

Lecture 21

Topics:

- I) Hypothesis Formalism
- II) approximate p-values
translate approx z-scores
to probabilities
- III) exact p-values

e.g. Suppose your data gives

$$\hat{\theta} = 29.2$$

where the sampling variability
in $\hat{\theta}$ is described by

$$\hat{\theta} \sim N(0, 100).$$

Question: Test the null hypothesis

$$H_0: \theta = 0.$$

Fancy syntax
for the words
"is the sample
 $\hat{\theta}$ consistent with
the population
value $\theta = 0?$ "

Answer:

Find the z-score
of $\hat{\theta} = 29.2$ under
the assumption $\theta = 0$.

$$z = \frac{\hat{\theta} - E(\hat{\theta})}{sd(\hat{\theta})} = \frac{29.2 - 0}{10}$$

$$\stackrel{H_0}{=} \frac{29.2 - 0}{10}$$

$$= 2.92$$

(1)

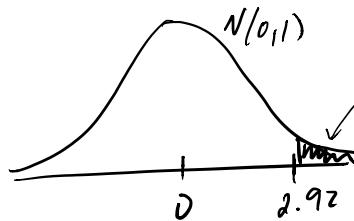
Since a z-score of 2.92 is kinda unlikely we are inclined to reject H_0 in favor for a different value of θ .

(2)

... But how unlikely is "kinda unlikely"?

Here is the picture:

if $\theta = 0$ then $z = \frac{\hat{\theta} - 0}{10}$
is a draw from



This area is 0.0018
and is called a
p-value for testing
 $H_0: \theta = 0$.

This is what we observed

- * P-values quantifies how hard it is to get $\hat{\theta} = 29.2$ (or even more rare) assuming H_0 is true.
- * Small p-values \Rightarrow more rare
 $\Rightarrow H_0$ is probably false

large p-values \Rightarrow less rare
 $\Rightarrow H_0$ is consistent with
the data $\hat{\theta} = 2.92$

- * P-values effectively translate a z-score to a rarity measure that others can understand. i.e. since the p-value = 0.0018 = $\frac{1.05}{600}$

we would need to sample from $N(0, 100)$ about 600 times before getting $\hat{\theta} = 2.92$ or bigger.

Using the CLT for approximating p-values

(3)

The CLT is the main tool for translating z-scores to p-values when testing a population mean μ .

e.g. Huge list of numbers

$$y_1, y_2, \dots, y_N \text{ where } N = 5 \times 10^{22}$$

You want to estimate

$$\mu = \frac{1}{N} \sum_{i=1}^N y_i = \bar{y}$$

i.e. the list average.

Suppose the list s.d. is given by

$$\sigma = \sqrt{\frac{1}{N} \sum_{i=1}^N (y_i - \bar{y})^2} = 7.2$$

Sometimes called the population s.d.

To construct $\hat{\mu}$ you randomly sample (w/pop) $n = 45$ numbers from the list

& get y_1, y_2, \dots, y_n

with

$$\hat{\mu} = \frac{1}{n} \sum_{i=1}^n y_i = 22.1$$

Question: Test the null hypothesis

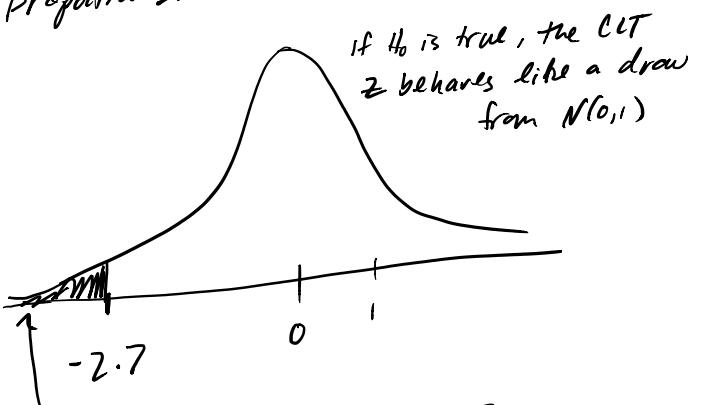
$$H_0: \mu = 25$$

Answer: First find the z-score of $\hat{\mu} = 22.1$ given $\mu = 25$ & $\sigma = 7.2$.

