.35
  0.25 0.61 1.1
  0.28 0.55 1.23
;
READ (NROW=2,TYPE=SYM) COVAR ;
  4.64 2.3 5.1
READ (NROW=3,TYPE=TRIANG) TMAT ;
  1
  2 3
  4 5 6
READ (NCOL=5,TYPE=DIAG) BAND ;
  110. 0. 35. 50. ;

The matrices stored by these four examples are the following:

**COEFMAT:**

<p>| | | |</p>
<table>
<thead>
<tr>
<th></th>
<th></th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td>0.32</td>
<td>0.5</td>
<td>1.3</td>
</tr>
<tr>
<td>0.3</td>
<td>0.4</td>
<td>1.35</td>
</tr>
<tr>
<td>0.25</td>
<td>0.61</td>
<td>1.1</td>
</tr>
<tr>
<td>0.28</td>
<td>0.55</td>
<td>1.23</td>
</tr>
</tbody>
</table>

**COVAR:**

<p>| | |</p>
<table>
<thead>
<tr>
<th></th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td>4.64</td>
<td>2.3</td>
</tr>
<tr>
<td>2.3</td>
<td>5.1</td>
</tr>
</tbody>
</table>

**TMAT:**

<p>| | | |</p>
<table>
<thead>
<tr>
<th></th>
<th></th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>2</td>
<td>3</td>
</tr>
<tr>
<td>4</td>
<td>5</td>
<td>6</td>
</tr>
</tbody>
</table>
Examples for spreadsheet files

We will use this SML.WKS file (it is the Lotus version of the SML.XLS file shown earlier) in the following examples:

```plaintext
04/30/57  23.2  34.5  10.9
05/31/57  23.6  35.1  11.0
06/30/57  23.9  35.8  11.2
07/31/57  24.0   11.5
```

```plaintext
READ (FILE='SML.WKS');
?
the series CJMTL and PMTL are defined.
?
FREQ Q and SMPL 48:1, 49:1 are set if there are
?
no current FREQ or SMPL.

READ (FILE='SML.WKS',TYPE=GEN);
?
creates the 5x2 matrix @LOTMAT, with the values
?
of CJMTL and PMTL in its columns.

READ (FILE='SML.WKS') PMTL;
?
only reads in PMTL. CJMTL is ignored.
```
Define the monthly series SF, LA, and SD from 57:4 to 57:7:
If there is no current sample, this is the new sample with FREQ M.
The series LA will be given a missing value in its last observation.
READ (FILE='NM3.WK1') SF LA SD;

READ (FILE='NM3.WK1') SF;
? An error message is printed because 3 series names are required.
RECOVER

RECOVER (Interactive)

RECOVER(s) the command stream from a previously aborted TSP session.

`RECOVER [<filename>] ;`

**Usage**

With RECOVER, the command stream from a previous session may be reinstated in the event it was terminated abnormally. In most cases TSP will automatically recover an aborted session, and this procedure is not necessary. However, if you have changed directories, or renamed INDX.TMP you will have to use this procedure to recover the session.

Note that the recovery process restores the commands entered, but does not execute them -- if it was a long session, this could be costly, and you may not need all results duplicated. It is highly recommended that you REVIEW the session after it is recovered, then EXEC ranges of lines that will restore what you need to proceed.

During your interactive session, TSP is maintaining the file INDX.TMP in your current directory which contains all the commands you have entered so far in a special format (indexed, keyed access). This file is referenced any time you REVIEW, EDIT, EXEC, etc.... Upon normal exit from the program, this file is used to create a sequential file BKUP.TSP containing the commands, and INDX.TMP is deleted. If the program terminates abnormally, INDX.TMP will still exist and BKUP.TSP will not. Every time you start up interactive TSP, recovery is automatically offered if an INDX.TMP file exists in the current directory.

If you use this command to recover some other file, it MUST be a file that had been created by TSP originally as an INDX.TMP file.

RECOVER follows the same conventions as INPUT and OUTPUT for accepting filenames. Although you may specify a filename on the command line, you will probably want to be prompted for it since it is likely that you will be recovering a file from a different directory, or with a different extension.

RECOVER produces no printed output other than the information that the session has been recovered.
REGOPT

Output  Options  Examples  References

REGOPT controls the calculation and output of the regression diagnostics for OLSQ and some output of other commands. It replaces the old SUPRES and NOSUPRES commands.

\texttt{REGOPT (BPLIST=<list of variables>, CALC, CHOWDATE=<date for splitting sample>, DWPVALUE=type, LMLAGS=<# of lags for LMAR test>, PRINT, PVCALC, PVPRINT, QLAGS=<# of Q-statistics>, RESETORD=value, SHORTLAB, STAR1=<value for *>, STAR2=<value for **>, STARS) <list of output names or keywords> ;}

Usage

\texttt{OLSQ} can produce a massive number of diagnostics. REGOPT provides the user with extensive customization of this output, so that irrelevant diagnostics do not crowd relevant ones or require extensive time to calculate. The \texttt{[PV]CALC} and \texttt{[PV]PRINT} options are used along with a list of the diagnostic codes (@names) that one wishes to control. The keywords AUTO, HET, REGOUT, and ALL may also be used to control groups of diagnostics (instead of listing all the names). Other options (such as BPLIST and LMLAGS) control individual diagnostics that have no clear default. OPTIONS LIMCOL= and SIGNIF= also control the display. Note that "robust" diagnostics are available with the HI option in OLSQ.

Output

The following three examples of controlling regression output with REGOPT illustrate the range of output available. The data for these examples is a regression squared on time:

\texttt{options crt; smpl 1,10; trend t; t2 = t\^2;}

Example 1: default option

\texttt{olsq t2 c t; ? default}

\begin{verbatim}
Equation 1
-----------------
Method of estimation = Ordinary Least Squares

Dependent variable: T2
Current sample: 1 to 10
Number of observations: 10
Mean of dep. var. = 38.5000  \texttt{LM het. test = .391605 [0.531]}
Std. dev. of dep. var. = 34.1736  \texttt{Durbin-Watson = .454545 [<.012]}
Sum of squared residuals = 528.000  \texttt{Jarque-Bera test = 1.01479 [0.602]}
Variance of residuals = 66.0000  \texttt{Ramsey's RESET2 = .850706E+38 [.000]}
Std. error of regression = 8.12404  \texttt{P (zero slopes) = 151.250 [.000]}
\end{verbatim}

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**Example 2: “short label” output**

```
regopt(shortlab);
olsq t2 c t;
```

**Example 3: maximal output**

```
regopt (pvp, print, bplist=(c, t), lmlags=2, qlags=2, noshort) all;
oeq 2; t c t;
```

---

**REGOPT**

<table>
<thead>
<tr>
<th>Variable</th>
<th>Coefficient</th>
<th>Error</th>
<th>t-statistic</th>
<th>P-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>C</td>
<td>-22.0000</td>
<td>5.54977</td>
<td>3.96412</td>
<td>.004</td>
</tr>
<tr>
<td>T</td>
<td>11.0000</td>
<td>.894427</td>
<td>12.2984</td>
<td>.000</td>
</tr>
</tbody>
</table>

---

**Example 2:**

```
regopt(shortlab);
oeq 2; t2 c t;
```

**Output:**

- **Method of estimation:** Ordinary Least Squares
- **Dependent variable:** T2
- **Current sample:** 1 to 10
- **Number of observations:** 10
- **YMEAN 38.5000**
- **Mean of dep. var. = 38.5000000**
- **Std. dev. of dep. var. = 34.1735765**
- **Sum of squared residuals = 528.000000**
- **Variance of residuals = 66.0000000**
- **Std. error of regression = 8.12403840**
- **R-squared = .949764521**
- **Adjusted R-squared = .943485086**
- **ARCH test = .258229904 [.611]**
- **Cusum test = 1.26364964 ** [.003]**
- **CusumSQ test = .46590091 [.051]**
- **Chow test = 53.5714286 ** [.000]**
- **Chow het. rob. test = 53.5714286 ** [.000]**
- **LR (wet. test) (w/ Chow) = 26.4920570 ** [.000]**
- **White het. test = 3.3893051 [.184]**
- **Breusch-Pagan het. test = 1.74588036 [.186]**
Jarque-Bera test = 1.0147853 [.602]
Shapiro-Wilk test = .86938609 [.098]
Ramsey's RESET2 = .85075917E+38 ** [.000]
F (zero slope) = 151.250000 ** [.000]
Schwarz B.I.C. = 36.3245264
Akaike Information Crit. = 36.0219413

Log likelihood = -34.0219413

Estimated Standard
Variable Coefficient Error t-statistic P-value
C -22.0000000 5.54977477 -3.96412484 ** [.004]
T 11.0000000 .894427191 12.2983739 ** [.000]

Variance Covariance of estimated coefficients

Correlation matrix of estimated coefficients

CUSUM PLOT

CUSUM SQ PLOT
show scalar;  ? list of scalar results showing @names and % names

<table>
<thead>
<tr>
<th>Class</th>
<th>Name</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>SCALAR</td>
<td>@NOB</td>
<td>constant 10.00000000</td>
</tr>
<tr>
<td></td>
<td>@FREQ</td>
<td>constant 0.00000000</td>
</tr>
<tr>
<td></td>
<td>@YMEAN</td>
<td>constant 38.50000000</td>
</tr>
<tr>
<td></td>
<td>@SDEV</td>
<td>constant 34.17357654</td>
</tr>
<tr>
<td></td>
<td>@SSR</td>
<td>constant 528.00000000</td>
</tr>
<tr>
<td></td>
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<tr>
<td></td>
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<tr>
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<td></td>
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<td>constant 0.0001491325</td>
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<td></td>
<td>@LRHET</td>
<td>constant 0.00000000</td>
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<td></td>
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<td></td>
<td>@WHITEHT</td>
<td>constant 0.18361479</td>
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<td>@BPHEMT</td>
<td>constant 0.18599239</td>
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<tr>
<td></td>
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</tr>
<tr>
<td></td>
<td>@SWILK</td>
<td>constant 0.098324680</td>
</tr>
</tbody>
</table>
Options

**BPLIST** = list of variables for the Breusch-Pagan heteroscedasticity test.

**CALC/NOCALC** indicates whether the listed diagnostics (list of output names) should or should not be calculated and stored under @names.

**CHOWDATE** = starting date of second period for Chow test. The default is to split the sample exactly in half (if the number of observations is odd, the extra observation will be in the second period).

**DWPVALUE** = APPROX or BOUNDS or EXACT specifies what method will be used for computing the P-value for the Durbin-Watson statistic. The default depends on the current FREQ: APPROX for FREQ N, BOUNDS for other frequencies, including Panel data.

**LMLAGS** = maximum number of lagged residuals for Breusch-Godfrey LM test of general autocorrelation (AR or MA). The default is zero.

**PRINT/NOPRINT** indicates whether the diagnostics should be printed. PRINT implies CALC.

**PVCALC/NOPVCALC** indicates whether p-values should be calculated and stored under %names. PVCALC implies CALC. See Method for the distributions used to compute these P-values in particular cases.

**PVPRINT/NOPVPRINT** indicates whether p-values should be printed. PVPRINT implies PVCALC, PRINT, and CALC. Using this option will sometimes cause regression output to be printed in one column instead of two, unless SHORTLAB is used. Other things like wide numbers (OPTIONS NWIDTH=, SIGNIF=) may also cause single column output.

**QLAGS** = maximum number of autocorrelations for Ljung-Box Q-statistics (Portmanteau test of residual autocorrelation). The default is zero.

**RESETORD** = order of Ramsey’s RESET test. The default is 2.

**SHORTLAB/NOSHORTL** indicates whether short or long labels are used when printing all diagnostics.

**STAR1** = upper bound on p-value for printing at least one star (*), when STARS option is on. The default is .05. There can be up to 5 pairs of (STAR1,STAR2) values, which can apply to different sets of diagnostics. This option only applies to the diagnostics listed for the REGOPT command.

**STAR2** = upper bound on p-value for printing two stars (**), when STARS option is on. The default is .01. This option only applies to the diagnostics listed for the REGOPT command.
REGOPT

STARS/NOSTARS indicates whether stars should be printed indicating significance of diagnostics. STARS implies PVCALC, except for regression coefficients (@T).

Examples

REGOPT (STARS, LMLAGS=5, QLAGS=5, BPLIST=(C,X,X2)) ALL;

turns on all possible diagnostic output, including VCOV matrix and residual plots.

REGOPT;

restores the default settings.

REGOPT (NOCALC) AUTO;

stops calculation of all the autocorrelation diagnostics (useful for pure cross-sectional datasets).

REGOPT (NOPRINT) RSQ FST;

suppresses printing of the R-squared and F-statistics. This is the same as the old TSP command SUPRES RSQ FST;

REGOPT (STARS, STAR1=.10, STAR2=.05) T ;
REGOPT (,STARS, STAR1=.05, STAR2=.02) AUTO ;

uses one set of significance levels for the t-statistics and another for the autocorrelation diagnostics.

