Introduction to Inference

Adapted by Juan Pablo Lewinger from textbook: Diez et al. OpenIntro Statistics 6/17/2021

Estimating a population proportion

Example: 1,000 people poll shows US President approval rating is 52%

- 52% is an *estimate* of the true unknown approval rating among the entire adult US population, which is the parameter of interest.
- · Unless we ask and get answer from every adult in the US (close to impossible), the true approval rating will remain unknown.
- We denote true unknown approval rating parameter by p and its estimate (52% = 0.52) by \hat{p}
- The error due to sampling is the difference between the parameter and its estimate: $p \hat{p}$
- · If we took a different sample we'll get a different estimate and a different error
- · What can we say in general about the error when estimating a proportion?

- Assume the true approval rating is 54% (p = 0.54)
- · Simulate the adult population of the US: 210,000,000 aged 18 and above
- · Simulate a poll: draw a random sample of 1,000 individuals
- · Calculate the proportion in the sample

```
pop_size <- 210000000; poll_size = 1000

USpop <- c(rep("Approve", 0.54*pop_size), rep("Disapprove", 0.46*pop_size))

length(USpop)

## [1] 210000000

head(USpop); tail(USpop)

## [1] "Approve" "Approve" "Approve" "Approve" "Approve" "Approve"

## [1] "Disapprove" "Disapprove" "Disapprove" "Disapprove" "Disapprove"

## [6] "Disapprove"</pre>
```

```
set.seed(2021)
poll <- sample(USpop, poll_size)</pre>
head(poll)
## [1] "Approve"
                     "Approve"
                                   "Approve"
                                                 "Approve"
                                                               "Approve"
## [6] "Disapprove"
table(poll)
## poll
      Approve Disapprove
##
          551
                      449
p_hat <- sum(poll=="Approve")/poll_size</pre>
p_hat
## [1] 0.55
```

If we took a different sample we'd get a different estimate:

```
poll2 <- sample(USpop, poll_size)

p_hat2 <- sum(poll2=="Approve")/poll_size

p_hat2

## [1] 0.54

poll3 <- sample(USpop, poll_size)

p_hat3 <- sum(poll3=="Approve")/poll_size

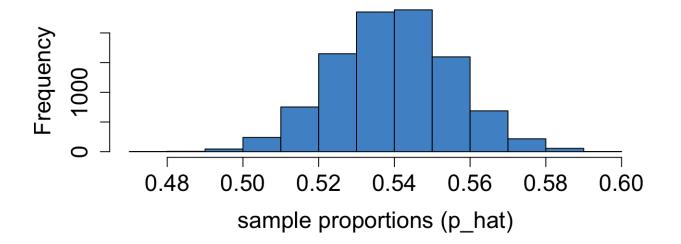
p_hat3

## [1] 0.53</pre>
```

• To get a sense of the distribution of possible values of \hat{p} let's repeat the simulation many times, say 10,000, and plot a histogram of the values of \hat{p} that we get:

```
sample_proportions <- replicate(10000, sum(sample(USpop, poll_size) == "Approve")/poll_size)
head(sample_proportions)
## [1] 0.54 0.56 0.55 0.54 0.56 0.53</pre>
```

```
hist(sample_proportions , col='steelblue3', xlab="sample proportions (p_hat)", main="", cex.lab=1.5, cex.axis=1.5)
```



```
c(mean = mean(sample_proportions), sd = sd(sample_proportions))
## mean sd
## 0.540 0.016
```

- · This is called the sampling distribution of the estimate \widehat{p}
- · The mean of the distribution, $\,\mu_{\widehat{p}}\,$, is 0.54, the same as the true population parameter!
- · This means that on average the population proportion estimates the true proportion without bias
- The spread (sd), called the standard error of \widehat{p} , and denoted $SE_{\widehat{p}}$ is quite small (0.02). This is the average error we make when we estimate the parameter p by its estimate \widehat{p} .
- When the true proportion is p=0.54 and the sample size n=1,000 the sample proportion tends to give a very good estimate of the population proportion
- The sampling distribution is symmetric and bell shaped, it looks like a normal distribution.

VERY IMPORTANT: The sampling distribution is never observed in real applications because we take a single sample of size n. Here we are simulating would would happen if we hypothetically took many, many samples of size n.

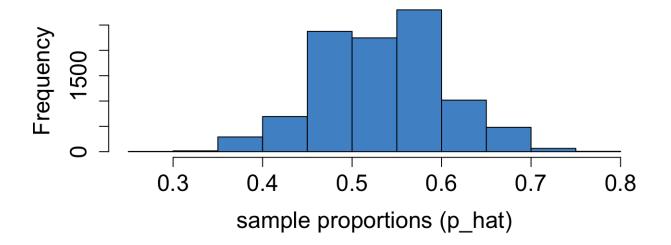
· How would the sampling distribution look like if the size of the sample was n=50 instead of n=1,000?

```
poll_size = 50

sample_proportions <- replicate(10000, sum(sample(USpop, poll_size) == "Approve")/poll_size)
head(sample_proportions)

## [1] 0.54 0.50 0.52 0.56 0.40 0.50</pre>
```

```
hist(sample_proportions , col='steelblue3', xlab="sample proportions (p_hat)", main="", cex.lab=1.5, cex.axis=1.5)
```



```
c(mean = mean(sample_proportions), sd = sd(sample_proportions))
## mean sd
## 0.54 0.07
```

- . The mean of the sampling distribution ($\mu_{\hat{p}}$) again equals the true population parameter p=0.54
- · However, the sandard error, $SE_{\widehat{p}}$, is now about 4 times larger! (0.071 vs 0.016)
- The larger the sample size the smaller the error (the error decreases proportionally to $\frac{1}{\sqrt(n)}$)

Central Limit Theorem (CLT)

When the sample size is sufficiently large, the sample proportion \hat{p} will tend to follow a normal distribution with the following mean and standard deviation:

$$\mu_{\hat{p}} = p$$

$$SE_{\hat{p}} = \sqrt{\frac{p(1-p)}{n}}$$

In order for the Central Limit Theorem to hold, the sample size is typically considered sufficiently large when:

$$np \ge 10$$

and

$$n(1-p) \ge 10$$

This is called the success-failure condition.