$$z = \frac{\hat{\mu} - E(\hat{\mu})}{\text{sd}(\hat{\mu})} = \frac{\hat{\mu} - \mu}{\left(\frac{\sigma}{\sqrt{n}}\right)} = \frac{22.1 - 25}{\frac{7.2}{\sqrt{45}}} = -2.7$$

Now find the p-value of this z-score to quantify rarity in terms of proportions.

(4)



$$\text{this area} = \text{p-value} = 0.0035$$

So, we would see a z-score of -2.7 or more extreme ~ 3 out of 1000 tries.

This is evidence that H_0 is wrong.

Remember: Hypothesis testing is only a technique for arguing against someone who believes H_0 .

Sometimes there is an agreed upon p-value cutoff (also called " α -level" or "the level of the test") where if the (p-value) < (p-value cutoff) then the observed z-score is rare enough that H_0 is rejected.

In the last example if the p-value cutoff is 0.01, H_0 is rejected. Alternatively if the p-value cutoff is 0.001 then H_0 can not be rejected.

Note: Hypothesis testing only works when H_0 is rejected.

(5)

What happens when σ is not known

e.g. Same example as before but suppose we are not given that $\sigma = 7.2$. To handle this case just estimate σ .

The Data:

$$Y_1, Y_2, \dots, Y_n \text{ where } n=45$$

where

$$\hat{\mu} = \frac{1}{n} \sum_{i=1}^n Y_i = 22.1$$

$$\hat{\sigma} = \sqrt{\frac{1}{n-1} \sum_{i=1}^n (Y_i - \hat{\mu})^2} = 6.9$$

Question: Test the Null

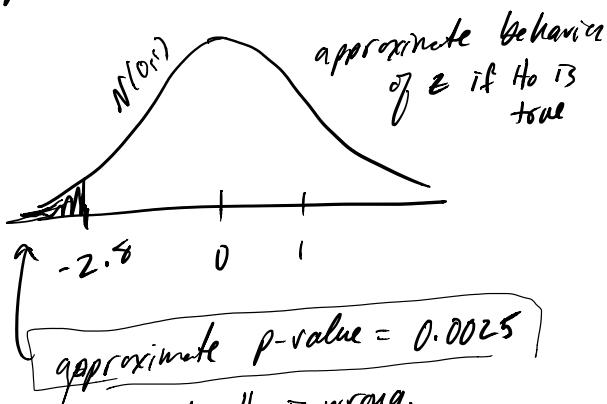
$$H_0: \mu = 25$$

Sometimes called
the sample
standard deviation
 s .

Answer: the z-score of the estimate:

$$z = \frac{\hat{\mu} - E(\hat{\mu})}{\text{sd}(\hat{\mu})} = \frac{\hat{\mu} - \mu}{\frac{\sigma}{\sqrt{n}}} = \frac{22.1 - 25}{\frac{7.2}{\sqrt{45}}} \approx \frac{22.1 - 25}{\frac{7.2}{\sqrt{45}}} = -2.8$$

Now, if we just ignore the fact that we have $\hat{\sigma}$ instead of σ we can quickly find an approximate p-value



which suggests H_0 is wrong.

Note: if n is small this approximation could be wrong.