Output

Summary table of diagnostics/OLSQ output (@Name = value, %Name = p-value)

<table>
<thead>
<tr>
<th>Group</th>
<th>Name</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>None</td>
<td>LHV</td>
<td>Dependent variable name</td>
</tr>
<tr>
<td></td>
<td>SMPL</td>
<td>Current sample</td>
</tr>
<tr>
<td></td>
<td>NOB</td>
<td>Number of observations</td>
</tr>
<tr>
<td></td>
<td>COEF</td>
<td>Regression coefficients</td>
</tr>
<tr>
<td></td>
<td>SES</td>
<td>Standard errors</td>
</tr>
<tr>
<td></td>
<td>T</td>
<td>t-statistics</td>
</tr>
<tr>
<td></td>
<td>VCOV</td>
<td>Variance-covariance matrix</td>
</tr>
<tr>
<td></td>
<td>VCOR</td>
<td>Correlation version of VCOV</td>
</tr>
<tr>
<td></td>
<td>NCOEF</td>
<td>Number of coefficients</td>
</tr>
<tr>
<td>REGOPT</td>
<td>Description</td>
<td></td>
</tr>
<tr>
<td>------------------------</td>
<td>-----------------------------------------------------------------------------</td>
<td></td>
</tr>
<tr>
<td>NCID</td>
<td>Number of identified coefficients (rank of VCOV)</td>
<td></td>
</tr>
<tr>
<td>REGOUT</td>
<td>Number of identified coefficients (rank of VCOV)</td>
<td></td>
</tr>
<tr>
<td>YMEAN</td>
<td>Mean of dependent variable</td>
<td></td>
</tr>
<tr>
<td>SDEV</td>
<td>Standard deviation of dependent variable</td>
<td></td>
</tr>
<tr>
<td>SSR</td>
<td>Sum of squared residuals</td>
<td></td>
</tr>
<tr>
<td>S2</td>
<td>Estimated variance of residuals (SSR/(NOB-NCID))</td>
<td></td>
</tr>
<tr>
<td>S</td>
<td>Standard error of residuals (SQRT(S2))</td>
<td></td>
</tr>
<tr>
<td>RSQ</td>
<td>R-squared (squared correlation between actual and fitted)</td>
<td></td>
</tr>
<tr>
<td>ARSQ</td>
<td>Adjusted R-squared (adjusted for number of RHS variables)</td>
<td></td>
</tr>
<tr>
<td>AUTO</td>
<td>Durbin-Watson statistic</td>
<td></td>
</tr>
<tr>
<td>DW</td>
<td>Durbin's h statistic (for single lagged dependent var.)</td>
<td></td>
</tr>
<tr>
<td>DH</td>
<td>Durbin's h statistic (for single lagged dependent var.)</td>
<td></td>
</tr>
<tr>
<td>DHALT</td>
<td>Durbin's h alternative (for any lagged dependent)</td>
<td></td>
</tr>
<tr>
<td>LMARx</td>
<td>Breusch-Godfrey LM test for autocorrelation of order x</td>
<td></td>
</tr>
<tr>
<td>QSTATx</td>
<td>Ljung-Box Q statistic for autocorrelation of order x</td>
<td></td>
</tr>
<tr>
<td>WNLAR</td>
<td>Wald test for nonlinear AR1 restriction vs. Y(-1), X(-1)</td>
<td></td>
</tr>
<tr>
<td>ARCH</td>
<td>Test for ARCH(1) residuals</td>
<td></td>
</tr>
<tr>
<td>RECRE</td>
<td>Recursive residuals</td>
<td></td>
</tr>
<tr>
<td>CUSUM</td>
<td>CUSUM plot</td>
<td></td>
</tr>
<tr>
<td>CUSUMSQ</td>
<td>CUSUMSQ plot</td>
<td></td>
</tr>
<tr>
<td>CSMAX</td>
<td>CUSUM test statistic</td>
<td></td>
</tr>
<tr>
<td>CSQMAX</td>
<td>CUSUMSQ test statistic</td>
<td></td>
</tr>
<tr>
<td>CHOW</td>
<td>F-test for stability of coefficients (split sample)</td>
<td></td>
</tr>
<tr>
<td>CHOWHET</td>
<td>Test for stability of coefficients with heteroskedasticity</td>
<td></td>
</tr>
<tr>
<td>LRHET</td>
<td>LR test for heteroscedasticity in split sample</td>
<td></td>
</tr>
<tr>
<td>HET</td>
<td>White het. test on cross-products of RHS variables</td>
<td></td>
</tr>
<tr>
<td>WHITEHT</td>
<td>White het. test on cross-products of RHS variables</td>
<td></td>
</tr>
<tr>
<td>BPHET</td>
<td>Breusch-Pagan het. test on user-supplied list of vars</td>
<td></td>
</tr>
<tr>
<td>LMHET</td>
<td>simple LM het. test on squared fitted values</td>
<td></td>
</tr>
<tr>
<td>None</td>
<td>F-statistic for zero slope coefficients</td>
<td></td>
</tr>
<tr>
<td>FST</td>
<td>Ramsey's RESET test of order x</td>
<td></td>
</tr>
<tr>
<td>RESETx</td>
<td>Ramsey's RESET test of order x</td>
<td></td>
</tr>
<tr>
<td>JB</td>
<td>Jarque-Bera (LM) normality test</td>
<td></td>
</tr>
<tr>
<td>SWILK</td>
<td>Shapiro-Wilk normality test</td>
<td></td>
</tr>
<tr>
<td>AIC</td>
<td>Akaike Information Criterion</td>
<td></td>
</tr>
<tr>
<td>SBIC</td>
<td>Schwarz Bayesian Information Criterion</td>
<td></td>
</tr>
</tbody>
</table>
**Method/Notes on specific diagnostics:**

**DW** ignores sample gaps except when there is PANEL data. The DWPVALUE option can be used to choose one of the 3 methods of calculating its P-value. EXACT computes the \((T-K)\) nonzero eigenvalues of the matrix:

\[
A - \Delta \chi (\chi' \chi)^{1/2} \Delta \chi'
\]

and then uses the Farebrother/Pan method to compute the P-value from the DW and these eigenvalues.

The APPROX method is a small sample adjustment to the asymptotic distribution, using a nonlinear regression fit to the 5% \(dL\) (lower bound) table:

\[
p-value = \Phi \left( DW - 2.0 + 0.58325 \times 10^{-4} + (-0.545221 + 1.5045 \times (K-1)T^{-0.0046}) \left( \frac{\sqrt{T}}{2} \right) \right)
\]

where \(\Phi\) is the cumulative normal. This usually provides a conservative test (i.e. P-value larger than the EXACT method, like the larger number from BOUNDS).

The BOUNDS method calculates the minimum and maximum possible P-values for a given DW, using the minimum and maximum possible sets of eigenvalues for \(K\) and \(T\), stored as %DWL and %DWU. See Bhargava et al (1982) for more details on bounds. For OLSQ with lagged dependent variables, DW is biased, so DH and/or DHALT are computed as well (if the dependent variable is lagged explicitly using a -lag expression).

The optional AUTO and HET diagnostics are not calculated for regressions with weights, instruments, or perfect fits; nor when there are any gaps in the SMPL (to simplify the processing of lags). Note that some of the later diagnostics grouped under AUTO are not strictly for autocorrelation but for heteroskedasticity or structural stability in datasets with a natural time ordering.

DH is not calculated when it involves taking the square root of a negative value. DHALT can be used in all cases (it uses the same regression as LMAR1).
LMARx prints a series of test statistics if LMLAGS is greater than 1. The sample size is adjusted downwards with each test, and the reported statistic is \((p+k-1)F\), asymptotically distributed as chi-squared\((p)\), where \(p\) is the number of lags. QSTATx also prints a series of test statistics (using QLAGS). See Harvey (1990), pp. 174-175.

WNLAR is a Wald test for AR(1) residuals versus mis-specified dynamics (left out lagged dependent and independent variables). If the original equation was \(Y = A + XB\), the regression

\[ Y = A2 +XB + RHO*Y(-1) + D*X(-1) \]

is run, and the restriction \(D = -B*RHO\) is tested. This is asymptotically distributed as chi-squared with degrees of freedom equal to the number of non-singular coefficients on the lagged \(Xs\). WNLAR is the same as COMFAC in \texttt{AR1} (\texttt{OBJFN=GLS}).

ADF is no longer computed here. See the \texttt{COINT} command.

ARCH is a regression of the squared residual on the lagged squared residual.

RECRRES are recursive residuals, calculated using a Kalman Filter (see the \texttt{KALMAN} command). You can display CUSUM and CUSUMQ plots by turning on the \texttt{PLOTS} option. RECRRES can also be used for the Von-Neumann ratio test for autocorrelation.

CHOW is an F-test for parameter stability. The default is to split the sample into equal halves, but the CHOWDATE option can be used to choose an unequal split. If there are insufficient degrees of freedom in one of the halves, the test is still valid, but it is usually not very powerful. The CHOWHET test is robust to simple heteroskedasticity and is the MAC2 test from Thursby (1992). Note that the Chow test does not have the assumed F distribution under heteroscedasticity.

LRHET is a likelihood ratio test for heteroscedasticity between the two periods in the same sample division as the Chow test.

\[
LRHET = T \log \left( \frac{SSR}{T-k} \right) - T_1 \log \left( \frac{SSR_1}{T_1-k} \right) - T_2 \log \left( \frac{SSR_2}{T_2-k} \right)
\]

WHITEHT is a regression of the squared residual on cross-products of the RHS variables. If the model is

\[ Y = B0 + B1*X1 + B2*X2 \]
and the residuals are $E$, the regression

$$E' E = A_0 + A_1 X_1 + A_2 X_2 + A_3 X_1 X_1 + A_4 X_1 X_2 + A_5 X_2 X_2$$

is calculated (if there are sufficient degrees of freedom).

$$TR^2 \sim \chi^2(5)$$

for this example.

BPHET is the same as WHITEHT, except the user specifies a presumably more general list of variables in the $E' E$ regression with the BPLIST option. Note that the ARCH command with the GT option can also be used to estimate such general heteroskedastic regression models.

LMHET is the same as WHITEHT and BPHET, where the squared residuals are regressed on a constant term and the squared fitted values.

RESET is Ramsey's RESET test, where the residuals are regressed on the original right hand side variables and powers of the fitted values. The default order (2) is basically a check for missing quadratic terms and interactions for the right hand side variables. It may also be significant if a quadratic functional form happens to fit outliers in the data.

JB is a powerful joint Lagrange Multiplier test of the residuals' skewness and kurtosis. It is asymptotically distributed as a chi-squared with two degrees of freedom under the null of normality. Small sample critical values are:

<table>
<thead>
<tr>
<th>#obs</th>
<th>5%</th>
<th>10%</th>
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</thead>
<tbody>
<tr>
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<td>3.26</td>
<td>2.13</td>
</tr>
<tr>
<td>30</td>
<td>3.71</td>
<td>2.49</td>
</tr>
<tr>
<td>40</td>
<td>3.99</td>
<td>2.70</td>
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<tr>
<td>50</td>
<td>4.26</td>
<td>2.90</td>
</tr>
<tr>
<td>75</td>
<td>4.27</td>
<td>3.09</td>
</tr>
<tr>
<td>100</td>
<td>4.29</td>
<td>3.14</td>
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<tr>
<td>125</td>
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<td>3.31</td>
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<td>150</td>
<td>4.39</td>
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<td>200</td>
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<td>3.48</td>
</tr>
<tr>
<td>250</td>
<td>4.51</td>
<td>3.54</td>
</tr>
<tr>
<td>300</td>
<td>4.60</td>
<td>3.68</td>
</tr>
<tr>
<td>400</td>
<td>4.74</td>
<td>3.76</td>
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<td>500</td>
<td>4.82</td>
<td>3.91</td>
</tr>
<tr>
<td>800</td>
<td>5.46</td>
<td>4.32</td>
</tr>
<tr>
<td>inf</td>
<td>5.99</td>
<td>4.61</td>
</tr>
</tbody>
</table>
SWLIK is a normality test based on normal order statistics, which has good power in small samples. Since it involves sorting the residuals, it may be quite slow in large samples. The test and its P-value are computed using Royston (1995), with code from Statlib.

AIC (Akaike Information Criterion) and/or SBIC (Schwarz Bayesian Information Criterion) can be minimized to select regressors in a model, such as choosing the length of a distributed lag. SBIC has optimal properties, see Geweke (1981). In general, these can be defined as

$$\hat{AIC} = -\hat{LOGL} + \hat{NCID}$$
$$\hat{SBIC} = -\hat{LOGL} + \hat{NCID} \cdot \log(\hat{NOB})/2$$

LOGL will include the sum of log weights if the OLSQ (WTYPE=HET,WEIGHT=x) option is used. The alternative is the default WTYPE=REPEAT.

Distributions used for P-values:

Note: in all cases, k is the number of identified coefficients in the model, including the intercept.

<table>
<thead>
<tr>
<th>Test Statistic</th>
<th>Null</th>
<th>Alternative</th>
<th>Distribution</th>
<th>Degrees of Freedom</th>
</tr>
</thead>
<tbody>
<tr>
<td>DW</td>
<td>No autocorrelation</td>
<td>Positive autocorrelation (usually)</td>
<td>ratio of Qform</td>
<td>--</td>
</tr>
<tr>
<td>DH</td>
<td>No autocorrelation</td>
<td>--</td>
<td>Normal</td>
<td>--</td>
</tr>
<tr>
<td>DHALT</td>
<td>No autocorrelation</td>
<td>--</td>
<td>Normal</td>
<td>--</td>
</tr>
<tr>
<td>LMARx</td>
<td>No autocorrelation</td>
<td>Autocorrelation of order x</td>
<td>Chi-squared</td>
<td>p+k-1</td>
</tr>
<tr>
<td>QSTATx</td>
<td>No autocorrelation</td>
<td>Autocorrelation of order x</td>
<td>Chi-squared</td>
<td>p ?</td>
</tr>
<tr>
<td>WNLAR</td>
<td>AR(1) disturbance</td>
<td>Other dynamics</td>
<td>Chi-squared</td>
<td># rhs vars</td>
</tr>
<tr>
<td>ARCH</td>
<td>Homoskedasticity</td>
<td>ARCH(1) disturbance</td>
<td>Chi-squared</td>
<td>1</td>
</tr>
<tr>
<td>CSMAX</td>
<td>Stable parameters</td>
<td>Parameters change</td>
<td>Durbin (1971)</td>
<td>--</td>
</tr>
<tr>
<td>CSQMAX</td>
<td>Stable parameters</td>
<td>Parameters change</td>
<td>Durbin (1969)</td>
<td>--</td>
</tr>
<tr>
<td>CHOW</td>
<td>Stable parameters</td>
<td>Parameters differ between two periods</td>
<td>F</td>
<td>(k, nob-2k) usually</td>
</tr>
<tr>
<td>CHOWHET</td>
<td>Stable parameters; variances differ</td>
<td>Parameters and variances differ between two periods</td>
<td>similar to F</td>
<td>(k, nob-2k) usually</td>
</tr>
<tr>
<td>LRHET</td>
<td>Homoskedasticity</td>
<td>Two variances for split sample</td>
<td>Chi-squared</td>
<td>1</td>
</tr>
<tr>
<td>LMHET</td>
<td>Homoskedasticity</td>
<td>Heteroskedasticity related to @FIT**2</td>
<td>Chi-squared</td>
<td>1</td>
</tr>
<tr>
<td>WHITEHT</td>
<td>Homoskedasticity</td>
<td>X-related Heteroskedasticity</td>
<td>Chi-squared</td>
<td>((k+1)k / 2) - 1</td>
</tr>
<tr>
<td>BPHET</td>
<td>Homoskedasticity</td>
<td>Heteroskedasticity</td>
<td>Chi-squared</td>
<td>#vars in</td>
</tr>
</tbody>
</table>
### Related to BPLIST

<table>
<thead>
<tr>
<th>Test</th>
<th>Description</th>
<th>Condition</th>
<th>Statistic</th>
<th>Test Type</th>
</tr>
</thead>
<tbody>
<tr>
<td>FST</td>
<td>Y= constant</td>
<td>Specified regression model</td>
<td>F</td>
<td>BPLIST-1 (k, nob-k)</td>
</tr>
<tr>
<td>JB</td>
<td>Normal disturbances</td>
<td>Non-normal</td>
<td>Chi-squared</td>
<td>2</td>
</tr>
<tr>
<td>SWILK</td>
<td>Non-normal disturbances</td>
<td>Normal</td>
<td>Shapiro-Wilk</td>
<td>--</td>
</tr>
<tr>
<td>RESETx</td>
<td>Higher order terms in Xs needed</td>
<td>F</td>
<td>(RESETORD, nob-k)</td>
<td>nob-k</td>
</tr>
<tr>
<td>T</td>
<td>Slope coefficient =0</td>
<td>Slope coefficient not zero</td>
<td>T (OLS, IV)</td>
<td>Normal (all other procs)</td>
</tr>
</tbody>
</table>

### References


Statlib, [http://lib.stat.cmu.edu/apstat/](http://lib.stat.cmu.edu/apstat/)
RENAME

Examples

RENAME changes the name of an old TSP variable (series, matrix, constant, etc.).

RENAME <old variable name> <new variable name> ;

Output

The name of the variable is changed. If a variable already exists with the new name, it is deleted.

