# Section 12: Hypothesis testing and p values

STOR 155.02, Spring '21

updated 2021-04-13

# What you will learn

• is in the title

#### Resources

• Textbook: 5.3

# Eyes on the schedule

#### Four lectures remain

• April 13, 20, 27, May 4

## Final project due May 11, 3p.m. sharp

- no excuses for lateness
- except by approved university exam excuses or exceptional circumstances
- assigned one week prior

# Formula: pct (decimal) CI

a. Find the low quantile  $z^*$ 

$$P\left(Z \leq -z^*
ight) = rac{1-pct}{2}$$

Excel: NORM.S.INV(.5\*(1-pct)) and Python: stats.norm.ppf(.5\*(1-pct))

with pct in decimals, e.g. 0.025 for 95% CI

b. or find the high quantile (same  $z^*$ )

$$P(Z \geq z^*) = 1 - rac{1-pct}{2}$$

c. Calculate the sample point estimates

$$\mathrm{low} = \hat{m}_n - z^* rac{\hat{s}_n}{\sqrt{n}}, \quad \mathrm{high} = \hat{m}_n + z^* rac{\hat{s}_n}{\sqrt{n}}$$

**Shortcut:** NORM.INV(.5\*(1-pct), mhat, shat/sqrtn) for low

Shortcut: stats.norm.interval(pct, mhat, shat/sqrtn) both



# Hypothesis testing

# Starter example

#### **Adults in Turkey**

- random sample
- mean caloric intake of 3500 calories / day
- with s.d. of 500 calories

(Source: FAO, 2006-08 via Wikipedia)

# Sample of 100 Ankara University students

- random sample
- average intake of 3450 calories / day

# Conclude Ankara U. student average calories is different from Turkish population average?

- could happen if mean for Ankara U. students is in reality smaller
- or if we got a different average just because of randomness in sampling

## Assume averages are the same

# What is the probability we would get the sample average 3450 or farther away from 3500 from a random sample size 100?

We have point estimates for the true mean (3500) and standard deviation (500).

By the CLT we know the distribution of the sample mean  $\bar{X}_{100}$  is approximately Normal  $(3500, \frac{500}{\sqrt{100}})$ ,

$$P\left(rac{|ar{X}_{100}-3500|}{50}>rac{|3450-3500|}{50}
ight)pprox P\left(Z>1 ext{ or } Z<-1
ight) \ =2P\left(Z>1
ight)pprox 0.32$$

That's a pretty high chance a random sample of size 100 would have average like the one we saw, or smaller!

Why the 2 above? Area under bell curve for z < -1 is the same as for z > 1.

## Hypothesis test recipe

#### Model

Data  $x_1 \dots x_n$  sampled randomly from an unknown population r.v. X, with true mean m and s.d. s.

#### **Research question**

Do our data give strong evidence that

$$m 
eq m_0$$

for some number  $m_0$ ?

Could also consider asking a one-sided question,  $m < m_0$  or  $m > m_0$ .

#### **Point estimates**

 $\hat{s}_n$ ,  $\hat{m}_n$  are point estimates for s,m based on the data, e.g.  $\hat{m}_n$  is the sample mean of  $x_1 \dots x_n$ .

#### $H_0$ 'null hypothesis', $H_1$ 'alternative'

- $H_0$  is the statement you want to reject (if possible)
- using the hypothesis test

in this class will always be about true mean m being equal to some other number  $m_0$ 

$$H_0: m=m_0, \qquad H_1: m
eq m_0$$

or can consider  $H_1: m < m_0$ ,  $H_1: m > m_0$ .

Will try to 'reject  $H_0$  in favor of  $H_1$ '

Make confidence interval for  $ar{X}_n$  assuming  $H_0$  is true

$$\mathrm{low} \, = \hat{m}_n - z^* rac{\hat{s}_n}{\sqrt{n}}, \quad \mathrm{high} \, = \hat{m}_n + z^* rac{\hat{s}_n}{\sqrt{n}}$$

If  $m_0 >$  high or  $m_0 <$  low, reject  $H_0$  in favor of  $H_1$ 

#### Which pct to choose for CI?

Rejection with higher pct confidence interval gives stronger evidence

## 'Significance level' of hypothesis test

 $\alpha = \text{test significance level} = 1 - \text{pct used in CI}$ 

#### intuitively

Rejecting a hypothesis test with  $\alpha$  level means if we redid this experiment we would incorrectly reject  $H_0$  only  $\alpha$  percent of the time

'incorrectly reject' means our test tells us to reject  $H_0$  when in fact it is true

## **Example**

 $\alpha = 0.05$  level corresponds to 0.95 CI



# Revisit example

True mean and s.d. daily caloric intake for Ankara students are the unknown numbers m,s.