(6)

Question: Can one compute an exact p-value for the observations

$$\hat{\mu} = 22.1, \hat{\sigma} = 6.9 ?$$

The approximate p-value 0.0025 is based on two approximations

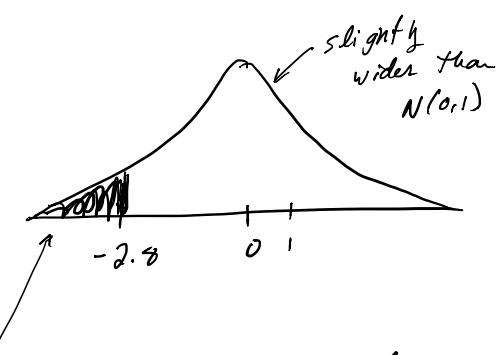
- I) the CLT that $\hat{\mu}$ fluctuates like a Normal distribution
- II) using $\hat{\sigma}$ instead of σ for computing the Z-score.

If you assume the population is normally distributed then the approximated z-score

$\frac{\hat{\mu} - \mu}{\hat{\sigma}/\sqrt{n}}$ is a $t^{(n-1)}$ random variable

with pdf

"a t distribution on
 $n-1$ degrees of freedom"



This area is the exact p-value under the assumption that the population is Normal.

using a calculator gives

the exact p-value = 0.0037

Comparing two population averages

e.g. Two huge lists of numbers

$$x_1, x_2, \dots,$$

$$y_1, y_2, \dots$$

Let

μ_x = the Ave of the x_i 's

μ_y = the Ave of the y_i 's.

σ_x = the "population s.d." of the x_i 's

σ_y = the "population s.d." of the y_i 's.

You want to estimate

$$\mu_x - \mu_y = ?$$

So you sample 25 numbers from the first list & get

$$X_1, X_2, \dots, X_{n_x}, n_x = 25$$

also sample 20 numbers from the second list & get

$$Y_1, Y_2, \dots, Y_{n_y}, n_y = 20$$

Suppose this data results in

$$\hat{\mu}_x = \bar{X} = -2.5$$

$$\hat{\mu}_y = \bar{Y} = 1.23$$

$$\begin{aligned} \hat{\sigma}_x &= 10.2 \\ \hat{\sigma}_y &= 5.1 \end{aligned} \quad \left\{ \begin{array}{l} \text{sample standard deviations of} \\ \text{the two samples.} \end{array} \right.$$

Question: Test the null hypothesis (7) (8)

$$H_0: \mu_x = \mu_y \text{ (i.e. } \mu_x - \mu_y = 0)$$

with a p-value cut off of 0.001.

Answer: Start by approximating the z-score of the estimate

$$\hat{\mu}_x - \hat{\mu}_y = -2.5 - 1.23$$

using the null hypothesis H_0 that $\mu_x - \mu_y = 0$

Note: MF1 & MF2 give

$$E(\hat{\mu}_x - \hat{\mu}_y) = \mu_x - \mu_y$$

$$sd(\hat{\mu}_x - \hat{\mu}_y) = \sqrt{\frac{\sigma_x^2}{25} + \frac{\sigma_y^2}{20}}$$

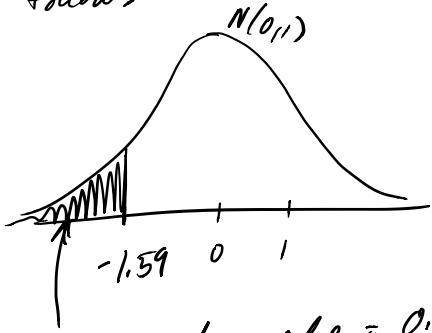
Therefore

$$\begin{aligned} \text{z-score} &= \frac{(\hat{\mu}_x - \hat{\mu}_y) - (\mu_x - \mu_y)}{\sqrt{\frac{\sigma_x^2}{25} + \frac{\sigma_y^2}{20}}} \\ &= \frac{(-2.5 - 1.23) - (0)}{\sqrt{\frac{\sigma_x^2}{25} + \frac{\sigma_y^2}{20}}} \quad \begin{array}{l} \text{hypothetical} \\ \text{value of} \\ \mu_x - \mu_y \end{array} \end{aligned}$$