Examples

Save the coefficients from a regression in the vector B1. Note: this is more efficient than COPY, if there is no reason to save the original @COEF.

OLSQ Y C X;
RENAMES @COEF B1;
REPL

Examples

REPL turns on the replacement mode option after it has been turned off with a NOREPL command. This option specifies that series are to be updated rather than completely replaced when the current sample under which they are being computed does not cover the complete series. OPTIONS REPL; is the same as REPL;.

REPL;

Usage

While in REPL mode, if data are created for a sample which overlaps with the SMPL previously used to create the same series, the old values which fall within the current SMPL definition will be replaced, but those outside the current SMPL will remain untouched.

REPL is the default mode and remains in effect until a NOREPL is executed.

Examples

The result of the following sequence of statements:

REPL;
SMPL 1 20; GERNR DATE=1;
SMPL 11 20; GERNR DATE=2;

will be a series DATE which is one for the first 10 observations and 2 for the second ten.

The result of this sequence of statements:

NOREPL;
SMPL 1 20; GERNR DATE=1;
SMPL 11 20; GERNR DATE=2;

is a series DATE which is missing for observations 1 to 10 and equal to two for observations 11 to 20.
RESTORE

Examples

RESTORE reads TSP variables from a file which has been created with the SAVE command.

RESTORE;
or
RESTORE 'filename string';

Usage

RESTORE reads a file named TSPSAV.SAV by default. If a filename string is supplied, the filetype .SAV is appended if it is not present. The RESTORE command is useful for restarting an interactive session which was stopped after issuing a SAVE command.

If a SMPL is already present in the current session, the SMPL in the save file is not restored. Any variables present in the current session with names equal to variables in the save file will be replaced by the save variables.

Output

The current SMPL is printed if it has been restored.

Examples

RESTORE; ? reads TSPSAV.SAV
RESTORE FOO; ? reads FOO.SAV
RETRY (Interactive)

RETRY combines the functions provided by the EDIT and EXEC commands. When editing is complete, execution is automatic.

RETRY [<line number>] ;

Usage

RETRY is identical to EDIT with two exceptions: execution of the modified command is automatic after RETRY, and execution of the EDIT command is not suppressed during collect mode.

The documentation for EDIT in this manual describes the editor, its commands, and provides examples.

In Oxmetrics, Mac and DOS/Windows TSP, it is easier to re-execute a single command using the cursor keys.
REVIEW

REVIEW (Interactive)

REVIEW displays a line or range of lines entered previously in the terminal session, or read from an external file.

REVIEW [<firstline>] , [<lastline>] ;

Usage

REVIEW provides a means of going back and re-examining earlier portions of your terminal session. If the second argument is omitted (indicating end of range) a single line will be displayed. If no arguments are present, the entire terminal session will be listed. The uses are numerous:

- Several commands use line numbers as arguments (EXEC, EDIT, DELETE, RETRY); you will probably need this procedure from time to time to locate commands you want to use again in some way.
- If you get results that puzzle you, you may want to review the sequence of commands that produced them.
- When using INPUT files, and directing the output to disk, you may want a reminder of what has just been executed before proceeding.
- When recovering a session that was terminated abnormally, you will want to REVIEW the command stream before executing selected portions of it.
SAMA performs seasonal adjustment of time series by the moving average method.

*SAMA (ARITH, PRINT) <input series> <output series> ;*

Usage

SAMA is followed by the name of the series to be seasonally adjusted and then the name to be given to the new series. The two series may be the same, in which case the new one will just replace the old one. SAMA should not be used on series which can be negative.

Output

When the print option is off, SAMA produces no printed output. The seasonally adjusted series is stored along with the intermediate results under the following names:

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@SFAC</td>
<td>vector</td>
<td>#periods</td>
<td>Seasonal factors which divide the old series to make new series</td>
</tr>
<tr>
<td>@MOVA</td>
<td>series</td>
<td>#obs</td>
<td>The ratio of the old series to its moving average</td>
</tr>
</tbody>
</table>

If the print option is on, these quantities are also printed in table form.

Method

Denote the series to be adjusted by $X$, indexed by $t$ and $T$ observations in length. The periodicity of the series (the number of periods per year) is $p$, which is customarily 4, in the case of quarterly series, or 12, in the case of monthly series. The ratio of the series to its moving average is formed in the following way:

$$MOVA(t) = \frac{X(t)}{(\text{Moving Average of } X)}$$

where the moving average of $X$ is defined as

- $p=4$: $\frac{1}{4}(X(t-2)/2 + X(t-1) + X(t) + X(t+1) + X(t+2)/2)$
- $p=12$: $\frac{1}{12}(X(t-6)/2 + X(t-5) + \ldots + X(t) + \ldots + X(t+5) + X(t+6)/2)$

This equation describes a weighted moving average over \( p+1 \) observations centered at each observation in turn. The vector \( \text{MOVA}(t) \) may be rewritten as a matrix where each row corresponds to a year and contains \( p \) elements. The total number of non-missing elements in \( \text{MOVA} \) is \( T-p \), because \( p/2 \) observations are dropped at the beginning and end of the series.

The \( p \) seasonal factors are formed by averaging each column in \( \text{MOVA} \):

\[
\text{SFAC}(1) = \frac{1}{(T-1)} \times (\text{MOVA}(2,1)+\text{MOVA}(3,1)+...+\text{MOVA}(T,1))
\]

\[\ldots\]

\[
\text{SFAC}(p) = \frac{1}{(T-1)} \times (\text{MOVA}(1,p)+\text{MOVA}(2,p)+...+\text{MOVA}(T-1,p))
\]

They are normalized to average to unity either arithmetically or geometrically, depending on the option specified. The seasonally adjusted series is then computed by dividing the old series by the seasonal factors.

**Options**

PRINT/NOPRINT specifies that the ratio of the series to its moving average and the computed seasonal factors be printed.

ARITH/NOARITH specifies that the arithmetic mean be used for normalization, rather than a geometric mean.

**Examples**

\[
\text{SAMA (PRINT,ARITH) GNPQ GNPQA;}
\]

\[
\text{SAMA (ARITH) GNPQ GNPQA; PRINT @SFAC @MOVA ;}
\]

These two examples have the same effect. The first prints the seasonal factors and the moving average ratio series. The second suppresses the printing, but then prints @SFAC and @MOVA, which are the same as what would have been printed.

**Reference**

SAMPSEL

Output  Options  Example  References

SAMPSEL estimates a generalized Tobit or sample selection model where both the regression and the latent variable which predicts selection are linear regression functions of the exogenous variables. Either a censored (regression variables not observed for non-selected observations) or truncated (all variables not observed) model may be estimated.

\[
\text{SAMPSEL (MILLS=<series>, nonlinear options) <probit dep var>}
\]
\[
<\text{probit indep vars}> \mid <\text{regression dep var}> <\text{regression indep vars}> ;
\]

Usage

The model estimated by SAMPSEL is the Tobit type II model described by Amemiya or the censored regression model with a stochastic threshold described by Maddala (see the references). It can be written as

\[
y_{2i} = \begin{cases} X_{2i} \beta + e_i & \text{if } y_{1i} > 0 \\ 0 & \text{otherwise} \end{cases}
\]
regression equation

\[
y_{1i} = X_{1i} \delta + u_i
\]
selection equation

e(i) and u(i) are assumed to be joint normally distributed:

\[
\begin{pmatrix} e_i \\ u_i \end{pmatrix} \sim N\left( \begin{pmatrix} 0 \\ 0 \end{pmatrix}, \begin{pmatrix} \sigma^2 & \rho \sigma \\ \rho \sigma & 1 \end{pmatrix} \right)
\]

In the output, the standard deviation of the regression equation is denoted SIGMA and the correlation coefficient is denoted RHO. The variance of the selection (probit) equation is normalized to one without loss of generality.

To use the procedure to estimate this model, supply the name of a zero/one variable which tells whether the observation was observed or not \((y(1) > 0)\) as the probit dependent variable, the regressors \(X_1\) as the probit independent variables, \(y(2)\) as the regression dependent variable, and \(X_2\) as the regression independent variables. Missing values for the regression variables are allowed for those observations for which \(y(1) = 0\).

If \(y(1)\) is always greater than zero, the truncated (conditional) model is estimated (Bloom and Killingsworth 1985). This is flagged with the message "Latent Selection Variable". The identifying condition that there be variables other than the constant in the probit equation is not checked.
Output

The output of SAMPSEL begins with an equation title and the name of the dependent variable. Starting values and diagnostic output from the iterations will be printed. Final convergence status is printed. This is followed by the mean of the dependent variable, number of positive observations, sum of squared residuals, R-squared, and a table of right hand side variable names, estimated coefficients, standard errors and associated t-statistics.

SAMPSEL also stores some of these results in data storage for later use. The table below lists the results available after a SAMPSEL command.

<table>
<thead>
<tr>
<th>name</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@LHV</td>
<td>list</td>
<td>1</td>
<td>Name of dependent variable</td>
</tr>
<tr>
<td>@RNMS</td>
<td>list</td>
<td>#vars</td>
<td>list of names of right hand side variables</td>
</tr>
<tr>
<td>@YMEAN</td>
<td>scalar</td>
<td>1</td>
<td>Mean of the probit dependent variable</td>
</tr>
<tr>
<td>@NOB</td>
<td>scalar</td>
<td>1</td>
<td>Number of observations</td>
</tr>
<tr>
<td>@NPOS</td>
<td>scalar</td>
<td>1</td>
<td>Number of positive observations in probit equation</td>
</tr>
<tr>
<td>@SSR</td>
<td>scalar</td>
<td>1</td>
<td>Sum of squared residuals (regression equation)</td>
</tr>
<tr>
<td>@LOGL</td>
<td>scalar</td>
<td>1</td>
<td>Log of likelihood function</td>
</tr>
<tr>
<td>@SBIC</td>
<td>scalar</td>
<td>1</td>
<td>Schwarz Bayesian Information Criterion</td>
</tr>
<tr>
<td>@AIC</td>
<td>scalar</td>
<td>1</td>
<td>Akaike Information Criterion</td>
</tr>
<tr>
<td>@IFCONV</td>
<td>scalar</td>
<td>1</td>
<td>1 if convergence achieved, 0 otherwise</td>
</tr>
<tr>
<td>@NCOEF</td>
<td>scalar</td>
<td>1</td>
<td>Number of coefficients = number in probit + number in regression + 2</td>
</tr>
<tr>
<td>@NCID</td>
<td>scalar</td>
<td>1</td>
<td>Number of identified coefficients</td>
</tr>
<tr>
<td>@COEF</td>
<td>vector</td>
<td>#coeffs</td>
<td>Coefficient estimates</td>
</tr>
<tr>
<td>@SES</td>
<td>vector</td>
<td>#coeffs</td>
<td>Standard errors</td>
</tr>
<tr>
<td>@T</td>
<td>vector</td>
<td>#coeffs</td>
<td>T-statistics</td>
</tr>
<tr>
<td>@GRAD</td>
<td>vector</td>
<td>#coeffs</td>
<td>Gradient of log likelihood at convergence</td>
</tr>
<tr>
<td>@VCOV</td>
<td>matrix</td>
<td>#coeffs*#coeffs</td>
<td>Variance-covariance of estimated coefficients</td>
</tr>
<tr>
<td>@DPDX</td>
<td>matrix</td>
<td>#selvars*2</td>
<td>Mean of probability derivatives for selection equation</td>
</tr>
<tr>
<td>@RES</td>
<td>series</td>
<td>#obs</td>
<td>Residuals for the observed sample</td>
</tr>
<tr>
<td>@MILLS</td>
<td>series</td>
<td>#obs</td>
<td>Inverse Mills ratios</td>
</tr>
</tbody>
</table>

If the regression includes a PDL variable, the following will also be stored:

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Method

The method used is maximum likelihood, obtained by means of a gradient method that uses the Hessian approximation given by the HITER option. Since this likelihood function is known to have multiple local optima frequently, the method of Nawata (1994, 1995, 1996) is used to find the global optimum. In Nawata's method, a grid search is done on the correlation coefficient RHO to find the set of local optima. Then further iterations are done to refine the optima to full precision and choose the global optimum. The grid points used are 0, .1, .2, ..., .8, .85, .9, .95, .99, .9999, -.1, -.2, ..., -.8, -.85, -.9, -.95, -.99, -.9999.

Sometimes the global optimum shows RHO = 1.0000 or -1.0000. In these cases, the actual estimate of RHO is slightly less than 1 in absolute value, and the residual covariance matrix is nearly singular. The standard error of RHO and its covariance with other parameters is set to zero in these cases.

Options

MILLS= name of variable where the inverse Mills ratio should be stored. The default is @MILLS.

Standard nonlinear options -- see NONLINEAR.

HITER=N, HCOV=N is the default.

Example

SAMPSSEL (PRINT, MAXIT=50, HCOV=NBW) IY C Z | Y C X ;

References


SAVE (Interactive)

**Examples**

SAVE writes all current user-defined TSP variables into a file which may be restored later with the RESTORE command.

**SAVE ['filename string'] ;**

**Usage**

SAVE creates a file named TSPSAV.SAV by default. If a filename string is supplied, the filetype .SAV is appended if it is not present. This file contains all user-defined variables including series, parameters, constants, matrices, formulas and lists. It also contains the current SMPL.

The SAVE command is useful for stopping an interactive session and restarting it at a later time. It is also useful for preserving a session environment if it is likely that later commands or a power failure may cause TSP to abort. In general, it is better to store TSP variables in databanks or regular files, since it is easier to determine the origins and contents of such files, and these files are transportable to other computers.

SAVE creates a binary file which contains TSP variable names and types, alternating with the actual data for each variable. The RESTORE command can read this file and recreate the TSP variables as they existed when the SAVE command was executed. Variables whose names begin with @ are not saved; it is assumed they are intermediate results from TSP procedures, like @RES which is usually created by an estimation procedure.

**Examples**

**SAVE;**

creates TSPSAV.SAV. If TSPSAV.SAV already exists, it is destroyed.

**SAVE FOO;**

creates FOO.SAV.
SELECT

**Options  Examples**

SELECT selects a subsample of observations from the last SMPL statement.

**SELECT (SILENT, PRINT) <logical expression> ;**

**Usage**

SELECT is exactly the same as SMPLIF, except it operates on the last SMPL statement instead of the last SMPLIF or SELECT statement. This means that consecutive SELECT statements are independent -- they do not create a nested sequence of subsamples. This is more convenient than saving and restoring the previous SMPL manually.

**Output**

Some pairs of sample observations are printed, unless the SILENT option or SUPRES @SMPL; has been used.

**Options**

PRINT/NOPRINT prints the full set of sample pairs resulting from SELECT. Normally only one line of output is printed.

SILENT/NOSILENT prints no output at all. Same as SUPRES SMPL; before SELECT.

**Examples**

**SMPL 1,10; TREND T;**
**SELECT T > 5; MSD T;**
**SELECT T > 2 & T < 9 ; MSD T;**

creates subsamples 6,10 and 3,8 . The second SELECT statement is equivalent to

**SMPL 1,10;**
**SMPLIF T > 2 & T < 9 ; MSD T;**

**SELECT 1;**

can be used to return to the last SMPL statement.
SET

Examples

SET performs computations on scalar variables and single elements of time series or matrices.

\[ \text{SET} \text{ <scalar>} = \text{<algebraic expression>} ; \]

or

\[ \text{<subscripted variable>} = \text{<algebraic expression>} ; \]

Usage

SET consists of the name of a scalar variable or an element of a series or matrix followed by an equal (=) sign and an arbitrary algebraic expression. The expression must follow the usual TSP rules for formulas.