#### **Hypotheses**

$$m_0 = 3500$$

$$H_0: m=3500, \qquad H_1: m 
eq 3500$$

#### **Confidence interval pct = 0.95**

Assume we get the same standard deviation for Ankara students,  $\hat{s}_n = 500$ .

$$\mathrm{low} \, = 3450 - z^* rac{500}{\sqrt{100}}, \quad \mathrm{high} \, = 3450 + z^* rac{500}{\sqrt{100}}$$

rounding to 3 decimals

$$(low, high) = (2470.018, 4429.982)$$

Q: Reject  $H_0$ ?

Q: Significance level  $\alpha$ ?

Q: If you consider a CI with pct =0.98 would your answer change?

# p-values

#### **Definition**

"The p-value is the probability of observing data at least as favorable to the alternative hypothesis as our current data set, **if the null hypothesis is true.**"

-- textbook ch 5.3.4

#### **Translation**

$$H_0: m = m_0, \quad H_1: m \neq m_0$$

p-value is the probability of observing data that is at least

 $|\hat{m}_n-m_0|$  away from  $m_0$  as what we saw...

meaning chance that if we redrew our sample we would see a sample mean at least as different from  $m_0$  as a sample mean at least as different from  $m_0$  as a sample mean at least as different from  $m_0$  as a sample mean at least as different from  $m_0$  as a sample mean at least as different from  $m_0$  as a sample mean at least as different from  $m_0$  as a sample mean at least as different from  $m_0$  as a sample mean at least  $m_0$  and  $m_0$  are sample mean at least  $m_0$  as  $m_0$  as  $m_0$  as  $m_0$  and  $m_0$  are  $m_0$  as  $m_0$  and  $m_0$  are  $m_0$  as  $m_0$  as  $m_0$  and  $m_0$  are  $m_0$  as  $m_0$  as  $m_0$  and  $m_0$  are  $m_0$  are  $m_0$  as  $m_0$  and  $m_0$  are  $m_0$  are  $m_0$ . a sample mean at least as different from  $m_{
m 0}$  as our current estimate  $\hat{m}_n$ 

#### **Calculation**

'two-sided test'

$$H_1: m 
eq m_0$$

$$p=P\left(rac{|ar{X}_n-m_0|}{\hat{s}_n/\sqrt{n}}>rac{|\hat{m}_n-m_0|}{\hat{s}_n/\sqrt{n}}
ight)=2P\left(Z<-rac{|\hat{m}_n-m_0|}{\hat{s}_n/\sqrt{n}}
ight)$$

'one-sided test'

$$H_1 : m < m_0$$

$$p = P\left(rac{ar{X}_n - m_0}{\hat{s}_n/\sqrt{n}} < rac{\hat{m}_n - m_0}{\hat{s}_n/\sqrt{n}}
ight) = P\left(Z < rac{\hat{m}_n - m_0}{\hat{s}_n/\sqrt{n}}
ight)$$

Often you'll see cutoff value referred to as the 'z-statistic'. These are for the two-sided or one-sided tests:

$$rac{|\hat{m}_n - m_0|}{\hat{s}_n/\sqrt{n}}$$
 or  $rac{\hat{m}_n - m_0}{\hat{s}_n/\sqrt{n}}$ 

#### Accept-reject hypothesis test using p values

$$H_0: m=m_0, \quad H_1: m 
eq m_0 \ lpha = ext{ test significance level}$$

reject  $H_0$  if

$$p \leq \alpha$$

This setup holds for one-sided tests too.

## Flip it: Significance in terms of p-value

p-value is the largest significance level at which you could reject the null

# How small a significance level / p-value is good enough?

Very much depends on the context

choose smaller  $\alpha$  when you really don't want to reject incorrectly

**Higgs boson discovery:** 

$$lpha = 3 imes 10^{-7} pprox 1/3,500,000$$

Scientific American blog post

generic standards you will see

$$\alpha = 0.05, 0.1$$

But why? Should instead think carefully about what makes sense for the question.

#### **Example for poll**

It is suspected that patients treated with a certain drug have systolic blood pressures (SBP) that are lower on average than 115, which is the mean for people not treated.

A random sample of 20 patients is treated with the drug. Their sample average SBP is 112 and the sample s.d. is 10.

## PollEv.com/brendanbrown849

poll closes at

