

Numerical linear algebra and optimization tools for bioinformatics

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Abstract

Computational models often require the solution of **large systems of linear equations $Ax = b$** or **least-squares problems $Ax \approx b$** or more challenging **optimization problems involving large sparse matrices**.

For example, the modeling of **biochemical reaction networks** in systems biology may depend on determining the **rank of large stoichiometric matrices**, and on accurate solution of **large multiscale linear programs**, as in Flux Balance Analysis (**FBA**) and Flux Variability Analysis (**FVA**). A thermodynamically feasible set of fluxes can be obtained by solving a similar **large optimization problem** that has a **negative entropy objective function**.

We describe some **general-purpose algorithms and software** that have provided efficient and reliable solutions for important problems in systems biology, and are likely to find broader application.

- 1 SOL
- 2 Sparse $Ax \approx b$
- 3 Stoichiometric matrices
- 4 Rank of stoichiometric matrices
- 5 SQOPT, SNOPT
- 6 PDCO
- 7 Conclusions

SOL

Systems Optimization Laboratory

Stanford University

SOL

- Founded 1974 by George Dantzig and Richard Cottle
- Dantzig, Alan Manne: economic models (linear & nonlinear)
- Gill, Murray, Saunders, Wright: Software for optimization

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Recent collaborators:

- **Philip Gill (UC San Diego)**
Optimization software NPSOL, QPOPT, SQOPT, SNOPT
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Flux balance analysis (FBA), Flux variability analysis (FVA)
Rank and nullspace of stoichiometric matrices
Nonequilibrium fluxes in metabolic networks
- **Bernhard Palsson (UCSD)**
FBA and FVA

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Funding: ONR, AFOSR, ARO, DOE, NSF, AHPCRC, . . . ,
DOE DE-FG02-09ER25917, NIH U01-GM102098

Sparse linear equations $Ax = b$ and least squares problems $Ax \approx b$

Problem types and software packages

Iterative methods for $Ax \approx b$

A may be a sparse matrix
or an operator for computing Av and/or $A^T w$

A may have any rank

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- Symmetric $Ax = b$

$$A = \square$$

CG, MINRES-QLP

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- Symmetric $Ax = b$

$$A = \square$$

CG, MINRES-QLP

- $\min \|Ax - b\|$

$$A = \square \text{ or } \square \text{ or } \square$$

LSMR

Iterative methods for $Ax \approx b$

A may be a sparse matrix
or an operator for computing Av and/or $A^T w$

A may have any rank

- Symmetric $Ax = b$ $A = \square$ CG, MINRES-QLP
- $\min \|Ax - b\|$ $A = \square$ or \square or \square LSMR
- Tall skinny $\min \|Ax - b\|$ $A = \square$ LSRN

Sparse direct methods for $Ax = b$

- $A = LDU$ LUSOL (Stanford)
- $A = QR$ SPQR (Tim Davis, UFL)
- Many sparse solvers HSL Library (RAL, UK)

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Dense factorizations

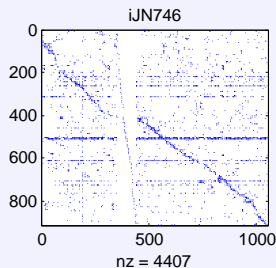
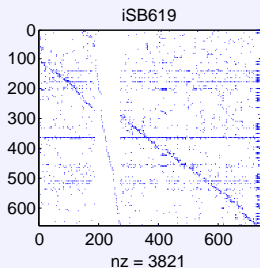
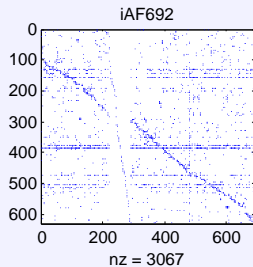
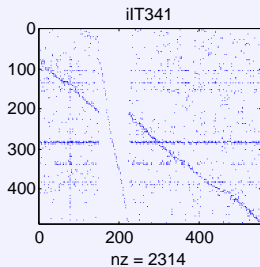
- SVD Golub & Reinsch
- GSVD (2 matrices) Van Loan; Paige & Saunders
- HOGSVD (N matrices) Ponnappalli, Saunders, Van Loan, Alter
Saturday 1:30pm session
- Parafac etc (tensors) Acar et al., previous talk!