The formula is evaluated using the current values of all the variables and the result is stored in the variable on the left hand side of the equation. If the left hand side is an element in a series and matrix, only that particular element is changed; the remainder of the variable is unchanged (whether or not the REPL mode is on).

Legal forms of scalar variables in TSP are the following (these forms can be used wherever scalars are allowed):

1. A simple variable name such as BETA, or D1. This could be already defined by a CONST or PARAM, but this is not required.

2. A subscripted series, such as GNP(I), GNP(72:1), GNP(72) or YOUNG(40). If the frequency is NONE, you can use a simple subscript in the same units as your sample. If the frequency is QUARTERLY (4), or MONTHLY (12), use a valid TSP date as the subscript. A variable (4 characters or less) can also be used on dated and undated series. Since variables do not take date values (except for annual frequency), a variable subscript is always relative to the start of the current SMPL. For example, the following loop fills the series X throughout the current SMPL:

\[
\text{SMPL 48:1,86:2; DO } l=1,@NOB; \text{ SET } X(l) = ... ; \text{ ENDDO;}\]

3. A subscripted matrix. A matrix may have a single numeric or variable subscript, which is computed by the following formula:
\begin{align*}
\text{subscript} &= (j-1)\times NROW + i \\
\text{where } i \text{ is the row index of the element and } j \text{ is the column index.}
\end{align*}

Matrices may also be doubly subscripted, but any variable subscripts must be 2 characters or less. Here are some examples of legal matrix elements:

\begin{align*}
XL(I,J) \quad XL(2,6) \quad MAT(345) \quad MAT(I) \quad XL(II,247) \quad A(K1,LL)
\end{align*}

These are illegal:

\begin{align*}
\text{MAT(VARSUB) } \quad XL(L20,2) \quad A(K,L+1) \quad GNP(I-1)
\end{align*}

The first and second are illegal because subscript names are limited to 4 or 2 characters. The third and fourth are illegal because expressions are not allowed as subscripts.

\text{SET} \text{ is not recommended for creating a series or matrix.} \text{ \textit{READ, GENR, TREND, DUMMY,} etc. should be used to create series.} \text{ \textit{READ, MFORM, MMMAKE, or COPY} should be used to create matrices. \text{ \textit{SET} can be used to update series and matrices, or to retrieve particular elements from them.}}

\text{Output}

\text{SET produces no printed output. A scalar is stored in data storage, or a series or matrix is updated.}

\text{Examples}

\begin{align*}
\text{SET VALUE} &= X(1,1) \\
\text{SET SE} &= @S \\
\text{MATRIX}(2,3) &= A+B^2 / (\text{LOG(LABW)}) \\
\end{align*}

\begin{align*}
\text{FREQ A;} \\
\text{SMPL 1983 1990;} \\
\text{GENR PFOR} &= 100; \\
\text{DO } I &= 1984 \text{ TO } 1990; \\
& \quad \text{SET } I1 = I-1; \\
& \quad \text{SET PFOR}(I) = PFOR(I1)\times \text{EXP}(1.0 + \text{DELTA}); \\
\text{ENDD;}
\end{align*}

\text{The last example above shows how \textit{SET} can be used to compute a series in which each observation is a dynamic function of a previous observation. This can be done more efficiently, however, by replacing the last DO loop with a dynamic GENR:}

\begin{align*}
\text{SMPL 84,90;} \\
\text{PFOR} &= \text{PFOR(-1)}\times \text{EXP}(1.0 + \text{DELTA});
\end{align*}
SHOW displays information about specific symbols, or classes of symbols. It can also be used to display the internal limits on data in TSP.

SHOW (DATE, DOC) [ <list of symbols>, SMPL, FREQ, ALL, EQUATION, LIST, MATRIX, MODEL, PROC, SCALAR, SERIES ] ;

Usage

SHOW may be used at any time during the interactive session to obtain information about symbols, and how they have been stored. Symbol names and class names may be freely mixed as arguments to the SHOW command. The classes are: EQUATION, LIST, MATRIX, MODEL, PROC, SCALAR, and SERIES. Providing a class name will list all symbols belonging to that class. SMPL and FREQ will display the current setting of each, and ALL will display all symbols, most recent entries first. Unique abbreviations of class names are allowed.

SHOW LIST ;  \( ? \) provides a list of all <listnames>

? provide information about all members of the particular list:
SHOW listname ;

SHOW PROC ;
or
SHOW procname ;

are useful if you want to call one of the procedures you have defined, but do not remember the number or order of the arguments to pass it.

SHOW by itself lists the internal array dimensions in TSP. This is helpful if you have a very large problem.

Output

SHOW SERIES stores a list of all the series under @RNMS.

<table>
<thead>
<tr>
<th>class</th>
<th>information provided by SHOW</th>
</tr>
</thead>
<tbody>
<tr>
<td>EQUATION</td>
<td>name, type (formula or identity), number of arguments and operations</td>
</tr>
<tr>
<td>LIST</td>
<td>name, # of members</td>
</tr>
<tr>
<td>MATRIX</td>
<td>name, dimensions, type</td>
</tr>
<tr>
<td>MODEL</td>
<td>name</td>
</tr>
</tbody>
</table>
SHOW

PROC       name, formal arguments
SCALAR     name, type, value
SERIES     name, #obs, beg date-end date, frequency

Options

DATE/NODATE prints the last date modified on a separate line (if the variable has documentation created with the DOC command).

DOC/NODOC prints any documentation on a separate line. When DOC is off, a portion of the documentation which fits on the end of the current line is printed.

Examples

Assume the following TSP commands have been given:

SMPL 1,10;
TREND T; T2=T*T;
OLSQ T2 C T;

SHOW would then produce the following results:

SHOW SERIES ;

Class Name Description
------- ------- --------------
SERIES @RES 10 obs. from 1-10, no frequency
          @FIT 10 obs. from 1-10, no frequency
          T2 10 obs. from 1-10, no frequency
          T  10 obs. from 1-10, no frequency

SHOW MATRIX ;

Class Name Description
------- ------- --------------
MATRIX @VCOV 2x2 symmetric
         @SES 2x1 general
         @COEF 2x1 general
         @SMPL vector, length 2
SIML solves linear and nonlinear simultaneous equation models using Newton's method with an analytic Jacobian. The model is solved period by period; if a dynamic simulation (the default) is specified, the solved values for lagged endogenous variables are fed forward to later periods.

SIML is a more powerful and more working space-intensive alternative to the SOLVE procedure. Use SIML if your model is highly nonlinear and difficult to solve with the conventional Gauss-Seidel recursive algorithms. SIML is also recommended for any small model (less than about 25 equations) because of its ease of use, requiring less setup cost than SOLVE. If the model is linear, SIML will solve it in one iteration (per time period).

**SIML (DEBUG, DYNAM or STATIC, ENDOG=(<list of endogenous variables>), METHOD=NEWTON or GAUSSN, PRNDAT, PRNRES, PRNSIM, SILENT, TAG=<tagname> or NONE, nonlinear options) <list of equation names> ;**

**Usage**

To simulate a model with the options set at default values, specify SIML with the ENDOG option to give the list of variables to be solved for and then a list of equations which are to be solved. These equations are specified earlier with FRML or IDENT statements. There is no difference between the two types of equations in SIML - for either one, the model solution tries to make the error as small as possible.

Equations for SIML can be the exact same ones which you estimated using FIML or LSQ. If you want to have linear equations in your model from OLSQ, INST, or AR1 estimation, use FORM to make them after the estimation.

Note that there must be as many equations as endogenous variables so that the Jacobian of the model is a square matrix.

SIML solves the model specified by the equations over the current SMPL, one period at a time. The starting values for the variables are chosen as follows:

1. If the variables already exist, the actual values for the current period are used as starting values, unless they are missing.
2. If the variables do not exist and this is the first period of the simulation, the value one is used as a starting value.
3. If the variables do not exist and this is not the first period of the simulation, the values of the last period solution are used as starting values.

Output

If no options are specified, the normal output from SIML begins with a title and listing of options. This is followed by a table of the data series if the PRNDAT option is on.

Next is the iteration output. If PRINT has not been specified, only one line per iteration is printed, showing the starting value of the objective function, the ending value, and the value of the steps size for this iteration. Even this information is not printed if you have specified the SILENT option.

If PRINT has been specified, considerably more output is produced, showing the values of the endogenous variables and the vector of changes at each iteration. If PRNRES is on, the residual error from each equation is printed when convergence is achieved or the maximum number of iterations is reached. The Jacobian is printed for the first two periods.

After solution of the model over the whole sample, a message is printed if the variables are being saved in data storage. Following this message a table of the results of the simulation is printed, labelled by the observation IDs. To suppress the table, use the NOPRNSIM option. @IFCONV is stored as a series with ones if the simulation for the observation converged, and zeroes otherwise. This may be useful for a convergence check, since this information may otherwise be hidden in a large output file full of iteration information.

Method

If a simultaneous model is linear, it can be written as \( Ax = b \), where \( A \) is a matrix of coefficients, \( x \) is the set of endogenous variables, and \( b \) is a vector of numbers (which may include functions of exogenous variables). This model can be solved directly by inverting \( A \) and multiplying \( b \) by it.

Newton's method applies this idea to the iterative solution of nonlinear models in the following way: At each iteration, the model is linearized in its variables around the values from the previous iteration. The linearized model is solved by matrix inversion. The resulting set of new values is treated as a direction vector for a linear search for a "better" set of values.
The criterion function for SIML is the sum of squared deviations of the equations. At the solution, all the deviations will be zero. Away from the solution, the deviations are computed by substituting the current values of the variables into the equations and evaluating them. This sum of squared deviations is the objective function printed out by SIML at each iteration. When the model is linear in the variables, the model is simply solved by the matrix equation above and no iteration is done.

TSP's implementation of Newton's method uses an analytic Jacobian evaluated at the current variable values as the matrix $A$; this is an extremely powerful method for finding the solution of a nonlinear model, but it can still run into trouble, primarily because of (near)-singularity of the Jacobian. If this happens, a message is printed, and you may wish to try the GAUSSN method, which uses a generalized inverse to try to get past a locally singular point.

**Options**

DEBUG/NODEBUG prints the endogenous variables and direction vector at each iteration - for debugging recalcitrant models.

DYNAM/NODYNAM specifies dynamic simulation. Earlier solved values of lagged endogenous variables are used in place of actual values. STATIC is the alternative to DYNAM. DYNAM is the default, unless there are no lagged endogenous variables.

ENDOG= *(a list of the endogenous variables in the model)*. This is the list of variables for which the model will be solved. They do not have to be predefined, unless you wish to supply starting values, or you are doing a static simulation with lagged endogenous variables.

METHOD= controls the action to be taken if the model becomes singular. For METHOD=NEWTON (the default) iteration stops at a singular point. For METHOD=GAUSSN, a generalized inverse is used (i.e., one or more variables are temporarily excluded from the model and are held constant for the iteration).

PRNRES/NOPRRES prints the residuals when solution is complete for each time period. All residuals will be small enough to satisfy the convergence criterion. The Jacobian for the first two time periods is also printed.

PRNDAT/NOPRNDAT prints all the endogenous and exogenous variables at the beginning of the simulation.

PRNSIM/NOPRNSIM prints a table containing the solved values of the endogenous variables.
SILENT/NOSILENT suppresses all the output.

STATIC/NOSTATIC specifies static simulation. Actual values of lagged endogenous variables are used, not earlier solved values.

TAG= tagname specifies that the solved values of all endogenous variables should be stored as series with names created by adding tagname to the variable names; tagname should be a single character, or perhaps two, to avoid creating excessively long names. (Names larger than the allowed length of a TSP name will be truncated). TAG=NONE stores under the original endogenous names.

The default values of the options are DYNAM, METHOD=NEWTON, and TAG=nothing. The print options are all off except PRNSIM, which prints only a one line summary of each iteration and a table of results.

Examples

This example solves the Illustrative Model described in the User's Manual:

SIML (PRNDAT,TAG=S,ENDOG=(GNP,CONS,I,R,LP))
CONSEQ,INVEQ,INTRSTEQ,GNPID,PRICEQ;

After this model has been solved, the solved series are stored under the names GNPS, CONSS, IS, etc. The input data and the solved series are printed.

The next example shows how to set up the well-known Klein Model I for simulation. The equations are linear, so they are formed after the corresponding instrumental variables estimation.

LIST Z C P(-1) K(-1) E(-1) TM W2 G TX ;
2SLS (INST=Z) CX C P P(-1) W ;
FORM CONS ;
2SLS (INST=Z) I C P P(-1) K(-1) ;
FORM INV ;
2SLS (INST=Z) W1 C E E(-1) TM ;
FORM WAGES ;
IDENT WAGE W = W1+W2 ;
IDENT BALANCE CX+I+G-(TX+W+P) ;
IDENT PPROD E-P-TX-W1 ;
IDENT CAPSTK K=K(-1)+I ;
SIML (TAG=S, ENDOG=(CX,I,W1,W,E,P,K)) CONS INV WAGES WAGE BALANCE PPROD CAPSTK ;
This model solves for $C_X$ (consumption), $I$ (investment), $W_1$ (wages in the private sector), $W$ (total wage bill), $E$ (production of the private sector), $P$ (profits), and $K$ (capital stock) using $T_M$ (time), $W_2$ (government wage bill), $T_X$ (taxes), and $G$ (government expenditures) as exogenous variables.

References


SMPL

Output   Examples

SMPL is used to define the observations of the data which will be used in the following TSP procedures. The SMPL vector is a set of pairs of observation identifiers which define the range(s) of observations which are in the current sample.

\[ \text{SMPL} \ \langle \text{beginning obs. id} \rangle \ \langle \text{ending obs. id} \rangle \ [\langle \text{beginning obs. id} \rangle \ \langle \text{ending obs. id} \rangle \ \ldots \ldots ] ; \]

or

\[ \text{SMPL} \ \text{SMPL\_vector\_name} ; \]

SMPL is often used in conjunction with FREQ, which sets the frequency of the data.

Usage

The sample of observations may be specified in four ways:

1. A SMPL statement listing the beginning and ending pairs of observations to be used.

2. A SMPL statement containing the name of a variable that contains a SMPL vector of pairs of observations ids.

3. A SMPLIF statement with an expression which is true for the observations to include in the sample (see the SMPLIF section).

4. A SELECT statement (same as SMPLIF except it selects observations from the previous SMPL statement instead of the current sample).

The first of these is by far the most common: the simplest form of the SMPL statement just specifies one continuous group of observations. For example, to request that observations 1 through 10 of the data be used, use the command

\[ \text{SMPL} \ 1 \ 10 ; \]

If you want to use more than one group of observations, specify the groups in any order on the SMPL statement. For example, the following SMPL skips observation 11:

\[ \text{SMPL} \ 1,10 \ 12,20; \]
The observation identifiers on the SMPL statement can be any legal observation identifier:

- Simple integers if the frequency is none.
- Years if the frequency is annual. If the year is less than 201 and greater than 0, 1900 will automatically be added; this can be reset with the BASEYEAR= option. (See the OPTIONS command entry for details.)
- Years followed by a colon and the period if the frequency is monthly or quarterly.

