STATS 200: Introduction to Statistical Inference Lecture 5: Testing a simple null hypothesis

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# Statistical inference = $Probability^{-1}$

Today: Does my data come from a prescribed distribution, *F*? This is oftentimes called testing **goodness of fit**.

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Example: You roll a 6-sided die n times, and observe 1, 3, 1, 6, 4, 2, 5, 3, ... Is this a fair die?

# Example: Einstein's theory of Brownian motion

Motion of a tiny (radius  $\approx 10^{-4}$  cm) particle suspended in water:



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Example: Einstein's theory of Brownian motion Albert Einstein (1905):  $P_{t+\Delta t} \sim \mathcal{N}\left(P_t, \begin{pmatrix} \sigma^2 & 0\\ 0 & \sigma^2 \end{pmatrix}\right)$ , where

$$\sigma^2 = \frac{RT}{3\pi\eta r N_A} (\Delta t).$$

- *P<sub>t</sub>*: position of particle at time t
- ► *R*: ideal gas constant
- ► *T*: absolute temperature
- η: viscosity of water
- r: radius of particle
- ► N<sub>A</sub>: Avogadro's number

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Jean Perrin (1909): Measured the position of a particle every 30 seconds to verify Einstein's theory (and to compute  $N_A$ ). For his experiment,  $\sigma^2 = 2.23 \times 10^{-7}$  cm<sup>2</sup>.

Does Perrin's data fit with Einstein's model?

# Null and alternative hypotheses

A **hypothesis test** is a binary question about the data distribution. Our goal is to either accept a **null hypothesis**  $H_0$  (which specifies something about this distribution) or to reject it in favor of an **alternative hypothesis**  $H_1$ .

If  $H_0$  (similarly  $H_1$ ) completely specifies the probability distribution for the data, then the hypothesis is **simple**. Otherwise it is **composite**.

Today we'll focus on testing simple null hypotheses  $H_0$ .

#### Simple vs. composite

Example: Let  $X_1, \ldots, X_6$  be the number of times we obtain 1 to 6 in *n* dice rolls. This null hypothesis is simple:

 $H_0: (X_1, ..., X_6) \sim \text{Multinomial} (n, (\frac{1}{6}, ..., \frac{1}{6})).$ 

#### Simple vs. composite

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$$H_0: (X_1, ..., X_6) \sim \text{Multinomial} (n, (\frac{1}{6}, ..., \frac{1}{6})).$$

We might wish to test this null hypothesis against the simple alternative hypothesis

$$H_1: (X_1, \ldots, X_6) \sim \text{Multinomial} \left( n, \left( \frac{1}{9}, \frac{1}{9}, \frac{1}{9}, \frac{2}{9}, \frac{2}{9}, \frac{2}{9} \right) \right),$$

or perhaps against the compositive alternative hypothesis

$$H_1: (X_1, \dots, X_6) \sim \text{Multinomial}(n, (p_1, \dots, p_6))$$
  
for some  $(p_1, \dots, p_6) \neq (\frac{1}{6}, \dots, \frac{1}{6})$ 

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#### Simple vs. composite

Example: Let  $(X_1, Y_1), (X_2, Y_2), (X_3, Y_3), \ldots$  be the displacement vectors  $P_{30} - P_0, P_{60} - P_{30}, P_{90} - P_{60}, \ldots$  where  $P_t \in \mathbb{R}^2$  is the position of a particle at time *t* in Perrin's experiment. Einstein's theory corresponds to the simple null hypothesis

$$H_0: (X_1, Y_1), \ldots, (X_n, Y_n) \stackrel{IID}{\sim} \mathcal{N}(0, 2.23 \times 10^{-7} I).$$

To test the theory qualitatively, but possibly allow for an error in Einstein's formula for  $\sigma^2$ , we might test the composite null hypothesis

$$H_0: (X_1, Y_1), \ldots, (X_n, Y_n) \stackrel{IID}{\sim} \mathcal{N}(0, \sigma^2 I)$$
 for some  $\sigma^2 > 0$ .

One can pose a number of different possible alternative hypotheses  $H_1$  to the above nulls.

#### Test statistics

A **test statistic**  $T := T(X_1, ..., X_n)$  is any statistic such that extreme values (large or small) of T provide evidence against  $H_0$ .