Stoichiometric matrices

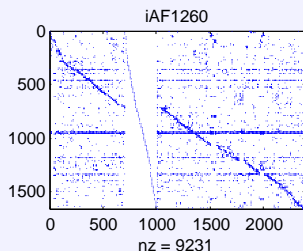
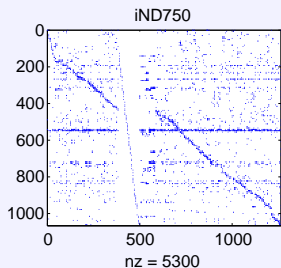
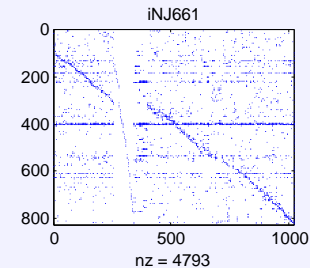
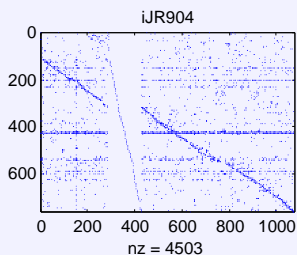
Rows: Chemical species

Cols: Chemical reactions

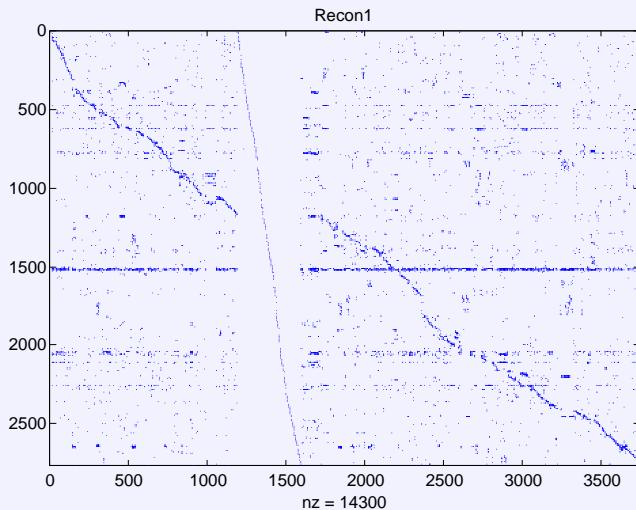
Models 1, 2, 3, 4 (all similar)



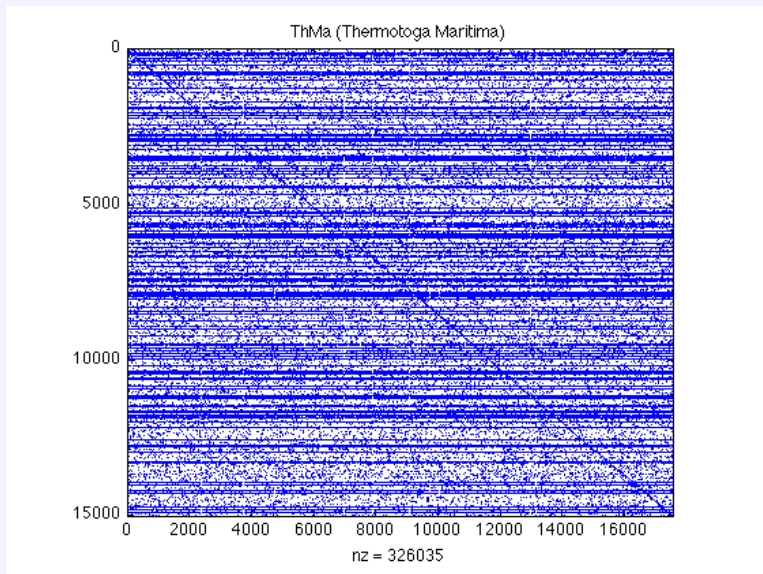
Models 5, 6, 7, 8 (all similar)



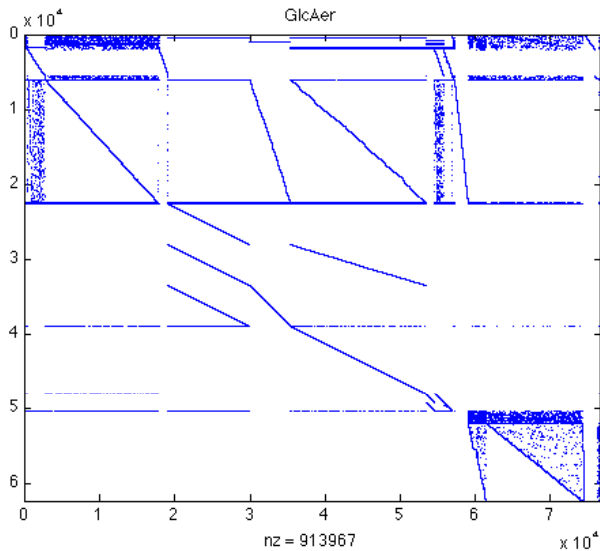
Model 9 (Recon1)



