STAT 88: Lecture 29

Contents

Section 9.3: Confidence Intervals: Method

Section 9.4: Confidence Intervals: Interpretation

Last time

A/B testing:

A/B testing is the shorthand for comparing the distributions of two random samples.

A = Control group; B = Treatment group.

It follows the same 5 steps for hypothesis testing:

- (a) H_0 : treatment has no effect on back pain.
- (b) H_A : treatment has an effect on back pain.
- (c) Test statistic X: # patient in the treatment group who had pain relief.

Under H_0 , any difference between treatment and control groups is due to the random assignment of elements to treatment and control, so X follows HG(N, G, n) where N=total number of patients; G=total number of patients who had pain relief; n=number of patients in the treatment group.

- (d) Find p-value.
- (e) Reject H_0 iff p-value $\leq 5\%$.

Type-I error: (From warm up in Lecture 28) The *type I error* is the probability of rejecting the null hypothesis H_0 given that it is true.

A population distribution has an SD of 20. You want to test if the population mean is equal to 50:

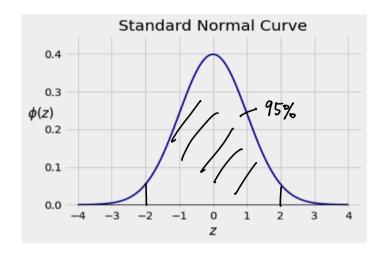
$$H_0: \mu = 50 \text{ vs } H_A: \mu < 50.$$

The average of a sample of 64 observations is \bar{x} .

- (a) Write down the expression for p-value.
- (b) Is p-value random or fixed? Why?
- (c) Suppose you reject H_0 if p-value is less than or equal to 5%. Find the region of \bar{x} where you reject H_0 .
- (d) Find the type-I error at 5% level, i.e. the probability of rejecting the null hypothesis H_0 given that it is true.

9.3. Confidence Intervals: Method

Preliminary The standard normal curve:



Confidence interval A confidence interval is an interval of estimates of a *fixed* but unknown parameter, based on data in a random sample.

Let X_1, \ldots, X_n be i.i.d. with mean μ and SD σ . We know \bar{X} is an unbised estimator of μ (i.e $E(\bar{X}) = \mu$), and $SD(\bar{X}) = \sigma/\sqrt{n}$ is a measure of the average spread of \bar{X} .

If n is large, the Central Limit Theorem tells us that the distribution of \bar{X} is roughly normal, so

$$P\left(-2 < \frac{\bar{X} - \mu}{\sigma/\sqrt{n}} < 2\right) \approx 0.95.$$

We rewrite this equation as follows:

$$P\left(\bar{X} - 2\frac{\sigma}{\sqrt{n}} < \mu < \bar{X} + 2\frac{\sigma}{\sqrt{n}}\right) \approx 0.95$$

$$\iff P\left(\mu \in \left(\bar{X} - 2\frac{\sigma}{\sqrt{n}}, \bar{X} + 2\frac{\sigma}{\sqrt{n}}\right)\right) \approx 0.95.$$

What is random and what is fixed?

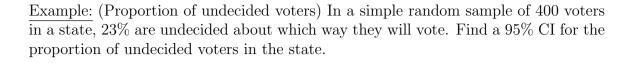
The *random* interval

$$\left(\bar{X} - 2\frac{\sigma}{\sqrt{n}}, \bar{X} + 2\frac{\sigma}{\sqrt{n}}\right)$$

is called an approximate 95% confidence interval for μ . It is a random interval because its endpoints depend on the sample mean \bar{X} which is a random variable whose value varies across samples.

Interpretation: the chance that this random interval contains the fixed parameter is about 95%.

Example: (From warm up in Lecture 28) A population distribution is known to have an SD of 20. The average of a sample of 64 observations is 55. What is your 95% confidence interval for the population mean?



Confidence Level

In above problem, find 99.7% confidence interval.