Example: Let  $X_1, \ldots, X_6$  count the results from *n* dice rolls, and let

$$T = \left(\frac{X_1}{n} - \frac{1}{6}\right)^2 + \ldots + \left(\frac{X_6}{n} - \frac{1}{6}\right)^2$$

Large values of  $\mathcal{T}$  provide evidence against the null hypothesis of a fair die,

$$H_0: (X_1, \ldots, X_6) \sim \mathsf{Multinomial}\left(n, \left(rac{1}{6}, \ldots, rac{1}{6}
ight)
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#### Test statistics

Example: Let  $(X_1, Y_1), \ldots, (X_n, Y_n)$  be the displacements from Perrin's experiment. For testing

$$H_0: (X_1, Y_1), \ldots, (X_n, Y_n) \stackrel{IID}{\sim} \mathcal{N}(0, 2.23 \times 10^{-7} I).$$

the following are possible test statistics:

$$\bar{X} = \frac{1}{n}(X_1 + \ldots + X_n)$$
  
$$\bar{Y} = \frac{1}{n}(Y_1 + \ldots + Y_n)$$
  
$$V = \frac{1}{n}(X_1^2 + Y_1^2 + \ldots + X_n^2 + Y_n^2)$$

(Values of  $\bar{X}$  or  $\bar{Y}$  much larger or smaller than 0, or values of V much larger or smaller than  $2 \times 2.23 \times 10^{-7}$ , provide evidence against  $H_0$  in favor of various alternatives  $H_1$ .)

Let  $R_i = X_i^2 + Y_i^2$ . Suppose we are interested in testing whether  $R_1, \ldots, R_n$  are distributed as  $2.23 \times 10^{-7} \chi_2^2$  (their distribution under  $H_0$ ). We can plot a histogram of these values:



Histogram of X^2+Y^2

X^2+Y^2

Deviations from  $2.23 \times 10^{-7} \chi_2^2$  are better visualized by a hanging histogram, which plots  $O_i - E_i$  where  $O_i$  is the observed count for bin *i* and  $E_i$  is the expected count under the  $2.23 \times 10^{-7} \chi_2^2$  distribution:



#### Hanging histogram of X^2+Y^2

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A test statistic can be  $T = \sum_{i=1}^{6} (O_i - E_i)^2$ .

Problem: Let  $p_i$  be the probability that the hypothesized chi-squared distribution assigns to bin *i*. If  $H_0$  were true, then  $O_i \sim \text{Binomial}(n, p_i)$  and  $E_i = np_i = \mathbb{E}[O_i]$ . So

$$\operatorname{Var}[O_i] = \mathbb{E}[(O_i - E_i)^2] = np_i(1 - p_i).$$

The variation in  $O_i$  is smaller, and scales approximately linearly with  $p_i$ , if  $p_i$  is close to 0. This might explain why the bars were smaller on the right side of the hanging histogram.

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Solution: We can "stabilize the variance" by looking at  $\frac{O_i - E_i}{\sqrt{E_i}} = \frac{O_i - E_i}{\sqrt{np_i}}$ .

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Solution: We can "stabilize the variance" by looking at 
$$\frac{O_i - E_i}{\sqrt{E_i}} = \frac{O_i - E_i}{\sqrt{np_i}}$$
.

Or alternatively, we can look at  $\sqrt{O_i} - \sqrt{E_i}$ . (Taylor expansion of  $\sqrt{x}$  around  $x = E_i$  yields  $\sqrt{O_i} - \sqrt{E_i} \approx \frac{1}{2\sqrt{E_i}}(O_i - E_i)$ , so this has a similar effect as  $\frac{O_i - E_i}{2\sqrt{E_i}}$  when  $O_i - E_i$  is small.)

The hanging chi-gram plots  $\frac{O_i - E_i}{\sqrt{E_i}}$ :

Hanging chi-gram of X^2+Y^2



The test statistic  $T = \sum_{i=1}^{6} \frac{(O_i - E_i)^2}{E_i}$  is called **Pearson's chi-squared statistic for goodness of fit**.

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Tukey's hanging rootogram plots  $\sqrt{O_i} - \sqrt{E_i}$ :

#### Hanging rootogram of X^2+Y^2

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We may take as test statistic  $T = \sum_{i=1}^{6} (\sqrt{O_i} - \sqrt{E_i})^2$ .

A **QQ plot** (or probability plot) compares the sorted values of  $R_1, \ldots, R_n$  with the  $\frac{1}{n+1}, \frac{2}{n+1}, \ldots, \frac{n}{n+1}$  quantiles of the hypothesized  $2.23 \times 10^{-7} \chi_2^2$  distribution:



Values close to the line y = x indicate a good fit.