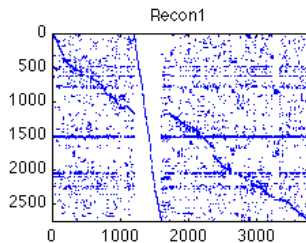
Model 10 (ThMa = *Thermotoga maritima*)



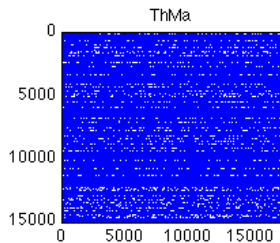
Model 11 (GlcAer)



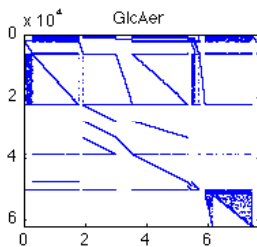
Models 9, 10, 11



nz = 14300



nz = 326035



nz = 913967 $\times 10^4$

Rank of stoichiometric matrices

Conservation analysis for biochemical networks

Conservation analysis

Goal: find subgroups conserved by biological systems

- Examples:
 - adenine nucleotide moiety (ADP, ATP, AMP)
 - NAD/NADH
 - CoA/Acetyl-CoA
- An important preliminary step in
 - evaluating drug targets
 - analyzing the transient behavior of biochemical networks

Finding rank(S) and null(S^T)

Conservation analysis reduces to finding rank(S) and null(S^T)

$$0 = \frac{d}{dt} \{z^T c(t)\} = z^T \frac{dc(t)}{dt} = z^T S v(t)$$

where z is a conserved moiety (group of chemical species)

Requires $S^T z = 0$

Finding $\text{rank}(S)$ and $\text{null}(S^T)$

Conservation analysis reduces to finding $\text{rank}(S)$ and $\text{null}(S^T)$

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Also part of conservation analysis:

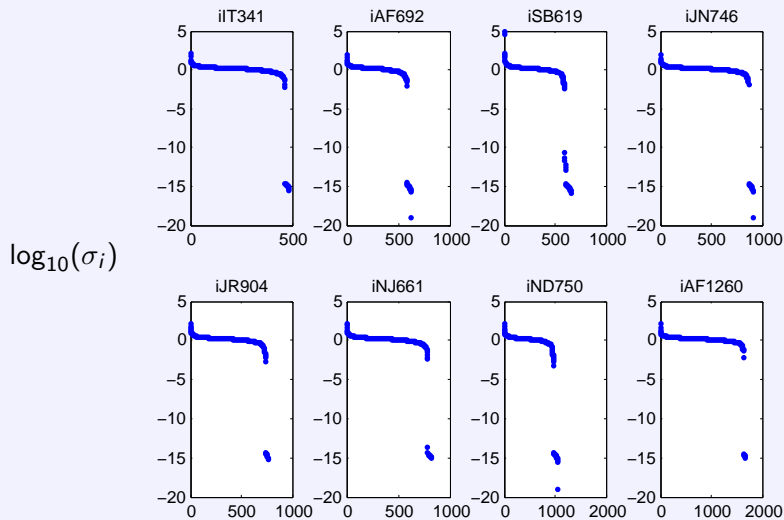
- Partitioning the rows (species) of S into **dependent** and **independent** rows (species)
- Computing a **link matrix** that describes the relations among the concentrations of dependent and independent species

rank(S) by SVD

Singular value decomposition	$S = UDV^T$
------------------------------	-------------

- $U^T U = I$ $V^T V = I$ D diagonal rank(S) = rank(D)
- Ideal for rank-estimation but U, V are dense
- model 9 (Recon1) 2800×3700 17 secs
 model 10 (ThMa) 15000×18000 11 hours
 model 11 (GlcAer) 62000×77000 ∞

Singular values of models 1–8 Dense SVD of S^T



rank(S) by QR

Householder QR factorization $SP = QR$

- $P = \text{col perm}$ $Q^T Q = I$ R diagonal $\text{rank}(S) = \text{rank}(R)$
- Nearly as reliable as SVD

rank(S) by QR

Householder QR factorization	$SP = QR$
------------------------------	-----------

