## Inference for a Single Proportion

Nate Wells

Math 141, 4/8/22

Hypothesis Testing Procedures 0000000 Confidence Intervals 000000

## Outline

In this lecture, we will...

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- Use theory to find the standard error for one sample proportions
- Calculate confidence intervals and perform hypothesis tests for proportions using the theory-based method
- Investigate the results of the lacroix taste-test

## Section 1

# Inference for a Single Proportion

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## The Sampling Distribution for Sample Proportion

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  - Suppose each person in the sample has their own binary variable  $X_i$ . Then the sum  $X_1 + \cdots + X_n$  is the number of A's in the sample, and the mean of the  $X_i$  is the proportion of A's.

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- By the Central Limit Theorem, if *n* is large, then  $\hat{p}$  is approximately Normal, with mean *p* and standard deviation  $\sqrt{\frac{p(1-p)}{n}}$

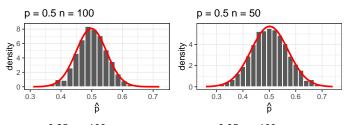
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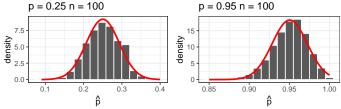
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### Examples

 Below are the sampling distributions for p̂ for a variety of values of p and n, along with the approximating Normal curve:





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- Why learn two methods?
  - The Theory-based method works best when modeling assumptions are true
  - Simulation-based methods can perform well in a variety of circumstances, but sometimes lack precision

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## Section 2

## Hypothesis Testing Procedures

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Hypothesis Testing Procedures

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### z-Scores

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By location-scale invariance,

$$P(X > x) = P\left(Z > \frac{x - \mu}{\sigma}\right)$$

 If we want to compute a P-Value for test statistic x, we can instead compute a P-value for its z-score z:

$$\begin{array}{rcl} \mathsf{P}\text{-value} &=& P(Z > z) & \text{if } H_a \text{ is one-sided right} \\ \mathsf{P}\text{-value} &=& P(Z < z) & \text{if } H_a \text{ is one-sided left} \\ \mathsf{P}\text{-value} &=& 2 \cdot P(Z > |z|) & \text{if } H_a \text{ is two-sided} \end{array}$$

### Hypothesis Tests

By the central limit theorem, if  $H_0: p = p_0$  is true, then for large n,  $\hat{p}$  is approximately Normal, with the standard error

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#### Theorem

To test  $H_0: p = p_0$  against  $H_a: p \neq p_0$  (or the one-sided alternative) we use the standardized test statistic

$$z = \frac{\hat{p} - p_0}{\sqrt{\frac{p_0(1-p_0)}{n}}}$$

If n is large enough so that both  $n\hat{p}$  and  $n(1-\hat{p})$  are at least 10, then the p-value for the test is computed using the standard Normal distribution.

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$$H_0: p = \frac{1}{3}$$
  $H_a: p > \frac{1}{3}$ 

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- That is, the observed  $\hat{p}$  was 2.5 standard errors above the mean.
- This seems unlikely to occur, if the null hypothesis were true (remember, 95% of all observations are within 2 standard errors of mean)

Confidence Intervals 000000

### Calculate P-Value

• If  $H_0$  is true, the z-score should be Normally distributed, with mean 0 and st. dev.

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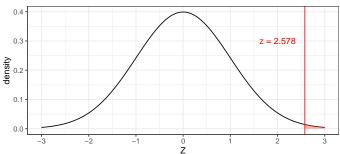
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Confidence Intervals 000000

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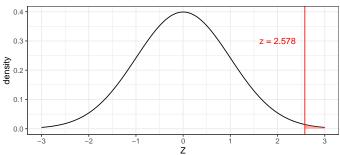


Distribution of z-scores

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Distribution of z-scores

The exact p-value is

1-pnorm(q=2.578, mean = 0, sd = 1)

## [1] 0.0049687

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```
set.seed(48)
lacroix %>% specify(response = correct, success = "yes") %>%
hypothesize(null = "point", p = 1/3) %>%
generate(reps = 5000, type = "simulate") %>%
calculate(stat = "prop") %>%
get_p_value(obs_stat = .5, direction = "right")
```

```
## # A tibble: 1 x 1
## p_value
## <dbl>
## 1 0.0038
```

# Section 3

# Confidence Intervals

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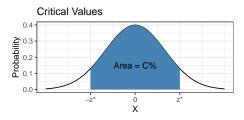
Confidence Intervals

# Critical Values

• The **critical value**  $z^*$  for a C% confidence interval is the value so that C% of area is between  $-z^*$  and  $z^*$  in the standard Normal distribution

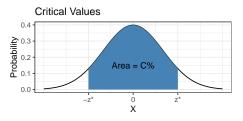
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- For Normal distributions, approximately 95% of observations are within 2 standard deviations of the mean.
  - So the critical value for 95% confidence is approximately

$$z^* = 2$$
 (exact value is  $z^* = 1.96$ )

Confidence Intervals

### **Confidence Intervals**

When a sample statistic is approximately Normally distribution, the C% confidence interval is

statistic  $\pm z^* \cdot SE$ 

where  $z^*$  is the critical value for C% confidence and SE is the standard error for the statistic.

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#### Theorem

Suppose an SRS of size n is collected from a population with parameter p. If n is large enough so that both  $n\hat{p}$  and  $n(1-\hat{p})$  are at least 10, then the confidence interval for p is

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Confidence Intervals

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Confidence Intervals

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qnorm(p = .95, mean = 0, sd = 1)

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• The standard error for  $\hat{p}$  is

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```
• How does this compare to the bootstrap method?
set.seed(84)
lacroix %>% specify(response = correct, success = "yes") %>%
generate(reps=5000, type = "bootstrap") %>%
```

```
calculate(stat = "prop") %>%
get_ci(level = .9, type = "percentile")
```

## # A tibble: 1 x 2
## lower\_ci upper\_ci
## <dbl> <dbl>
## 1 0.390 0.593

Confidence Intervals

# Reflections on Experiment Design

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  - Why did we ask students to identify which of 3 cups was different, rather than giving 2 unmarked cups (1 lemon, 1 lime) and asking students to identify which is lime?
  - Why would observing  $\hat{p} < 0.33$  be unlikely under both the null and alternative hypotheses?