How do we get a test statistic from a QQ plot? One way is to take the maximum vertical deviation from the y = x line: Let  $R_{(1)} < \ldots < R_{(n)}$  be the sorted values of  $R_1, \ldots, R_n$ . Take

$$T = \max_{i=1}^{n} \left| R_{(i)} - F^{-1} \left( \frac{i}{n+1} \right) \right|,$$

where F is the CDF of the  $2.23 \times 10^{-7} \chi_2^2$  distribution so  $F^{-1}(t)$  is its  $t^{\text{th}}$  quantile.

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where F is the CDF of the  $2.23 \times 10^{-7} \chi_2^2$  distribution so  $F^{-1}(t)$  is its  $t^{\text{th}}$  quantile.

Problem: For values of R where the distribution has high density, the quantiles are closer together, so we expect a smaller vertical deviation. This explains why we see more vertical deviation in the upper right of the last QQ plot.

Solution: We may stabilize the spacings between quantiles by considering instead

$$T = \max_{i=1}^n \left| F(R_{(i)}) - \frac{i}{n+1} \right|.$$

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Solution: We may stabilize the spacings between quantiles by considering instead

$$T = \max_{i=1}^n \left| F(R_{(i)}) - \frac{i}{n+1} \right|.$$

This is almost the same as the **one-sample Kolmogorov-Smirnov (K-S) statistic**,

$$T_{\mathcal{KS}} = \max_{i=1}^{n} \max\left( \left| F(R_{(i)}) - \frac{i}{n} \right|, \left| F(R_{(i)}) - \frac{i-1}{n} \right| \right).$$

(You can show  $\frac{i-1}{n} < \frac{i}{n+1} < \frac{i}{n}$ , and the difference between T and  $T_{KS}$  is negligible for large n.)

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Supposing that we've picked our test statistic T, how large (or small) does T need to be, before we can safely assert that  $H_0$  is false?

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Supposing that we've picked our test statistic T, how large (or small) does T need to be, before we can safely assert that  $H_0$  is false?

In most cases we can never be 100% sure that  $H_0$  is false. But we can compute T from the observed data and compare with the sampling distribution of T if  $H_0$  were true. This is called the **null distribution** of T.

Example: Consider

$$H_0: (X_1, Y_1), \ldots, (X_n, Y_n) \stackrel{IID}{\sim} \mathcal{N}(0, 2.23 \times 10^{-7} I).$$

Under  $H_0$ ,  $\bar{X} \sim \mathcal{N}(0, 2.23 \times 10^{-7}/n)$ . This normal distribution is the null distribution of  $\bar{X}$ .

Here's the PDF for the null distribution of  $\bar{X}$ , when n = 30:

Null distribution of Xbar



If, for the observed data,  $\bar{X} = 0.5 \times 10^{-4}$ , this would not provide strong evidence against  $H_0$ . In this case we might accept  $H_0$ .

Here's the PDF for the null distribution of  $\bar{X}$ , when n = 30:

Null distribution of Xbar



If, for the observed data,  $\bar{X} = 2.5 \times 10^{-4}$ , this would provide strong evidence against  $H_0$ . In this case we might reject  $H_0$ .

The **rejection region** is the set of values of T for which we choose to reject  $H_0$ . The **acceptance region** is the set of values of T for which we choose to accept  $H_0$ .

We choose the rejection region so as to control the probability of **type l error**:

 $\alpha = \mathbb{P}_{H_0}[\text{reject } H_0]$ 

This value  $\alpha$  is also called the **significance level** of the test.

If, under its null distribution, T belongs to the rejection region with probability  $\alpha$ , then the test is level- $\alpha$ .

(Notation: For a simple null hypothesis  $H_0$ , we write  $\mathbb{P}_{H_0}[\mathcal{E}]$  to denote the probability of event  $\mathcal{E}$  under  $H_0$ , i.e. the probability of  $\mathcal{E}$  if  $H_0$  were true.)



Null distribution of Xbar

Example: A (two-sided) level- $\alpha$  test might reject  $H_0$  when  $\bar{X}$  falls in the above shaded regions. Mathematically, let  $z(\alpha)$  denote the  $1 - \alpha$  quantile, or "upper  $\alpha$  point", of the distribution  $\mathcal{N}(0, 1)$ . As  $\bar{X} \sim \mathcal{N}(0, \sigma^2/n)$  under  $H_0$  (where  $\sigma^2 = 2.23 \times 10^{-7}$ ), the rejection region should be  $(-\infty, -\frac{\sigma}{\sqrt{n}} \times z(\alpha/2)] \cup [\frac{\sigma}{\sqrt{n}} \times z(\alpha/2), \infty)$ .