$$\approx \frac{(-2.5 - 1.23) - (0)}{\sqrt{\frac{10.2^2}{25} + \frac{5.1^2}{20}}} \quad \begin{array}{l} \text{plug in} \\ \hat{\sigma}_x = 10.2 \text{ &} \\ \hat{\sigma}_y = 5.1 \text{ for} \\ \sigma_x^2 \text{ & } \sigma_y^2 \end{array}$$

$$= -1.59$$

The approximate z-score -1.59 (9) translates to an approximate p-value as follows



$$\text{approximate p-value} = 0.056$$

This is much bigger than the cut-off \Rightarrow can't reject H_0 .

If you want an exact p-value in this case you must assume both populations are exactly normal & $\sigma_x = \sigma_y = \sigma$

when $\sigma_x = \sigma_y$ notice

$$\begin{aligned} \text{sd}(\hat{\mu}_x - \hat{\mu}_y) &= \sqrt{\frac{\sigma_x^2}{n_x} + \frac{\sigma_y^2}{n_y}} \\ &= \sqrt{\frac{\sigma^2}{n_x} + \frac{\sigma^2}{n_y}} \\ &= \sqrt{\sigma^2 \left(\frac{1}{n_x} + \frac{1}{n_y} \right)} \\ &= \sigma \sqrt{\frac{1}{n_x} + \frac{1}{n_y}} \end{aligned}$$

Now approximate σ with a "pooled estimate of s.d." denoted $\hat{\sigma}_p$ or s_p & given by

$$\hat{\sigma}_p = \sqrt{\underbrace{\frac{n_x-1}{n_x+n_y-2} \hat{\sigma}_x^2 + \frac{n_y-1}{n_x+n_y-2} \hat{\sigma}_y^2}_{\text{a weighted avg of } \hat{\sigma}_x^2 \text{ & } \hat{\sigma}_y^2}}$$

Now under the normal population assumption & $\sigma_x = \sigma_y = \sigma$ the approximate z-score (10)

$\frac{(\hat{\mu}_x - \hat{\mu}_y) - (\mu_x - \mu_y)}{\hat{\sigma}_p \sqrt{\frac{1}{n_x} + \frac{1}{n_y}}}$ is a t random variable on $n_x + n_y - 2$ degrees of freedom.

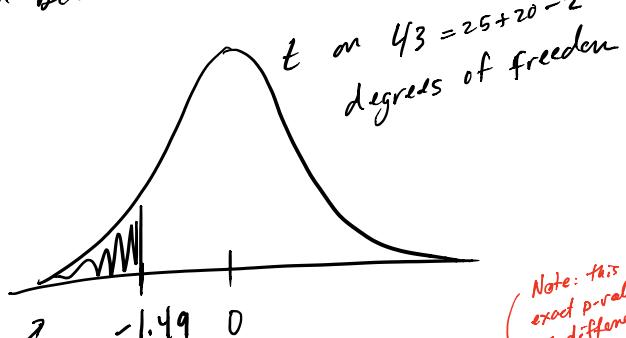
In the above example

$$\hat{\sigma}_p = \sqrt{\frac{25-1}{25+20-2} (10.2)^2 + \frac{20-1}{25+20-2} (5.1)^2} = 8.34$$

And therefore the approximate z-score of the estimates, assuming $H_0: \mu_x - \mu_y = 0$, is given by

$$\frac{(\hat{\mu}_x - \hat{\mu}_y) - (\mu_x - \mu_y)}{\hat{\sigma}_p \sqrt{\frac{1}{n_x} + \frac{1}{n_y}}} = \frac{-2.5 - 1.23 - 0}{8.34 \sqrt{\frac{1}{25} + \frac{1}{20}}} = -1.49$$

which behave like a draw from



This is the exact p-value

$$= 0.07$$

\therefore do not reject H_0 since $\underline{\text{p-value}} > \underline{\text{cut-off}}$

Note: this exact p-value is different from the "quick guess" method but not different enough to change the conclusion