SMPL can be changed as often as you like during a TSP program. While a SMPL is in force, no observations on series outside that SMPL will be stored or can be retrieved, unless they are specified with lags (or leads) and the lagged (led) value is within the sample. For example if the sample runs from 48 to 72 and the variable GNP(-1) is specified, the 1947 value of GNP will be used for the 1948 observation of GNP(-1).

**Output**

Every time the SMPL is changed, TSP prints out the current sample unless SUPRES SMPL; has been specified earlier in the program. The sample vector is also stored in data storage under the name @SMPL. The number of observations in the current sample is stored as a scalar, under the name @NOB. This can be quite convenient if you do not know exactly how many observations a SMPLIF or SELECT command will yield, for example.

**Examples**

```plaintext
FREQ A
SMPL 56,80
SMPL 21,40 46,82
FREQ Q SMPL 72:1,82:4
FREQ M SMPL 78:5,81:9
FREQ N SMPL 2,7 9,14 16,21 23,28 30,35
```

The last example specifies groups of six observations at a time, skipping every seventh observation beginning with the first. This is a common arrangement for panel data with a single lagged endogenous variable, but it is easier to use FREQ (PANEL) which automates this sample selection.

Suppose we have a vector called SAMPLE loaded with 2,7,9,14,16,21,23,28, 30,35. Then the last example could also be done with

```plaintext
SMPL SAMPLE
```
SMPLIF

Options  Examples

SMPLIF is used to select a sample of observations based on the values of an expression. The observations in the current sample for which the expression is true are selected.

**SMPLIF (PRINT, SILENT) <logical expression> ;**

Usage

SMPLIF is simply followed by an expression. This expression can be a series, such as a dummy variable, or it can involve several series with logical operators. The expression is used to select observations from the current sample in the following way: if the value of the expression for an observation is greater than zero, the observation is kept, otherwise it is dropped. The resulting SMPL vector replaces the previous one.

Note that SMPLIF, unlike SMPL, chooses only observations within the current sample; you should reset the sample to cover the whole data set if you want to select a different group of observations later. Successive SMPLIFs will nest within each other, resulting in a non-increasing set of observations. Use SELECT for non-nested observation selection.

If the expression is false for all current observations, an empty sample would result. @NOB is stored as zero for this case, but the sample is left unchanged. @NOB should be tested when empty samples are possible.

Output

SMPLIF produces the same output as SMPL: the resulting sample is printed if printing has not been suppressed and the variables @SMPL and @NOB are stored in data storage.

Options

PRINT/NOPRINT prints the full set of sample pairs resulting from SMPLIF. Normally only one line is printed.

SILENT/NOSILENT prints no output at all.
Examples

Delete the first observation for every individual in a panel data set which has six years of data for each of 20 people:

```
SMPL 1 120 ;
TREND(PERIOD=6) YEAR;
SMPLIF YEAR > 1 ;
```

This example shows how logical expressions can be used to select data for estimation:

```
SMPLIF P>0 & R>0 & DER<ELAG ;
```
SOLVE

SOLVE solves linear and nonlinear simultaneous equation models using the Gauss-Seidel method or the Fletcher-Powell method for minimization. The model is solved period by period; if a dynamic simulation (the default) is specified, the solved values for lagged endogenous variables are fed forward to later periods.

SOLVE is suitable for large, mostly linear, loosely structured models. Since the algorithms are designed to operate on the model in blocks or groups of equations, you can obtain cost savings by using SOLVE instead of SIML when your model is large but not very interrelated - for example, it includes several different sectors.

The Gauss-Seidel algorithm, which is fundamentally a recursive loop through the equations, is the least powerful of the algorithms available in TSP for model simulation. The Fletcher-Powell algorithm, which solves simultaneous blocks by minimizing the sum of squared residuals from each equation, is somewhat more powerful, but not as good as the Newton's method implementation in SIML, since it does not use an analytic Jacobian.

SOLVE (CONV2=<secondary convergence criterion>, DEBUG, DYNAM or STATIC, KILL, MAXPRT=<iterations to be printed>, METHOD=GAUSS or FLPOW or JACOBI, PRNDAT, PRNRES, PRNSIM, TAG=<tagname> or NONE, nonlinear options) <name of collected model> ;

Usage

To simulate a model with SOLVE, first specify all the equations of the model in normalized form, that is, with each endogenous variable appearing once and only once on the left hand side of an equation. These equations may be specified with FRML or IDENT statements. There is no difference between the two types of equations in SOLVE - for either one, the model solution tries to make the error as small as possible. There must be as many equations as endogenous variables.

After the equations are specified, form and order the model with the MODEL procedure. This procedure takes the list of equations and endogenous variables in the model and produces a collected and ordered model which is stored under a name which you supply. This is the name which should appear on the SOLVE statement.

SOLVE solves the model specified over the current SMPL, one period at a time. The starting values for the variables are chosen as follows:
1. If the variables already exist, the actual values for the current period are used as starting values, unless they are missing values.

2. If the variables do not exist and it is the first period of the simulation, the value zero is used as a starting value.

3. If the variables do not exist and this is not the first period of the simulation, the values of the last period solution are used as starting values.

Output

If no options are specified, the normal output from SOLVE begins with a title and listing of options. This is followed by a table of the data series if the PRNDAT option is on. Then comes the iteration output. If PRINT has not been specified, only convergence or non-convergence messages are printed for each block of the model as they are solved, along with the iteration count. If the PRINT option is on, a one line message for each iteration is printed, showing the iteration number, the block number, the objective function (the sum of squared residuals), and the value of the stepsize for this iteration.

If PRNRES is on, the residual error from each equation is printed for the first MAXPRT iterations on each block, and also at convergence of the block. This can help in identifying problem equations if your model is difficult to solve.

After solution of the model over the whole sample, a message is printed if the variables are being saved in data storage. Following this message a table of the results of the simulation is printed, labelled by the observation ids. This table can be suppressed by specifying the NOPRNSIM option. @IFCONV is stored as a series which contains ones if the simulation for the observation converged, and zeroes otherwise. This is useful to check the convergence. The sum of @IFCONV should be equal to @NOB if all periods converged.

Method

The model to be solved is broken into a series of blocks by the procedure MODEL. These blocks are alternately recursive and simultaneous. A recursive block is one which can be solved simply by specifying the value(s) of a set of previously determined endogenous variables and then computing each equation in the block in turn. Each equation must depend only on endogenous variables which are input to the block or computed previously within the block.
Obviously, a recursive block is easily solved on the conditional values of the input endogenous variables. A simultaneous block, on the other hand, does not have a triangular Jacobian and thus requires either the inversion of the Jacobian or some sort of iterative technique. The two methods available in SOLVE are the Gauss-Seidel and the Fletcher-Powell. The first is simply a generalization of the method for computing recursive blocks: the equations are computed in order, each endogenous variable being evaluated in turn. Then the new values of the endogenous variables are used to start the process over again until convergence (no change in the variables) is achieved. This process works best on mildly simultaneous and fairly linear models; it does not guarantee convergence.

The criterion function for SOLVE is the sum of squared deviations of the equations. At the solution, all deviations will be zero. Away from the solution, the deviations are computed by substituting the current values of the variables into the equations and evaluating them. This sum of squared deviations is the objective function printed out by SOLVE at each iteration.

The Fletcher-Powell algorithm solves simultaneous blocks by minimizing this criterion function with respect to the endogenous variables in the block. The method uses numerical first derivatives of the objective function and a rank one updating technique to build up the second derivative matrix. See the references for further information on this method.

Options

**CONV2=** specifies a secondary convergence criterion which is applied to the sum of squared residuals for each block of equation after the variables have passed the standard TOL convergence test.

**DEBUG/NODEBUG** prints the endogenous variables at each iteration. It is useful for debugging recalcitrant models.

**DYNAM/NODYNAM** specifies dynamic simulation. Earlier solved values of lagged endogenous variables are used in place of actual values. STATIC is the alternative to DYNAM. DYNAM is the default, unless there are no lagged endogenous variables.

**KILL/NOKILL** specifies whether the simulation is to be stopped if convergence fails for any block or period. If you use the DYNAM option, you may wish to specify the KILL option since the simulated results will feed forward.

**MAXPRT=** the number of iterations for which printout of the residuals is to be produced. This has no effect unless PRNRES is on. In that case, the default value is 5.
METHOD=GAUSS or JACOBI or FLPOW specifies the iteration method to be used. The JACOBI method is a variation of the Gauss-Seidel method in which the endogenous variables for any simultaneous block are all updated at once at the beginning of the iteration (see Ortega and Rheinboldt, pp. 217-220).

PRNRES/NOPRNRES prints the residuals when solution is complete for each time period. All of the residuals will be small enough to satisfy the convergence criterion.

PRNDAT/NOPRNDAT prints the starting values for the endogenous variables at each time period.

PRNSIM/NOPRNSIM prints a table of the solved values of the endogenous variables at the completion of the simulation.

STATIC/NOSTATIC specifies static simulation. Actual values of lagged endogenous variables are used, not earlier solved values.

TAG= tagname specifies that the solved values of all endogenous variables should be stored as series with names created by adding tagname to the variable names; tagname should be a single character, or perhaps two, to avoid creating excessively long names. (Names larger than the allowed length of a TSP name will be truncated). TAG=NONE stores under the original endogenous names.

The default values of the options are DYNAM, CONV2=.001, MAXPRT=5, METHOD=GAUSS, and no TAG. The print options are off except PRNSIM, so a one line summary of each iteration and a table of results will be printed.

Example

This example shows how to set up the well-known Klein Model I for simulation. At the end of this simulation, the solved variables are stored under the names CXS, IS, etc.

\[
\begin{align*}
\text{LIST } & Z \ C \ P(-1) \ K(-1) \ E(-1) \ TM \ W2 \ G \ TX ; \\
\text{2SLS (INST=Z) } & CX \ C \ P(-1) \ W ; \\
\text{FORM CONS} ; \\
\text{2SLS (INST=Z) } & I \ C \ P(-1) \ K(-1) ; \\
\text{FORM INV} ; \\
\text{2SLS (INST=Z) } & W1 \ C \ E(-1) \ TM ; \\
\text{FORM WAGES}; \\
\text{IDENT WAGE } & W = W1+W2 ; \\
\text{IDENT BALANCE } & E=E+CX+I+G-(TX+W+P) ; \\
\text{IDENT PPROD } & P = E-TX-W1 ; \\
\text{IDENT CAPSTK } & K=K(-1)+I ;
\end{align*}
\]
LIST KENDOG I W E P CS W1 K ;
LIST KLEIN CONS WAGES BALANCE PPROD INV WAGE CAPSTK ;
MODEL KLEIN KENDOG KLEINC ;
SOLVE (TAG=S,TOL=.0001,METHOD=FLPOW) KLEINC ;

References


SORT

Options  Examples

SORT sorts the observations of a series in increasing order. Other series or all series currently defined may also be reordered in the same order as the first series.

SORT (ALL, METHOD=BUBBLE/[QUICK], REVERSE) <key series> [<list of other series>] ;

or

SORT (AVETIES, METHOD=BUBBLE/[QUICK], MINTIES, RANK, REVERSE) <key series> <rank of series> ;

Usage

The simplest form is SORT followed by the name of a series to be sorted. A QuickSort is performed, so any equal values may be reordered relative to each other.

To sort several series in the same order as the first, just list them in the command. To sort all currently defined series, use the ALL option. All series must have the same length.

To sort series individually, use a DOT loop:

DOT X Y Z;
   SORT . ;
ENDDOT;

To "sort" in random order, draw a random variable and sort based on its values.

To obtain the rank order of a series use the second form of the command. The options AVETIES and MINTIES specify how tied series are to be treated.

Output

There is no printed output, but the series are stored after reordering their observations.

Options

ALL/NOALL causes all currently defined series to be sorted in the order defined by the named series.
**SORT**

**AVETIES/NOAVETIE** specifies that the average rank is to be stored for tied series. Used with RANK option.

**METHOD=[QUICK]/BUBBLE** specifies the sorting method to be used. The default QUICK is much faster than using a BUBBLE sort. However, if you wish to sort on multiple keys using a series of SORT commands, BUBBLE guarantees that the order of the initial sorts will be preserved.

**MINTIES/NOMINTIE** specifies that the minimum rank is to be stored for tied series. Used with RANK option.

**RANK/NORANK** stores the ordinal rank of the first series in the second series (the order of the first series is not changed).

**REVERSE/NOREVERS** sorts in decreasing order.

**Examples**

If \(X = 20 \ 40 \ 30 \ 50 \ 10\) and \(Y = 1 \ 2 \ 3 \ 4 \ 5\),

\[
\text{SORT (RANK) } X \ RX; \quad \text{? yields } RX = 2 \ 4 \ 3 \ 5 \ 1 \ (\text{and } X \text{ is unchanged})
\]
\[
\text{SORT } X Y; \quad \text{? yields } X = 10 \ 20 \ 30 \ 40 \ 50 \text{ and } Y = 5 \ 1 \ 3 \ 2 \ 4
\]
\[
\text{SORT (ALL) } X; \quad \text{? yields the same as the previous command}
\]
\[
\text{SORT (REVERSE) } X; \quad \text{? yields } X = 50 \ 40 \ 30 \ 20 \ 10
\]

If \(SCORE = 10 \ 20 \ 20 \ 40\),

\[
\text{SORT (RANK,AVETIES) } SCORE \ RA; \quad \text{? yields } RA=1.5 \ 2.5 \ 2.5 \ 4
\]
\[
\text{SORT (RANK,MINTIES) } SCORE \ RM; \quad \text{? yields } RM=1 \ 2 \ 2 \ 4
\]
STOP

Examples

STOP causes the TSP program to stop. If any variables are marked for storage on output databanks, they are written to the databank before the program stops.

STOP ;

Usage

Often, older TSP programs include a STOP statement at the end of the program section before the END statement that separates the TSP program section from the data section. This STOP statement is no longer required, since the END statement itself implies a STOP.

However, if you want to stop somewhere else in your TSP program, you can do this by using a STOP statement at any time. This can be convenient if you encounter an error and wish to abort the program. If you put a STOP statement at the beginning of your TSP program, TSP will check your whole program for syntax and then stop as soon as it reaches execution. This can be useful for debugging long programs.

Output

STOP produces no printed output. If output databanks have been used, variables are stored on them before stopping the program.

Examples

Here is an example of using STOP to check a TSP program for syntax:

STOP ;  ? Abort execution before doing anything
SMPL 1 1000 ;
....... long involved TSP program including complex equations, etc.
.......  
END ;

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STORE

Example  Reference

STORE writes microTSP-format (or EViews format) databank files.

STORE <list of series> ;
STORE <drive letter>:<series name> ......; (PC version only)

where drive letter is A, B, C, D, etc.