## **P-values**

The *p***-value** is the smallest significance level at which your test would have rejected  $H_0$ .

For a one-sided test that rejects for large T, letting  $t_{obs}$  denote the value of T computed from the observed data, the *p*-value is  $\mathbb{P}_{H_0}[T \ge t_{obs}]$ .

For a two-sided test that rejects at the  $\alpha/2$  and  $1 - \alpha/2$  quantiles of the null distribution of T, the *p*-value is 2 times the smaller of  $\mathbb{P}_{H_0}[T \ge t_{obs}]$  and  $\mathbb{P}_{H_0}[T \le t_{obs}]$ .

The *p*-value provides a quantitative measure of the extent to which the data supports (or does not support)  $H_0$ . It is preferable to report the exact *p*-value, rather than to just say "we rejected at level-0.05".

# A word of caution

Accepting (or failing to reject)  $H_0$  **does not** imply there is strong evidence that  $H_0$  is true. Both of the following are possible:

- The particular test statistic you chose is not good at distinguishing the null hypothesis H<sub>0</sub> from the true distribution. Or equivalently, the true distribution is not well-captured by the alternative H<sub>1</sub> that your test statistic is targeting. (For example, in Perrin's data, if there is significant drift in the y direction, you would not detect this using the test statistic X̄.)
- You do not have enough data to reject H<sub>0</sub> at the significance level that you desire. In this case your study might be underpowered—we'll discuss this issue a couple weeks from now.

# Determining the null distribution

To figure out the rejection region, we must understand the null distribution of the test statistic. There are three methods:

- ► Sometimes we can derive the null distribution exactly, for example in the previous slides where the test statistic is X̄ and X<sub>1</sub>,..., X<sub>n</sub> are normally distributed under H<sub>0</sub>.
- Sometimes we can derive an asymptotic approximation, using tools such as the CLT and continuous mapping theorem.
- ▶ When *H*<sup>0</sup> is simple, we can always obtain the null distribution by simulation.

## Using an asymptotic null distribution

Example: Let  $(X_1, \ldots, X_6)$  denote the counts of 1 to 6 from *n* rolls of a die, and consider testing the simple null of a fair die

$$H_0: (X_1, \ldots, X_6) \sim \mathsf{Multinomial}\left(n, \left(\frac{1}{6}, \ldots, \frac{1}{6}\right)\right)$$

using the test statistic

$$T = \left(\frac{X_1}{n} - \frac{1}{6}\right)^2 + \ldots + \left(\frac{X_6}{n} - \frac{1}{6}\right)^2.$$

Recall from last lecture that for large *n*, *T* is approximately distributed as  $\frac{1}{6n}\chi_5^2$ .

To perform an **asymptotic level**- $\alpha$  **test**, we may reject  $H_0$  when  $t_{obs}$  exceeds  $\frac{1}{6n}\chi_5^2(\alpha)$ , where  $\chi_n^2(\alpha)$  denotes the  $1 - \alpha$  quantile, or "upper  $\alpha$  point", of the  $\chi_n^2$  distribution.

# Using a simulated null distribution

Example: Let T be Pearson's chi-squared statistic for goodness of fit for the values  $X_1^2 + Y_1^2, \ldots, X_{30}^2 + Y_{30}^2$  from Perrin's experiments, discussed previously. We may simulate the null distribution of T:



This shows the 1000 values of T across 1000 simulations. The observed value  $t_{obs} = 2.83$  for Perrin's real data is in red.

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#### Using a simulated null distribution

Example: Let T be the K-S statistic for  $X_1^2 + Y_1^2, \ldots, X_{30}^2 + Y_{30}^2$ , discussed previously. We may simulate the null distribution of T:



The observed value  $t_{obs} = 0.132$  for Perrin's real data is in red.

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### Using a simulated null distribution

We obtain an approximate *p*-value as the fraction of simulated values of T larger than  $t_{obs}$ . (For a two-sided test, we would take either the fraction of simulated values of T larger than  $t_{obs}$  or smaller than  $t_{obs}$ , and multiply this by 2.)

For Perrin's data, the Pearson chi-squared *p*-value is 0.754, and the K-S *p*-value is 0.612. We accept  $H_0$  in both cases, and neither test provides significant evidence against Einstein's theory of Brownian motion.