To find 90% confidence interval,

So 90% CI is
$$\left(\bar{X}- \quad \frac{\sigma}{\sqrt{n}}, \bar{X}+ \quad \frac{\sigma}{\sqrt{n}}\right).$$

9.4. Confidence Intervals: Interpretation

95% CI for μ :

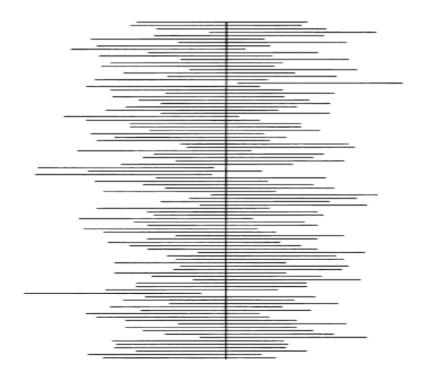
$$\left(\bar{X}-2\frac{\sigma}{\sqrt{n}},\bar{X}+2\frac{\sigma}{\sqrt{n}}\right).$$

It satisfies the property

$$P\left(\mu \in \left(\bar{X} - 2\frac{\sigma}{\sqrt{n}}, \bar{X} + 2\frac{\sigma}{\sqrt{n}}\right)\right) \approx 0.95.$$

The probability statement above is interpreted in terms of long run frequencies:

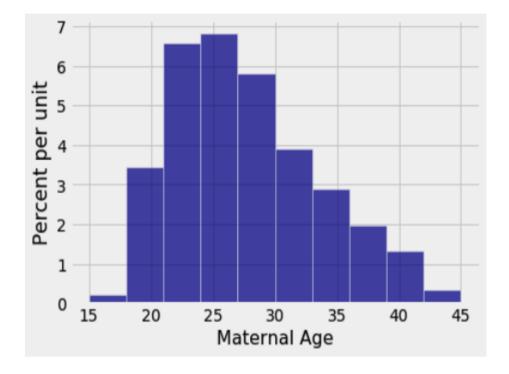
If you repeat the sampling process 100 times, and construct a 95% confidence interval each time, then about 95 of the 100 intervals will contain the parameter μ .



Ex: Suppose your observed instance of 95% CI is (79,82). What is the chance that $\mu \in (79,82)$?

Comparison with the Bootstrap The interpretation of CI is the same as in Data 8.

Example: Here is a distribution of 1174 maternal ages (years) from a random sample.



The sample mean is about 27.23 years and the sample SD is about 5.8 years. Find the approximate 95% CI of μ and interpret.

This works because \bar{X} is normally distributed by CLT. But if n is too small \bar{X} may not be normal and we have to bootstrap your 95% CL. How do you do this?

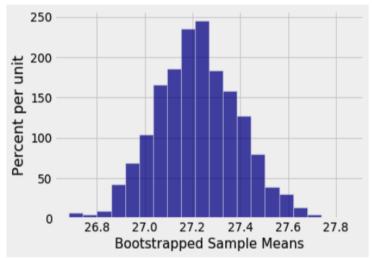
```
def one_resampled_mean():
    return np.average(births.sample().column('Maternal Age'))
```

We then called this function repeatedly to create an array of 2,000 bootstrap means:

```
means = make_array()

for i in np.arange(2000):
    means = np.append(means, one_resampled_mean())

Table().with_column('Bootstrapped Sample Means', means).hist(0, bins=2)
```



Finally, we found the "middle 95%" of the bootstrapped means. That was our empirical bootstrap 95% confidence interval for the population mean.

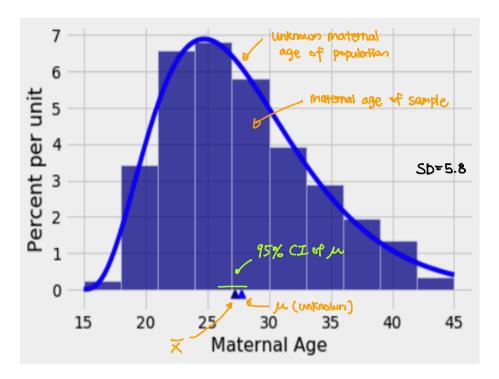
```
left = percentile(2.5, means)
right = percentile(97.5, means)
left, right

(26.89182282793867, 27.572402044293014)

(26.89182282793867, 27.572402044293014)
```

What the Confidence Interval Measures

CI is an interval of estimates of μ :



 \bar{X} is close to μ . On average it is $\mathrm{SD}(\bar{X}) = \frac{\sigma}{\sqrt{n}}$ away from μ . Is there a 95% chance that maternal ages are between (26.89, 27.57)?