Usage

MicroTSP-format databank files may be useful for transferring data between microTSP and TSP. They are plain (editable, non-binary) files containing comments, frequency, starting and ending dates, and data values (one per line). See the microTSP documentation for details. They are not efficient in terms of disk space usage or the time required to read or write them. However, they are easy to edit, for manual data revision. TSP does not support the use of complex path names such as C:\foo\ser on the PC (if the data are in the same directory as your program, or in the working directory for interactive use, this should not be a problem). To move regular TSP databanks between machines, use the DBCOPY command. However, spreadsheet files are usually the easiest way to exchange small or medium-sized datasets between different programs.

When STORE writes to an existing file, it preserves any existing comments and always writes the current date. Only time series may be stored -- matrices, parameters, etc. can only be stored on regular TSP databanks.

The FETCH command reads files created by STORE.

Example

STORE X Y;

writes the series X and Y to the files X.DB and Y.DB.

Reference


http://www.eviews.com
SUPRES

Examples

SUPRES suppresses the printing of output results globally. It is an alias for REGOPT (NOPRINT):

SUPRES <list of output results> ;
REGOPT (NOPRINT) <list of output results> ;

NOSUPRES undoes any previous SUPRES that has been issued.

Usage

The arguments to SUPRES can be any of the output names beginning with @ described in this manual. The printing of the output associated with these names will be suppressed throughout the TSP program unless a NOSUPRES or REGOPT command with these codes is issued. The output results are still stored in memory and may be accessed.

Examples

To suppress all regression output, use the following command:

SUPRES REGOUT SMPL NOB COEF FST SBIC LOGL ;

This can be done more simply by using the SILENT option on an estimation (OLSQ, INST, etc.) command.

If you wish to see only a particular result, such as the Durbin-Watson, follow the SUPRES command with a NOSUPRES command:

NOSUPRES DW ;

To suppress the printing of the sample every time it changes, use

SUPRES SMPL ;

To cancel all previous SUPRES commands, use SUPRES with no arguments:

SUPRES ;
SUR

Options    References

SUR obtains seemingly unrelated regression estimates of a set of nonlinear equations. It is a special case of LSQ with the options set for SUR estimation. The LSQ command has a more complete description of the procedure.

\texttt{SUR (COVU=OWN or <covariance matrix of residuals>, MAXITW=0, HETERO, NOITERU, NOROBUST, nonlinear options) <list of equation names> ;}

Method

Seemingly unrelated regression estimates are obtained by first estimating a set of nonlinear equations with cross-equation constraints imposed, but with a diagonal covariance matrix of the disturbances across equations. These parameter estimates are used to form a consistent estimate of the covariance matrix of the disturbances, which is then used as a weighting matrix when the model is reestimated to obtain new values of the parameters. These estimates are consistent and asymptotically normal, and, under some conditions, asymptotically more efficient than the single equation estimates.

The seemingly unrelated regression method is a special case of generalized least squares with a residual covariance matrix of a particular structure:

\[ V = \Sigma \otimes I_T \]

where \( T \) = the number of observations and Sigma is the matrix of cross-equation variances and co-variances. It is sometimes called Zellner's method since it was originally proposed for linear models by Arnold Zellner, or the Aitken estimator (of which it is a special case).

Options

These are the same as for LSQ, except that NOITERU and MAXITW=0 are the defaults, in order to obtain SUR estimates.

References


SYMTAB

Example
SYMTAB prints the TSP symbol table, showing the characteristics of all the variables in a TSP program. It is useful primarily to programmers for debugging TSP programs. Others may prefer the SHOW command, which does not print out information on program variables.

SYMTAB ;

Usage
SYMTAB can be used anywhere in the program; it will print out the names, locations, types, lengths, and file pointers for all the variables used up to that point in the program. A description of the table is given in the output section below.

Output
The symbol table printout has 6 items for each variable:

1. **Variable name** - you will see all the variables you have created, as well as all the @ variables which contain results of procedures. In addition, there are a large number of variables which begin L 0001, or F 0001, and so forth. These variables are the TSP program lines and the equations which are created by the GENR and SET commands.

2. **Location** - this is the address of the variable in the upper end of blank common in single precision words.

3. **Type** - this is the variable type. Legal types are the following:

<table>
<thead>
<tr>
<th>type</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>scalar or constant</td>
</tr>
<tr>
<td>2</td>
<td>double precision (time) series (see OPTIONS DOUBLE)</td>
</tr>
<tr>
<td>3</td>
<td>(time) series</td>
</tr>
<tr>
<td>4</td>
<td>parameter</td>
</tr>
<tr>
<td>6</td>
<td>equation (output of FRML command)</td>
</tr>
<tr>
<td>7</td>
<td>identity (output of IDENT command)</td>
</tr>
<tr>
<td>8</td>
<td>model (output of MODEL command)</td>
</tr>
<tr>
<td>9</td>
<td>text string (for FILE=, FORMAT=, and TITLE)</td>
</tr>
<tr>
<td>10</td>
<td>program variable (a command, DO, IF, or PROC information)</td>
</tr>
</tbody>
</table>

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### SYMTAB

<table>
<thead>
<tr>
<th></th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>11</td>
<td>general matrix</td>
</tr>
<tr>
<td>12</td>
<td>symmetric matrix</td>
</tr>
<tr>
<td>13</td>
<td>triangular matrix</td>
</tr>
<tr>
<td>14</td>
<td>diagonal matrix</td>
</tr>
<tr>
<td>20</td>
<td>variable name list</td>
</tr>
</tbody>
</table>

4. **Length** - this is the length of the variable in single precision words. The length includes two extra items for time series and matrices which hold dating and dimension information.

5. **LDOC** - length of documentation, if any (see the DOC command).

6. **DB** - a flag for storage on the current OUT databank(s).

**Example**

Placement of the SYMTAB command at the end of the run will cause all the variables of the run to be printed out:

```
NAME USER ;
.
TSP program statements
.
SYMTAB ;  ? causes information on all variables used in the program to be displayed.
END ;
.
TSP data section
.
END ;
```
SYSTEM

SYSTEM provides interaction with the operating system without having to terminate the interactive session or batch job. If you are using a multitasking windowing system, such as Windows, you can do the same thing just by switching between windows.

**SYSTEM ['command string'];**

**Usage**

In batch mode, the SYSTEM command will execute the command you give in quotes. This may be useful for deleting files or running a program that processes files you have created with your TSP job. Note that you may have to explicitly CLOSE files before you can manipulate them with SYSTEM commands.

In interactive mode, SYSTEM usually takes no arguments, and simply produces the message

```
Enter system commands. Type EXIT or CONTINUE to resume TSP session.
$ (or some other system prompt)
```

when you have the system prompt, you may enter as many commands as you like.

**$ EXIT**

will resume your interactive session where you left off with no loss of continuity.

Each command you type is passed to the operating as a spawned subprocess. There are things that the subprocess will not be allowed to do, such as modify things about the environment within which the parent process (TSP) is executing.

.
TERMINAL (Interactive)

Redirect output that has previously been routed to an output file with the OUTPUT command to the terminal or screen.

TERMINAL ;

Usage

The TERMINAL command takes no arguments and performs a very simple operation: subsequent output is directed to the terminal. An important result of the TERMINAL command is that the previous file being used for output is closed. The output may not be examined (after using the SYSTEM command) until it has been closed. An output file may always be reopened, and output appended to it with another OUTPUT command and the same filename.
THEN

Example

THEN is part of the compound statement IF .... ; THEN ; .... ; ELSE ; ...... It comes before the statement (or group of statements surrounded by DO ; .... ; ENDDO ;) which are to be executed if the result of the expression on the IF statement is true.

THEN ;

Usage

THEN has no arguments; it is required as the next statement immediately following an IF statement. The statement immediately following the THEN statement will be executed if the result of IF is true. If you want more than one statement executed when the IF clause is true, enclose all the statements in a DO ; .... ENDDO ; group.

Example

IF LOGL>OLDL ; THEN ; DO ;
   SET OLDL = LOGL ;
   COPY @COEF SAVEB ;
ENDDO ;

See also the examples for the ELSE statement.
3SLS obtains three stage least squares estimates of a set of nonlinear equations. It is a special case of LSQ with the options set for 3SLS estimation. The LSQ entry has a more complete description of the command.

Three stage least squares estimates are consistent and asymptotically normal, and, under some conditions, asymptotically more efficient than single equation estimates. In general, 3SLS is asymptotically less efficient than FIML, unless the model is linear in the parameters and endogenous variables.

3SLS (COVU=OWN or <residual covariance matrix>, DEBUG, FEI, HETERO, INST=(<list of instrumental variables>), ITERU, MAXITW=0, ROBUST, nonlinear options) <list of equation names> ;

Usage

Three stage least squares is a combination of multivariate regression (SUR estimation) and two stage least squares. It obtains instrumental variable estimates, taking into account the covariances across equation disturbances as well. The objective function for three stage least squares is the sum of squared transformed fitted residuals.

Specification of the 3SLS command is the same as that of the LSQ command, except that the INST list is required. The variables in the INST list will be used to instrument all the equations, so that the actual instrumental variable matrix has the form given by Jorgenson and Laffont (1975), rather than that given by Amemiya (1977). In a simultaneous equations model, this means that a variable cannot be exogenous to one equation and endogenous to another. See the GMM command if you wish to relax this restriction.

Method

Three stage least squares estimates are obtained by first estimating a set of nonlinear (or linear) equations with cross-equation constraints imposed, but with a diagonal covariance matrix of the disturbances across equations. This is the constrained two stage least squares estimator. The parameter estimates thus obtained are used to form a consistent estimate of the covariance matrix of the disturbances, which is then used as a weighting matrix when the model is reestimated to obtain new values of the parameters.
The actual method of parameter estimation is the Gauss-Newton method for nonlinear least squares described under LSQ. If the model is linear in the parameters and endogenous variables, only two iterations will be required, one to obtain the covariance matrix estimate, and one to obtain parameter estimates.

For further details on the properties of the linear three stage least squares estimator see the Theil text or Zellner and Theil (1962). For the nonlinear three stage least squares estimator, see Amemiya (1977) and Jorgenson and Laffont (1975). The method of estimation in TSP is described more fully in Berndt, Hall, Hall, and Hausman (1975), also available at [this website](#).

Options

**COVU** = residual covariance matrix (same as the old **WNAME**= option below).

**DEBUG/NODEBUG** specifies whether detailed computations of the model and its derivatives are to be printed out at every iteration. This option produces extremely voluminous output and is not recommended for use except by systems programmers maintaining TSP.

**FEI/NFEI** specifies that models with additive individual fixed effects are to be estimated. The panel structure must have been defined previously with the **FREQ** (PANEL) command. The equations specified must be linear in the parameters (this will be checked) and variables. Individual-specific means will be removed from both variables and instruments.

**INST** = (list of instrumental variables). The list of instrumental variables supplied is used for all the equations. See the **INST** section of this manual and the references for further information on the choice of instruments.

**ITERU/NOITERU** specifies iteration on the **COVU** matrix; provides the same function as the old **MAXITW**= option.

**MAXITW** = the number of iterations to be performed on the parameters of the residual covariance matrix estimate. If **MAXITW** is zero the covariance matrix of the residuals is held fixed at the initial estimate (which is specified by **WNAME**). This option can be used to obtain estimates that are invariant to which equation is dropped in a shares model like translog.

**HETERO/NOHETERO** causes heteroskedastic-consistent standard errors to be used. See the **GMM** (NMA=) command for autocorrelation-consistent standard errors. Same as the old **ROBUST** option, or **HCOV=R**.

**WNAME** = the name of a matrix to be used as the starting value of the covariance matrix of the residuals.
**WNAME=OWN** specifies that the initial covariance matrix of the residuals is to be obtained from the residuals corresponding to the initial parameter values. If neither form of **WNAME=** is used, the initial covariance matrix is an identity matrix.

**Nonlinear options** control the iteration methods and printing. They are explained in the **NONLINEAR** section of this manual. Some of the common options are **MAXIT, MAXSQZ, PRINT/NOPRINT, and SILENT/NOSILENT**.

The only legal choice for **HITER=** is **G** (Gauss). **HCov=G** is the default method for calculating standard errors; **R** (Robust) is the only other valid option.

**Example**

Klein-I model:

```plaintext
FORM(VARPREF=C_) CONS CX C P(-1) W;
FORM(VARPREF=L_) INV I C P(-1) K(-1);
FORM(VARPREF=W_) WAGES W1 C E E(-1) TM;
3SLS(INST=(C,TM,W2,G,TX,P(-1),K(-1),E(-1))) CONS INV WAGES;
```

**References**


The TITLE statement is used to change the title on the top of the page of TSP printed output, or to print a centered title underlined with equals signs on the current page.

*TITLE (PAGE) 'Text string to be used as title';*

Usage

To change the TSP title, follow the word TITLE with up to sixty characters of text enclosed in single quotes.

If OPTIONS HARDCOPY; (the default for non interactive jobs) is in effect, a new page is started, with the new title printed at the top of the page. All further pages will use the new title until it is changed. If you do not want to start a new page, use the NOPAGE option. This option makes use of the PAGE command unnecessary.

If OPTIONS CRT; is in effect, the title is centered and printed underlined with stars on the current page (or screen).

Output

Under OPTIONS HARDCOPY, a new page is started with the new title at the top. Under OPTIONS CRT, the title is centered and printed underlined with equals signs (===).

Options

PAGE/NOPAGE tells whether a new page should be started, when OPTIONS HARDCOPY is in effect When OPTIONS CRT (the default) is being used, PAGE has no effect.

Examples

*TITLE 'Results for small firms';
SELECT SALES<10;
DOT X Y Z;
TITLE . ; ? prints variable names from DOT as titles.
OLSQ . C R ; LAD . C R ;
ENDDOT ;
TOBIT obtains estimates of the linear Tobit model, where the dependent variable is either zero or positive. The method used is maximum likelihood under the assumption of homoskedastic normal disturbances. For non-normal censored regression, see LAD.

\[
\text{TOBIT (LOWER=lower limit, MILLS=\text{name for output inverse Mills ratio}, UPPER=upper limit, WEIGHT=\text{weighting series}, <nonlinear options>) <dependent variable> <list of independent variables> ;}
\]

Usage

The basic TOBIT statement is like the PROBIT or OLSQ statements: first list the dependent variable and then the independent variables. If you wish to have an intercept term in the regression (usually recommended), include the special variable C or CONSTANT in your list of independent variables. You may have as many independent variables as you like subject to the overall limits on the number of arguments per statement and the amount of working space, but of course the number is limited by the number of data observations you have available.

The observations over which the regression is computed are determined by the current sample. If any of the observations have missing values within the current sample, TOBIT will print a warning message and will drop those observations.

The list of independent variables on the TOBIT command may include variables with explicit lags and leads as well as PDL (Polynomial Distributed Lag) variables. These distributed lag variables are a way to reduce the number of free coefficients when you are entering a large number of lagged variables in a regression by imposing smoothness on the coefficients. See PDL for a description of how to specify such variables.

The dependent variable need not be a strictly zero/positive variable. Negative values are treated as zero. The standard Tobit model involves truncation of the dependent variable below zero. Models with upper and/or lower truncation can be estimated by using the UPPER and/or LOWER option(s). See the Examples for more details.