Central Limit Theorem (CLT)

In our scenario p = 0.54 so:

$$np = 1000 \times 0.54 = 540 > 10$$

and

$$n(1-p) = 1000 \times 0.46 = 460 > 10$$

So the Central Limit Theorem Holds with:

$$\mu_{\hat{p}} = 0.54$$

$$SE_{\hat{p}} = \sqrt{\frac{p(1-p)}{n}} = \sqrt{\frac{0.54 \times (1-0.46)}{1000}} = 0.016$$

These CLT-based calculations of the mean of the sample distribution and the standard error are consistent with our simulation results.

The CLT allows us to theoretically derive the standard error of sample distrubutions (no need to simulate)!

Confidence interval for a proportion

- When we estimate a proportion we can report the 'point estimate' \hat{p} along its standard error, which quantifies the uncertainty about the estimate.
- · A point estimate will never 'hit' the true parameter exactly
- So, we would like to provide a range of values, an interval, that contains the true parameter with high confidence
- · We build the confidence interval around the most plausible value, the sample proportion
- · When the central limit applies, the sampling distribution is close to a normal distribution
- · And normal distribution always has 95% of the data within 1.96 standard deviations of the mean
- So we construct a 95% confidence interval as:

Point Estimate $\pm 1.96 \times SE$

$$\hat{p} \pm 1.96 \times \sqrt{\frac{\hat{p}(1-\hat{p})}{n}}$$

Confidence interval for a proportion

In the presidential approval example (first poll):

$$\hat{p} = 0.55$$

$$SE_{\hat{p}} = \sqrt{\frac{p(1-p)}{n}} = \sqrt{\frac{0.55 \times (1-0.55)}{1000}} = 0.016$$

95% CI:
$$\hat{p} \pm 1.96 \times SE_{\hat{p}} = 0.55 \pm 1.96 \times 0.016 = (0.53, 0.57)$$

If the sample size was n = 50 instead:

$$SE_{\hat{p}} = \sqrt{\frac{p(1-p)}{n}} = \sqrt{\frac{0.55 \times (1-0.55)}{50}} = 0.07$$

95% CI:
$$\hat{p} \pm 1.96 \times SE_{\hat{p}} = 0.43 \pm 1.96 \times 0.07 = (0.48, 0.62)$$

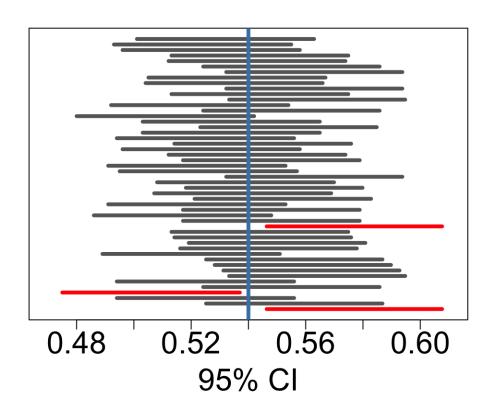
Interpretation of a 95% confidence interval

Suppose we took many samples of size n and built a 95% confidence interval from each. Then about 95% of those intervals would contain the parameter p

VERY IMPORTANT In a real application/analysis we draw a single sample of size n, compute a single point estimate, and a single confidence interval based on the sample. Here, using theory (CLT) and simulations, we are exploring what would happen if we hypothetically repeated the process of drawing a sample and computed estimates and CIs many times.

Confidence interval for a proportion

50 95% confidence intervals based on 50 different samples of size n=1,000



Your turn

- 1. Estimate the proportion of each shape of bead based on your sample
- 2. Compute the corresponding standard error for each estimate of proportion
- 3. Check whether the success-failure condition for the CLT holds.
- 4)Compute a corresponding 95% CI confidence interval (regardless of whether the CLT applies)

I'll reveal the true population proportions so that you can see how good your estimates and CIs are

Estimating a population mean

- · We estimate the population mean μ by the population sample \bar{x}
- The CLT also applies to the sample mean in the vast majority of cases:

When the sample size is sufficiently large, the sample mean \bar{x} will tend to follow a normal distribution with the following mean and standard deviation:

$$\mu_{\bar{x}} = \mu$$

$$SE_{\bar{x}} = \frac{\sigma}{\sqrt{n}}$$

where σ is the population standard deviation

Confidence interval for the mean

Point Estimate $\pm 1.96 \times SE$

$$\bar{x} \pm 1.96 \times \frac{\hat{\sigma}}{\sqrt{n}}$$

where $\,\widehat{\sigma}$ is the sample standard deviation, which estimates the population standard deviation $\,\sigma$

Your turn

- 1. Estimate the mean weight of the beads in the population
- 2. Compute the corresponding standard error
- 3. Compute the corresponding 95% CI
- 4. Estimate the mean weight of the rondelle beads, and corresponding SE and 95% CI

I'll reveal the true population mean weights so that you can see how good your estimates and CIs are.