- $P = \text{col perm}$ $Q^T Q = I$ R diagonal $\text{rank}(S) = \text{rank}(R)$
- Nearly as reliable as SVD
- Dense QR used by Vallabhajosyula, Chickarmane, Sauro (2005)
- Sparse QR (SPQR) now available: Davis (2013)
- model 9 (Recon1) 2800×3700 0.0 secs
 model 10 (ThMa) 15000×18000 2.5 secs
 model 11 (GlcAer) 62000×77000 0.2 secs(!)

rank(S) by LDU

Sparse LU with Threshold Rook Pivoting (TRP) $P_1SP_2 = LDU$

- $P_1, P_2 =$ perms D diagonal $\text{rank}(S) \approx \text{rank}(D)$
 L, U well-conditioned
- $L_{ij} = U_{ij} = 1$
 $|L_{ij}|$ **and** $|U_{ij}| \leq \text{facto1} = 4$ or 2 or 1.2, 1.1, ...

rank(S) by LDU

Sparse LU with Threshold Rook Pivoting (TRP)	$P_1 S P_2 = L D U$
--	---------------------

- $P_1, P_2 =$ perms D diagonal rank(S) \approx rank(D)
 L, U well-conditioned
- $L_{ij} = U_{ij} = 1$
 $|L_{ij}|$ **and** $|U_{ij}| \leq \text{facto1} = 4$ or 2 or 1.2, 1.1, ...
- LUSOL: Main engine in sparse linear/nonlinear optimizers
 MINOS, SQOPT, SNOPT
- model 9 (Recon1) 2800×3700 0.0 secs
 model 10 (ThMa) 15000×18000 4.0 secs
 model 11 (GlcAer) 62000×77000 158 secs

model	m	n	rank(S)		nnz(S)	nnz(Q)	nnz(R)	time	
			SVD	SPQR				SVD	SPQR
Recon1	2766	3742	2674	2674	14300	2750	21093	17.5	0.1
ThMa	15024	17582	14983	14983	326035	844096	10595016	11hrs	2.5
GlcAer	62212	76664	?	62182	913967	1287	916600	infty	0.2

model	m	n	rank(S)		nnz(S)	nnz(Q)	nnz(R)	time	
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ThMa	15024	17582	14983	14983	326035	844096	10595016	11hrs	2.5
GlcAer	62212	76664	?	62182	913967	1287	916600	infy	0.2

factol = 2.00 S = LDU

model	m	n	rank(S)	nnz(S)	nnz(L)	nnz(U)	time	
Recon1	2766	3742	2674	14300	4280	16463	0.1	
ThMa	15024	17582	14983	326035	30962	346122	4.1	
GlcAer	62212	76664	62182	913967	635571	1810491	186.2	

factol = 4.00 S = LDU

model	m	n	rank(S)	nnz(S)	nnz(L)	nnz(U)	time	
Recon1	2766	3742	2674	14300	2701	12896	0.1	
ThMa	15024	17582	14983	326035	36350	330485	4.0	
GlcAer	62212	76664	62182	913967	427456	1584188	157.9	

model	m	n	rank(S')		nnz(S)	nnz(Q)	nnz(R)	time	
			SVD	SPQR				SVD	SPQR
Recon1	3742	2766	2674	2674	14300	107935	36929	17.2	0.1
ThMa	17582	15024	14983	14983	326035	624640	605888	11hrs	0.7
GlcAer	76664	62212	?	62182	913967	3573696	4038988	infy	2.7

factol = 2.00 S' = LDU

model	m	n	rank(S')	nnz(S)	nnz(L)	nnz(U)	time	
Recon1	3742	2766	2674	14300	12832	7421	0.3	
ThMa	17582	15024	14983	326035	501198	358601	37.8	
GlcAer	76664	62212	62182	913967	1996892	709448	586.0	

factol = 4.00 S' = LDU

model	m	n	rank(S')	nnz(S)	nnz(L)	nnz(U)	time	
Recon1	3742	2766	2674	14300	9811	6093	0.2	
ThMa	17582	15024	14983	326035	410290	355475	14.8	
GlcAer	76664	62212	62182	913967	1823067	711906	791.2	