Output
The output of TOBIT begins with an equation title and the name of the dependent variable. Then the starting values and diagnostic output from the iterations are printed, followed by the convergence status.

The results printed are the mean of the dependent variable, the number of lower censored, uncensored, and upper censored observations, and a table of right hand side variable names, estimated coefficients, standard errors and associated t-statistics. The estimated standard deviation of the residual, SIGMA, is listed last in this table.

TOBIT also stores some of these results in data storage for your later use. The table below lists the results available after a TOBIT command.

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@LHV</td>
<td>list</td>
<td>1</td>
<td>Name of dependent variable</td>
</tr>
<tr>
<td>@YMEAN</td>
<td>scalar</td>
<td>1</td>
<td>Fraction of positive observations</td>
</tr>
<tr>
<td>@NOB</td>
<td>scalar</td>
<td>1</td>
<td>Number of observations</td>
</tr>
<tr>
<td>@NPOS</td>
<td>scalar</td>
<td>1</td>
<td>Number of positive observations</td>
</tr>
<tr>
<td>@LOGL</td>
<td>scalar</td>
<td>1</td>
<td>Log of likelihood function</td>
</tr>
<tr>
<td>@IFCONV</td>
<td>scalar</td>
<td>1</td>
<td>1 if convergence achieved, 0 otherwise</td>
</tr>
<tr>
<td>@NCOEF</td>
<td>scalar</td>
<td>1</td>
<td>Number of parameters (#params) including SIGMA</td>
</tr>
<tr>
<td>@NCID</td>
<td>scalar</td>
<td>1</td>
<td>Number of identified coefficients</td>
</tr>
<tr>
<td>@RNMS</td>
<td>list</td>
<td>#params</td>
<td>list of names of independent variables</td>
</tr>
<tr>
<td>@COEF</td>
<td>vector</td>
<td>#params</td>
<td>Coefficient estimates</td>
</tr>
<tr>
<td>@SES</td>
<td>vector</td>
<td>#params</td>
<td>Standard errors</td>
</tr>
<tr>
<td>@T</td>
<td>vector</td>
<td>#params</td>
<td>T-statistics</td>
</tr>
<tr>
<td>%T</td>
<td>vector</td>
<td>#params</td>
<td>p-values for T-statistics</td>
</tr>
<tr>
<td>@GRAD</td>
<td>vector</td>
<td>#params</td>
<td>Gradient of log likelihood at convergence</td>
</tr>
<tr>
<td>@VCOV</td>
<td>matrix</td>
<td>#params*#params</td>
<td>Variance-covariance of estimated coefficients</td>
</tr>
<tr>
<td>@DPDX</td>
<td>matrix</td>
<td>#vars*2</td>
<td>Means of probability derivatives</td>
</tr>
<tr>
<td>@RES</td>
<td>series</td>
<td>#obs</td>
<td>Residuals for non-truncated observations</td>
</tr>
<tr>
<td>@MILLS</td>
<td>series</td>
<td>#obs</td>
<td>Inverse Mills' ratios</td>
</tr>
</tbody>
</table>

If the regression includes a PDL variable, the following will also be stored:

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@SLAG</td>
<td>scalar</td>
<td>1</td>
<td>Sum of the lag coefficients</td>
</tr>
<tr>
<td>@MLAG</td>
<td>scalar</td>
<td>1</td>
<td>Mean lag coefficient (number of time periods)</td>
</tr>
<tr>
<td>@LAGF</td>
<td>vector</td>
<td>#lags</td>
<td>Estimated lag coefficients, after &quot;unscrambling&quot;</td>
</tr>
</tbody>
</table>
Method

TOBIT uses analytic first and second derivatives to obtain maximum likelihood estimates via the Newton-Raphson algorithm. This algorithm usually converges fairly quickly. Starting values for the parameters are obtained from a regression on the observations with positive values of the dependent variable. See Greene (1981), p. 508, formula (13) and footnote 5 for the details. Alternative starting values may be supplied in @START (see NONLINEAR). A globally concave parameterization of the likelihood function is used for iterations. Multicollinearity of the independent variables is handled with generalized inverses, as in all TSP regression procedures.

The numerical implementation involves evaluating the normal density and cumulative normal distribution functions. The cumulative normal distribution function is computed from an asymptotic expansion, since it has no closed form. See the references under CDF for the actual method used to evaluate CNORM(). The ratio of the density to the distribution function is also known as the inverse Mills ratio. This is used in the derivatives and with the MILLS= option.

Options

LOWER= the value below which the dependent variable is not observed. The default is zero.

MILLS= the name of a series used to store the inverse Mills ratio series evaluated at the estimated parameters. The default is @MILLS.

WEIGHT= the name of a weighting series. The weights are applied directly to the likelihood function, and no normalization is performed.

UPPER= the value above which the dependent variable is not observed. The default is no limit.

Nonlinear options - see NONLINEAR.

Examples

Standard Tobit model with truncation below zero:

**TOBIT CAR C INCOME RURAL MSTAT;**

Truncation below two:

**TOBIT (LOWER=2) CAR C INCOME RURAL MSTAT;**

Truncation above ten:
**TOBIT**

**TOBIT (UPPER=10) CAR C INCOME RURAL MSTAT;**

**References**


TREND

Options  Examples

TREND generates a series with a linear growth trend. The trend may be repeated under the control of several options.

TREND (FREQ, PERIOD=<value>, PSTART=<value>, START=<initial value>, STEP=<increment>) <series> ;

or

TREND <series> [<initial value>] [<increment>] ;

Usage

TREND followed by a series name makes a simple time trend variable and stores it under that name. This variable is equal to one in the beginning observation of the current sample and increases by one in every period.

The starting value of the series may be changed by including the second argument and the increment may also be changed by including the third argument. If you want to change the increment, but not the starting value, use the STEP option.

Since the TREND command creates a series under control of SMPL, you must be careful to specify a SMPL which covers the whole period in which you are interested, so you don't have missing values or strange jumps due to gaps in the sample.

Output

TREND produces no printed output. A single series is stored.

Options

FREQ/NOFREQ causes the trend to be restarted every time there is a new year. This option is valid only when FREQ Q; or FREQ M; has been specified.

PERIOD= specifies the number of observations after which the trend starts repeating itself. For example, PERIOD=4 would have the same effect as FREQ Q.

PSTART= specifies the starting period (for the first observation in the sample), when PERIOD is used. The default is one. For example,

FREQ Q; SMPL 70:2,79:4 ;
TREND (FREQ) QT ;
TREND

is equivalent to

SMPL 2,40;
TREND (PERIOD=4,PSTART=2) QT;

START= gives an initial value to the trend. The default is one.

STEP= supplies a value for the trend increment. The default is one.

Examples

FREQ A ; SMPL 46 75;
TREND TIME;

The above example creates a time trend variable called TIME for the illustrative model. The variable is 1 in 1946, 2 in 1947, 3 in 1948, and so forth.

FREQ Q ; SMPL 70:2,79:4 ;
TREND (FREQ) QT;

This example makes a series QT equal to 2,3,4, 1,2,3,4, 1,2,3,4,...etc.

SMPL 1 400;
TREND (PERIOD=5) TIME;

This example makes a series which is 1,2,3,4,5 starting in every fifth observation; this might be useful for panel data, where a trend was needed.

SMPL 1,400;
TREND (PERIOD=5,START=71) YEAR;

This example is the same as above, except the trend is equal to 71,72,73,74,75, 71,72,73,74,75, ... instead of 1,2,3,4,5, 1,2,3,4,5, ...
TSTATS

Output  Option  Example

TSTATS prints a table of names, coefficients, standard errors, t-statistics, and (if it is not suppressed using REGOPT) a variance-covariance matrix. TSTATS is useful when you compute your own estimate of a variance-covariance matrix, since it is tedious to take the square roots of the diagonal elements and generate a readable printout of the results.

\[ \text{TSTATS (NAMES=(<list of names>)) <coefficient vector> <variance-covariance matrix> ;} \]

Usage

The simplest form is TSTATS followed by the name of a vector containing the estimated values of a set of coefficients and the name of a symmetric matrix which contains the estimated covariance matrix of those coefficients. If the vector of coefficients is \( N \) long, the matrix must be of order \( N \). The NAMES option allows you to label the coefficients in the table conveniently.

Output

A table of regression coefficients, etc. is printed unless it has been SUPRESed. @RNMS, @COEF, @SES, and @VCOV are stored.

Option

NAMES= <list of coefficient names>. The default is just to number the coefficients 1,2, etc.

Example

Suppose, for example, that we have manually created PDL variables for use in a nonlinear regression, and then unscrambled the regression coefficients and their covariance matrix (by using the PDL transformation matrix). The estimated lag coefficients are called BETA and their covariance matrix estimate is VARB. A table of estimates with the corresponding t-statistics is printed by implementing the following command:

\[ \text{TSTATS(NAMES=(BETA1-BETA7)) BETA VARB ;} \]
UNIT

UNIT is a synonym for COINT, which performs unit root and cointegration tests.

UNIT (ALL, NOCOINT, CONST, DF, FINITE, MAXLAG=<number of lags>, MINLAG=<number of lags>, PP, RULE=AIC2, SEAS, SEAST, SEASTSQ, SILENT, TREND, TSQ, UNIT, WS) <list of variables> [ | <list of special exogenous trend variables> ] ;
UNMAKE

Examples
UNMAKE takes a matrix and splits it column by column into a set of series. The matrix must have a number of rows equal to the number of observations in the current sample and a number of columns equal to the number of series whose names are supplied. Similarly, UNMAKE will split a vector into a set of scalars, if the number of scalars is equal to the length of the vector. UNMAKE is the reverse of **MMAKE** which makes a matrix from series, or a vector from scalars.

\[ \text{UNMAKE \ matrix} \ <\text{list of series}>; \]
\[ \text{or} \]
\[ \text{UNMAKE \ vector} \ <\text{list of scalars}>; \]

Usage
UNMAKE's first argument is the name of the matrix to be broken up; the second a list of the names of the series where the columns of the matrix will be put. The number of series is limited only by the maximum size of the argument list (usually about 100 or more) and the space available in data storage for the new matrix.

The series named will be replaced if they already exist; if the **NOREPL** option is being used, observations outside the current sample will be deleted. The new series have the frequency and starting observation of the current sample.

The matrix to be unmade may be of any type; it will be expanded before UNMAKE is executed. An exception to this rule is in the case of a diagonal matrix -- if the length of the current sample is equal to the length of the diagonal, and only one series name is supplied, the diagonal of the matrix will be stored in this series.

UNMAKE can also be used to split a vector into scalars; this is useful for rearranging coefficient vectors and setting up starting value vectors. This is easier than specifying several **SET** statements or a tricky **DOT** loop.

Output
UNMAKE produces no printed output. A set of series (or scalars) are stored in data storage.
Examples

If the current sample is SMPL 1 5, and there is a 5 by 2 matrix X with the following elements:

<p>| | |</p>
<table>
<thead>
<tr>
<th></th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>9</td>
</tr>
<tr>
<td>2</td>
<td>8</td>
</tr>
<tr>
<td>3</td>
<td>7</td>
</tr>
<tr>
<td>4</td>
<td>6</td>
</tr>
<tr>
<td>5</td>
<td>5</td>
</tr>
</tbody>
</table>

the command

`UNMAKE X X1 X2;`

results in the following two series being stored:

<table>
<thead>
<tr>
<th>X1</th>
<th>X2</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>9</td>
</tr>
<tr>
<td>2</td>
<td>8</td>
</tr>
<tr>
<td>3</td>
<td>7</td>
</tr>
<tr>
<td>4</td>
<td>6</td>
</tr>
<tr>
<td>5</td>
<td>5</td>
</tr>
</tbody>
</table>

Any submatrix within a matrix can also be obtained by unmaking the matrix under one SMPL and remaking it under another. For example, given an 8 by 8 covariance matrix, a new covariance matrix that contains only the elements corresponding to the 3rd, 4th and 5th variables can be extracted by the following commands:

`SMPL 1 8;`  
`UNMAKE VAR COL1-COL8;`  
`SMPL 3 5;`  
`MMAKE VAR35 COL3-COL5;`

Here is an example of UNMAKE with a vector:

`OLSQ Y C X1-X6;`  
`UNMAKE @COEF B0-B6;`

See also the example under `MMAKE` for manipulating a vector of starting values for parameters.
UPDATE (Interactive)

Update allows you to specify portions of a series, or list of series whose observations you wish to modify. It is for interactive use only.

UPDATE <list of series> ;

Usage

UPDATE is a special form of the ENTER command. The only difference is that UPDATE is expecting to modify an existing series. You will be prompted to specify the observations you wish to update -- your response must conform to the rules for specifying a SMPL. A single number will be interpreted as a request to update a single observation. Prompting and storage will be defined by the SMPL you specify, and the most recent FREQ prior to the UPDATE command. If the frequency is not appropriate for the series, you will encounter errors.

In response to the prompt for data, you may enter as many items per line as you like -- the prompts will adjust accordingly. Prompting will cease when the observations you specified have been updated, and the series will be restored. If more than one series is being updated, you will be prompted to update them sequentially.
**USER**

**USER (Mainframe)**

**Reference**

USER allows a user-written TSP subroutine to be linked to the program. The list of arguments you have written is passed directly to the subroutine USER.

```
USER <list of arguments> ;
```

**Usage**

To use the USER feature, you will have to write your own subroutine in Fortran and link it to the TSP program. To do this, you should consult the *TSP Programmer's Guide* and your local TSP consultant. Attempting to write a TSP subroutine requires understanding how the program works internally and is not recommended for novices or those who are unfamiliar with Fortran.

A default USER subroutine comes with the program; currently it takes the generalized inverse of a symmetric matrix and stores the corresponding eigenvalues and eigenvectors. Note that this feature requires that you have the source code version of TSP (usually not available for PC or Mac, but sometimes available on request for unix).

**Output**

The USER procedure will produce whatever output you print or store in data storage.

**Reference**

VAR

Output  Options  Example  Reference

VAR performs vector autoregressions, which are a set of unrestricted "reduced form" linear regressions with lags of the dependent variables on the right hand side. Impulse response functions (dynamic simulations based on the estimated coefficients), forecast error decompositions, and block exogeneity tests are also performed.

VAR (BOUNDS=<sign. level of confidence intervals>,NBOOT=<num of bootstrap replications>,NHORIZ=<length of impulse response>,NLAGS=<number of lags in VAR>, SHOCK=ALL or CHOL or GEN or STDDEV or UNIT or <matrix name>, SILENT, TERSE) <list of dependent variables> [ | <list of exogenous variables> ] ;

Usage

First list the dependent variables, and specify the number of lags desired in the options list. If there are any exogenous variables, give their names after a |. If you want to have intercept terms in the regressions (usually recommended), include the special variable C or CONSTANT in the list of exogenous variables. You may have as many independent variables as you like subject to the overall limits on the number of arguments per statement and the amount of working space; obviously, the number is limited by the number of data observations available. The observations over which the regression is computed are determined by the current sample. If any observations have missing values within the current sample, VAR drops the missing observations and prints an error message.