Large-scale LP, QP, NLP

SQOPT

$$\begin{array}{ll} \text{QP} & \text{minimize}_{x \in \mathbb{R}^n} \quad c^T x + \frac{1}{2}(x - x_0)^T H(x - x_0) \\ & \text{subject to} \quad \ell \leq \begin{pmatrix} x \\ Ax \end{pmatrix} \leq u \end{array}$$

- Includes LP ($H = 0$)
- H symmetric positive semidefinite
- A large and sparse
- LUSOL is main engine (for satisfying the constraints)
- Fortran 77 (\Rightarrow C implementation via f2c translator)
- Fortran 90 version can compile with **double precision (normal)** or **quadruple precision (for astounding results!)**

SNOPT

NLP

minimize $\phi(x)$
 $x \in \mathbb{R}^n$

subject to $l \leq \begin{pmatrix} x \\ Ax \\ c(x) \end{pmatrix} \leq u$

- $\phi(x)$ smooth nonlinear objective function
- $c(x)$ = vector of smooth nonlinear functions
- Best if gradients of nonlinear functions are known
- Uses SQOPT to solve sequence of QP subproblems

Flux Balance Analysis (FBA) on *Thermotoga maritima*

$$\min c^T v \quad \text{subject to} \quad Sv = 0, \quad l \leq v \leq u$$

S rows and cols 18210×17535

Nonzero S_{ij} 33602

max and min $|S_{ij}|$ 2×10^4 and 3×10^{-6}

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SQOPT in double precision (15 digits)

Feasibility tol 1e-6

Optimality tol 1e-6

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SQOPT in double precision (15 digits)

Feasibility tol $1e-6$

Optimality tol $1e-6$

SQOPT in quad precision (32 digits)

Feasibility tol $1e-15$

Optimality tol $1e-15$

Flux Balance Analysis (FBA) on *Thermotoga maritima*

$$\min c^T v \quad \text{subject to} \quad Sv = 0, \quad l \leq v \leq u$$

S rows and cols 18210×17535

Nonzero S_{ij} 33602

max and min $|S_{ij}|$ 2×10^4 and 3×10^{-6}

SQOPT in double precision (42 secs)

SQOPT EXIT 10 -- the problem appears to be infeasible

Problem name	ThMa		
No. of iterations	18500	Objective value	8.2286249495E-07
No. of infeasibilities	9	Sum of infeas	1.9606461069E-03
No. of degenerate steps	11611	Percentage	62.76
Max x (scaled)	3482 8.2E+00	Max pi (scaled)	18210 9.8E-01
Max x	5134 5.9E+00	Max pi	18210 1.0E+00
Max Prim inf(scaled)	32832 1.3E-03	Max Dual inf(scaled)	16417 1.0E+00
Max Primal infeas	32832 5.6E-06	Max Dual infeas	32669 2.3E+02

Flux Balance Analysis (FBA) on *Thermotoga maritima*

$$\min c^T v \quad \text{subject to} \quad Sv = 0, \quad l \leq v \leq u$$

S rows and cols 18210×17535

Nonzero S_{ij} 33602

max and min $|S_{ij}|$ 2×10^4 and 3×10^{-6}

Restart SQOPT in quad precision (36 secs)

SQOPT EXIT 0 -- finished successfully

Problem name	ThMa		
No. of iterations	498	Objective value	8.7036461686E-07
No. of infeasibilities	0	Sum of infeas	0.0000000000E+00
No. of degenerate steps	220	Percentage	44.18
Max x (scaled)	3482 8.2E+00	Max pi (scaled)	2907 1.3E+00
Max x	5134 5.9E+00	Max pi	15517 1.1E+00
Max Prim inf(scaled)	16475 5.2E-28	Max Dual inf(scaled)	13244 1.9E-32
Max Primal infeas	16475 5.2E-29	Max Dual infeas	13244 4.8E-33

Flux Balance Analysis (FBA) on GlcAer

$$\min c^T v \quad \text{subject to} \quad Sv = 0, \quad l \leq v \leq u$$

S rows and cols	68300 × 76664
Nonzero S_{ij}	926357
max and min $ S_{ij} $	8×10^5 and 5×10^{-5}

SQOPT in quad precision cold start, no scaling (30786 secs)

SQOPT EXIT 0 -- finished successfully

Problem name	GlcAer		
No. of iterations	84685	Objective value	-7.0382454070E+05
No. of degenerate steps	62127	Percentage	73.36
Max x	61436 6.3E+07	Max pi	25539 2.4E+07
Max Primal infeas	72623 3.0E-21	Max Dual infeas	17817 2.7E-21