Output

VAR output begins with an equation title and the names of the dependent variables, followed by the log likelihood value and a table of the regression coefficients. Various statistics on goodness-of-fit are printed for each equation: the sum of squared residuals, the standard error of the regression, the R-squared, and the Durbin-Watson statistic for autocorrelation of the residuals (biased unless NLAGS=0 because of the presence of lagged dependent variables). Block exogeneity tests (Granger causality tests) are computed as F tests to see if lagged values of other dependent variables are significant in each equation. Next are the impulse response functions (see Method) and variance decompositions.
The command PLOTS (PREVIEW) IR; can be used in TSP/Oxmetrics to automatically display color plots of the impulse responses for all subsequent VAR commands, even when running in batch mode. For NEND endogenous variables, there are NEND such plots per window - responses of a given endogenous variable to shocks in all variables. There will also be NEND VAR plot windows created, labelled VAR, VAR2, etc.

The variance decomposition is for residual variances only (it does not include sampling error in the regression coefficients). VAR stores most of these results in data storage for later use. Here are the results available after a VAR command:

<table>
<thead>
<tr>
<th>variable</th>
<th>type</th>
<th>length</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>@LHV</td>
<td>list</td>
<td>#eqs</td>
<td>Name of the dependent variable</td>
</tr>
<tr>
<td>@RNMS</td>
<td>list</td>
<td>#eqs*#vars</td>
<td>list of names of right hand side variables</td>
</tr>
<tr>
<td>@LOGL</td>
<td>scalar</td>
<td></td>
<td>Log of likelihood function</td>
</tr>
<tr>
<td>@SBIC</td>
<td>scalar</td>
<td></td>
<td>Schwarz Bayesian Information Criterion</td>
</tr>
<tr>
<td>@AIC</td>
<td>scalar</td>
<td></td>
<td>Akaike Information Criterion</td>
</tr>
<tr>
<td>@SSR</td>
<td>vector</td>
<td>#eqs</td>
<td>Sum of squared residuals</td>
</tr>
<tr>
<td>@S</td>
<td>vector</td>
<td>#eqs</td>
<td>Standard error of the regression</td>
</tr>
<tr>
<td>@YMEAN</td>
<td>vector</td>
<td>#eqs</td>
<td>Mean of the dependent variable</td>
</tr>
<tr>
<td>@SDEV</td>
<td>vector</td>
<td>#eqs</td>
<td>Standard deviation of the dependent variables</td>
</tr>
<tr>
<td>@DW</td>
<td>vector</td>
<td>#eqs</td>
<td>Durbin-Watson statistic</td>
</tr>
<tr>
<td>@RSQ</td>
<td>vector</td>
<td>#eqs</td>
<td>R-squared</td>
</tr>
<tr>
<td>@ARSQ</td>
<td>vector</td>
<td>#eqs</td>
<td>Adjusted R-squared</td>
</tr>
<tr>
<td>@FBEX</td>
<td>vector</td>
<td>#eqs</td>
<td>F-statistics for block exogeneity</td>
</tr>
<tr>
<td>@COEF</td>
<td>vector</td>
<td>#eqs*#vars</td>
<td>Coefficient estimates</td>
</tr>
<tr>
<td>@SES</td>
<td>vector</td>
<td>#eqs*#vars</td>
<td>Standard Errors</td>
</tr>
<tr>
<td>@VCOV</td>
<td>matrix</td>
<td>(#eqs*#vars)**2</td>
<td>Variance-covariance of estimated coefficients.</td>
</tr>
<tr>
<td>@COVU</td>
<td>matrix</td>
<td>#eqs*#eqs</td>
<td>Residual covariance matrix = E'Ε/(T-K).</td>
</tr>
<tr>
<td>@IMPRES</td>
<td>matrix</td>
<td>nhoriz*#eqs**2</td>
<td>Impulse Response function.</td>
</tr>
<tr>
<td>@FEVD</td>
<td>matrix</td>
<td>nhoriz*(1+#eqs)**eqs</td>
<td>Forecast error variance decomposition</td>
</tr>
<tr>
<td>@RES</td>
<td>matrix</td>
<td>#obs*#eqs</td>
<td>Residuals</td>
</tr>
</tbody>
</table>
Method

OLS is performed, equation by equation. This is efficient because the right hand side variables are identical for every equation. The impulse response function is just a dynamic simulation with shocks in the first period for some variables and zeros for the other variables.

Options

**BOUNDS=** specifies the significance level of the confidence intervals. The default is .95, for 95%.

**NBOOT=** specifies the number of bootstrap replications used in computing confidence intervals for the impulse response functions. The default is 200. A nonparametric bootstrap is used - NBOOT resamples are taken from the rows of the original data matrix. The regression coefficients, residual covariance matrix, shock matrix, and impulse response functions are computed for each resample.

**NHORIZ=** number of time periods for the impulse response function (default 10). Specify NHORIZ=0 or SILENT to suppress the impulse response output.

**NLAGS=** number of lags of the dependent variables to include on the right hand side of the equations (default zero). @SBIC or @AIC can be used to choose the number of lags (minimize @SBIC -- see the REGOPT command).

**SHOCK=** **ALL** or **CHOL** or **GEN** or **STDDEV** or **UNIT** or **matrix name** specifies the type of shock for the impulse response function. CHOL, the default, is a Choleski factorization (matrix square root) using the current ordering of the dependent variables. A shock to the first factor affects the first variable initially, while a shock to the second factor affects the first two variables, etc. ALL computes Choleski factorizations for all (n!) orderings of the variables; since different orderings can produce markedly different results, this option is useful for making sure the results are robust to ordering. You can also supply your own (square) matrix factorization. STDDEV and UNIT specify (residual) standard deviation or unit shocks to single variables; variance decompositions are not computed for these shocks.
Shocks specified that a Generalized Impulse Response should be computed, which is invariant to the ordering of the equations. See the Pesaran and Shin (1998) reference. For each dependent variable, the impulse responses and confidence intervals are printed in a table. Labels are of the form Y_X. That is, if the column is labelled GNP_M1, then it is the impulse response of GNP to a shock in M1. The printing of the table can be suppressed by using SUPRES @IMPRES; or the option SILENT in the VAR command.

The columns are stored as series of length NHORIZ, starting in the first observation of the estimation sample. In the example above, the series names would be @IRGNP_M1, @IRGNP_M1LB95%, and @IRGNP_M1UB95%.

SILENT/NOSILENT suppresses all printed output. This is useful for running regressions for which you only want selected output (which can be obtained from the @ variables, which will be stored).

TERSE/NOTERSE causes minimal output (results only) to be printed.

Example

```
VAR(NLAGS=5) Y1 Y2 Y3 Y4 | C X1 X2 X3;
```

is equivalent to the following regressions:

```
OLSQ Y1 Y1(-1)-Y1(-5) Y2(-1)-Y2(-5) Y3(-1)-Y3(-5) Y4(-1)-Y4(-5) C X1-X3;
OLSQ Y2 Y1(-1)-Y1(-5) Y2(-1)-Y2(-5) Y3(-1)-Y3(-5) Y4(-1)-Y4(-5) C X1-X3;
OLSQ Y3 Y1(-1)-Y1(-5) Y2(-1)-Y2(-5) Y3(-1)-Y3(-5) Y4(-1)-Y4(-5) C X1-X3;
OLSQ Y4 Y1(-1)-Y1(-5) Y2(-1)-Y2(-5) Y3(-1)-Y3(-5) Y4(-1)-Y4(-5) C X1-X3;
```

Example of VAR with generalized impulse response functions, showing how to plot the results by hand (TSP/Oxmetrics will do this automatically - see above).

```
VAR (NLAG=MAXLAG,SHOCK=GEN,NHORIZ=HORIZ) Y CONS IN | C T ;
SELECT .NOT.MISS(@IRGNP_M1) ;
PLOT @IRGNP_M1 @IRGNP_M1LB95% @IRGNP_M1UB95% ;
```

See the TSP User's Guide for more examples.

Reference

WRITE

WRITE

Options  Examples

WRITE is used to write variables to the screen, output file, or an external file. The output may be labelled, free format, or a format of your specification. WRITE can create Lotus, Excel, and binary files. PRINT is synonymous with WRITE.

WRITE (COLS=<list of column names>,FILE='filename string', FORMAT=BINARY or DATABANK or EXCEL or FREE or LABELS or LOTUS or RB4 or RB8 or (format text string), FULL, ROWS=<list of rownames>, UNIT=<I/O unit number> <list of variables> or <matrixname>;

Usage

WRITE is the inverse of the READ statement: series or other variables which are read with a particular READ statement may be written by a WRITE statement of the same form.

When the list of variables contains only series, WRITE writes one record for each observation in the current sample (unless the format statement specifies more than one record); this record contains the value of all the series listed on the statement for that observation. If only one series is listed, WRITE writes at least one observation per record.

The first WRITE command to a file creates a new file or overwrites an existing file. Subsequent WRITEs to the same text file (FREE or formatted), in the same batch run will append to the file if the file has remained open (this is usually the case unless you have a lot of open files). Subsequent writes to the same spreadsheet file (EXCEL or LOTUS) overwrite the existing file.

If the list of variables on the WRITE statement includes some which are not series, the variables are written to the file one at a time, with one record per variable. Symmetric, triangular, and diagonal matrices are written in compressed storage mode unless the FULL option is specified. You are responsible for making sure your format allows for enough data points if you use the FORMAT='format string' option. Subscripted matrices are treated like scalars.

In labelled or FREE format: series, scalars, and matrices, missing values are shown as the character . (period) and a warning is printed. FREE format values are written to all significant digits and each observation starts on a new line. Minimal spacing is used to produce as compact output as possible.

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WRITE

A single WRITE command will write either a single matrix or several series to a spreadsheet file. If the file specified already exists, it is overwritten by the new file. If you write a matrix, the file will consist only of numbers. If you write series, their names will be put in the first row. The first column will contain dates (or observation numbers), and each series will be put in a column below its name. Series are written under the control of the current sample. If there are gaps in the sample, observation numbers will be used instead of dates in the first column. Dates are written as the last day of each period, and formatted as Month-Year.

Output

WRITE produces printed output in the output file or screen, unless the FILE= option is used, in which case data are written to an external data file.

Options

COLS= list of column names used to label a matrix when it is written (useful for printed output).

FILE= 'filename string' specifies the actual name of the file where the data is to be written.

FORMAT=BINARY or DATABANK or EXCEL or FREE or LABELS or LOTUS or RB4 or RB8 or '(format text string)' specifies the format in which the data is to be written or printed. The default is LABELS unless the FILE option is also specified, in which case the default is FREE. Each format option is described below.

FORMAT=BINARY specifies that the data is to be written in single precision (REAL*4) format on the external file. This format for data is by far the most efficient if you do not plan to move the data to another computer and should be used, if possible, if you have a large amount of data. To read such a data file, use the READ command with the FORMAT=BINARY option. This format cannot be used for equations.

FORMAT=DATABANK specifies that the data are to be written to a TSP databank. This option requires the FILE= option also.

FORMAT=EXCEL writes an Excel spreadsheet file (similar to FORMAT = LOTUS). If the filename ends .XLS, this is the default. This option requires the FILE= option also.
WRITE

FORMAT=FREE specifies that the data is to be written to an external file in a format determined by TSP. This option causes the data to be represented by six numbers per record with a field width of 15 characters and at least 7 significant digits. The exact format is chosen by the program to represent the numbers most conveniently.

FORMAT=LABELS specifies that the data is to be formatted as for printed output, with observation labels if they are series, and row and column numbers and titles if they are vectors or matrices. This option is the default if the output is being written to the output file or screen, or if the item being written is an equation.

FORMAT=LOTUS writes a Lotus 123 .WKx worksheet file. The variable names are written in the first row atop the series columns. If a FREQ other than N is in effect, dates are written in the leftmost column. FORMAT=LOTUS is the default if the filename string includes .WK. (See the READ command.)

FORMAT=RB4 is the same as FORMAT=BINARY (single precision binary)

FORMAT=RB8 is used for double precision binary.

FORMAT=’(format text string)’ specifies a fixed format with which the data is to be written. The quotes are required and should surround a Fortran FORMAT statement, including the parentheses but excluding the word FORMAT. If you are unfamiliar with the construction of a Fortran FORMAT statement, see FORMAT.

FULL/NOFULL specifies whether symmetric, triangular, and diagonal matrices are to be expanded before being written out. This option is ignored when the FORMAT=DATABANK or FORMAT=LABELS are used.

ROWS= list of row names used to label the rows of a matrix when it is printed (useful for printed output).

UNIT= an integer number (usually between 1 and 4, or 8 and 99) which is the Fortran input/output number of an external file from which the variables listed will be written. This is rarely used.

Examples

This example is the inverse of the example for the READ command.

WRITE (FILE=’FOO.DAT’) X Y Z ;

To look at some transformed series:

PRINT X,LX,DX ;

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The following example creates a spreadsheet file that can be read by the corresponding READ command examples:

```
FREQ Q;
SMPL 48:1,49:1;
READ CJMTL; 183.4 185.2 192.1 193.3 206.9; )
READ PMTL; .436 .562 .507 .603;
WRITE (FILE='SML.WKS') CJMTL,PMTL;
```

The next example creates a spreadsheet file with the same data columns, but no dates or series names (since a matrix is used).

```
MMAKE M CJMTL,PMTL;
WRITE (FILE='SMM.WKS') M;
```

The SMM.WKS file that is created:

```
A   B
1 183.4 NA
2 185.2 .436
3 192.1 .562
4 193.3 .507
5 206.9 .603
```

Alternatively, you could print M with the rows and columns labelled:

```
       COLS=(CJMTL,PMTL)) M ;
```

```
CJMTL PMTL
YR1996 183.4 NA
YR1997 185.2 .436
YR1998 192.1 .562
YR1999 193.3 .507
YR2000 206.9 .603
```
YLDFAC

Example References

YLDFAC factors a symmetric matrix $X$ into a triangular matrix $L'$ and a diagonal matrix $D$ such that $X=LDL'$ and the diagonals of $D$ are functions of the characteristic roots of $X$.

\[ YLDFAC \text{<symmetric matrix>} \text{<diagonal matrix>} \text{<triangular matrix>} ; \]

Usage

Three required arguments to YLDFAC are: name given to the matrix to be factored (must be symmetric or an error message will be printed); name given to the diagonal matrix; and name given to the upper triangular matrix. Most symmetric matrices can be factored in this way. The elements of $D$ are functions of the characteristic roots of the input matrix. If all the diagonal elements of $D$ are positive, the input matrix is positive definite. If all of them are nonnegative, the input matrix is positive semi-definite. If there are some positive elements of $D$ and some negative, the input matrix is indefinite. Zero elements on the diagonal of $D$ imply that the input matrix is singular or near singular. The diagonal elements of $L$ are normalized to 1.

Output

YLDFAC produces no printed output. Two matrices are stored in data storage.

Method

A modified Choleski method of factorization is used where the diagonal element is extracted from the square root matrix as the factorization is performed. The underlying method is described in the Faddeev reference.

Example

\[ YLDFAC \text{A DIAG UPPER ;} \]
\[ \text{MAT ANEW = UPPER'DIAG*UPPER ;} \]

generates a matrix ANEW which is identical to the original matrix A. Note that because the triangular matrix is stored as an upper triangle, it must be transposed to obtain $L$.

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