Flux Balance Analysis (FBA) on GlcAer

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S rows and cols	68300 × 76664
Nonzero S_{ij}	926357
max and min $ S_{ij} $	8×10^5 and 5×10^{-5}

SQOPT in quad precision cold start, with scaling (4642 secs)

SQOPT EXIT 0 -- finished successfully

Problem name	GlcAer		
No. of iterations	37025	Objective value	-7.0382454070E+05
No. of infeasibilities	1	Sum of infeas	6.9661927856E-16
No. of degenerate steps	28166	Percentage	76.07
Max x (scaled)	59440 3.7E+00	Max pi (scaled)	40165 8.1E+11
Max x	61436 6.3E+07	Max pi	25539 2.4E+07
Max Prim inf(scaled)	81918 7.0E-16	Max Dual inf(scaled)	59325 1.5E-17
Max Primal infeas	81918 1.3E-07	Max Dual infeas	27953 2.0E-22

PDCO

Primal-dual interior method for convex optimization

PDCO (Matlab primal-dual convex optimizer)

$$\begin{array}{ll} \underset{x}{\text{minimize}} & \phi(x) \\ \text{subject to} & Ax = b, \quad \ell \leq x \leq u, \end{array}$$

where $\phi(x)$ is convex with known gradient and Hessian.

A may be a sparse matrix or an operator for Av and $A^T w$

e.g. Basis Pursuit (BP and BPDN) Chen, Donoho, Saunders 2001

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To ensure unique solutions, PDCO solves regularized problems:

$$\begin{aligned} & \underset{x, r}{\text{minimize}} && \phi(x) + \frac{1}{2} \|D_1 x\|^2 + \frac{1}{2} \|r\|^2 \\ & \text{subject to} && Ax + D_2 r = b, \quad \ell \leq x \leq u, \end{aligned}$$

where D_1, D_2 are diagonal and positive-definite.

PDCO (Matlab primal-dual convex optimizer)

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where D_1, D_2 are diagonal and positive-definite.

Typically $D_1 = \gamma I$ $\gamma = 10^{-3}$ or 10^{-4}

Same for D_2 if $Ax = b$ should be satisfied accurately

For least-squares problems $D_2 = I$

PDCO applied to FBA

$$\begin{array}{ll} \text{FBA} & \text{minimize } d^T v_e \\ & \text{subject to } S v_f - S v_r + S_e v_e = 0 \\ & v_f, v_r \geq 0, \quad \ell \leq v_e \leq u \end{array}$$

- Flux Balance Analysis = LP problem (Palsson 2006)
- d optimizes a biological objective
e.g., maximize replication rate in unicellular organisms
- v_e = exchange fluxes = sources and sinks of chemicals

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- Flux Balance Analysis = LP problem (Palsson 2006)
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- PDCO works with $A = \begin{bmatrix} S & -S & S_e \end{bmatrix}$ then $LL^T = AD^2A^T$
(sparse Cholesky with D increasingly ill-conditioned)

PDCO applied to FBA

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- d optimizes a biological objective
e.g., maximize replication rate in unicellular organisms
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- PDCO works with $A = [S \quad -S \quad S_e]$ then $LL^T = AD^2A^T$
(sparse Cholesky with D increasingly ill-conditioned)
- Solution is $v^* = v_f^* - v_r^*$ and v_e^*

PDCO applied to Entropy problem

$$\begin{aligned}
 \text{EP} \quad & \underset{v_f, v_r}{\text{minimize}} && v_f^T (\log v_f + c - e) + v_r^T (\log v_r + c - e) \\
 & \text{subject to} && Sv_f - Sv_r = -S_e v_e^* \\
 & && v_f, v_r > 0
 \end{aligned}$$

- $c =$ any vector, $e = (1, 1, \dots, 1)^T$
 $v_e^* =$ optimal exchange fluxes from FBA
- Entropy objective function is strictly convex
- Solution v_f^*, v_r^* is thermodynamically feasible
 (satisfies energy conservation and 2nd law of thermodynamics)

Fleming, Maes, Saunders, Ye, Palsson (2012)

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- <http://www.stanford.edu/group/SOL/>
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Future work

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- **Randomized numerical linear algebra**
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 - How to parallelize algorithms to handle truly massive data sets?
 - For example, **LSRN**
- **High-dimensional statistics**
 - How to make valid inference when the number of problem parameters is much larger than the sample size?
 - How to construct confidence regions and obtain p-values in this setting?

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