6.1 and 6.3 Inference About Proportions

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Outline

- Formulas for Standard Errors
- Normal-based Inference for Proportions

Case Study: Hitchhiker Snails



hitchhiker snail

Case Study: Hitchhiker Snails

- A type of small snail (*Tornatellides boeningi*) is very widespread in Japan, and colonies of the snails that are genetically very similar have been found very far apart.
- How could the snails travel such long distances?
- Biologist Shinichiro Wada fed 174 live snails to birds, and found that 26 were excreted live out the other end. (The snails are apparently able to seal their shells shut to keep the digestive fluids from getting in).

Yong, E. "The Scatological Hitchhiker Snail," Discover, October 2011, 13.

Research Question

What proportion of "hitchhiker" snails survive when ingested by a bird?

Interval for Proportion

- 1. Calculate sample statistic
- 2. Find z* for desired level of confidence
- 3. Calculate standard error (see formula)
- 4. Use statistic $\pm z^* \times SE$
- 5. Interpret in context

From Last Time: Central Limit Theorem

For random samples with a *sufficiently large* sample size, the distribution of sample statistics for a mean or a proportion is approximately normal.

- For proportions, "sufficiently large" can be taken as at least 10 in each category.
- The larger the sample sizes, the more normal the distribution of the statistic will be.

Accuracy

- The accuracy of simulation methods depends on the number of simulations (more simulations \Rightarrow more accurate).
- The accuracy of the normal distribution depends on the sample size (larger sample size \Rightarrow more accurate).
- If the distribution of the statistic is normal and you have generated many simulated statistics, the answers should be very close.

Normal Confidence Interval Formula

sample statistic
$$\pm z^* \times SE$$

SE from bootstrap distribution.

Normal Formula for *p*-values

$$z = \frac{\text{sample statistic - null parameter}}{\text{SE}}$$

- Compare z to N(0, 1) to get p-value.
- · SE from randomization distribution.

Standard error

- The *classical approach* uses formulas rather than simulations to find standard errors.
- · Methods should agree when classical conditions are met.

Standard Error Formulas

Parameter	Distribution	Standard Error
Proportion	Normal	$\sqrt{\frac{p(1-p)}{n}}$
Difference in Proportions	Normal	$\sqrt{\frac{p_1(1-p_1)}{n_1} + \frac{p_2(1-p_2)}{n_2}}$
Mean	t, df= $n-1$	$\sqrt{\frac{\sigma^2}{n}}$
Difference in Means	t , df= min $(n_1, n_2) - 1$	$\sqrt{\frac{\sigma_1^2}{n_1} + \frac{\sigma_2^2}{n_2}}$

SE Formula Considerations

- *n* is always in the denominator (larger sample size gives smaller standard error).
- Standard error related to square root of 1/n.
- Standard error formulas use population parameters (unknown!)
 - For intervals, plug in the sample statistic(s) as your best guess at the parameter(s)
 - For tests, plug in the null value for the parameter(s), because you want the distribution assuming H_0 true (if the parameters have anything to do with H_0)

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Interval Calculation

- Study results: 26 of 174 snails survived
- Calculate a 90% confidence interval for p.

Conclusion

We are 90% confident that between 0.105 and 0.193 of hitchhiker snails survive being ingested by a bird.

Comparing Proportions: Bird Species Effect?

- The snails were fed to two different birds
- 14.3% of the 119 snails fed to Japanese white-eyes survived
- 16.4% of the 55 snails fed to brown-eared bulbuls survived

Japanese White-eye



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Brown-eared bulbul



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Interval Calculation

• Use the study results and the standard error formula to calculate a 95% confidence interval for the difference in proportions, $p_W - p_B$.

Case Study: Hormone Replacement Therapy

Does hormone replacement therapy cause increased risk of breast cancer?

Hormone Replacement Therapy

- Until 2002, hormone replacement therapy (HRT), estrogen and/or progesterone, was commonly prescribed to post-menopausal women.
- This changed in 2002, when the results of a large clinical trial were published
 - 8506 women were randomized to take HRT and 8102 were randomized to placebo.
 - 166 HRT and 124 placebo women developed any kind of cancer.

Hypothesis Test

- 1. State hypotheses
- 2. Calculate sample statistic
- 3. Calculate SE
- 4. Calculate z = (statistic null)/SE
- 5. Compare z to standard normal distribution to find p-value.
- 6. State a conclusion in context.

Hypotheses

Does hormone replacement therapy cause increased risk of invasive breast cancer?

- · Let p_1 = proportion of women taking HRT who get invasive breast cancer
- · Let $p_2 =$ proportion of women not taking HRT who get invasive breast cancer

What are the appropriate hypotheses here?

Calculate *z*-Statistic

Compute:

$$z = \frac{\text{sample statistic} - \text{null parameter}}{\text{SE}}$$

where

SE =
$$\sqrt{\frac{p_1(1-p_1)}{n_1} + \frac{p_2(1-p_2)}{n_2}}$$

- But what to use for p_1 and p_2 ?
- How to assume H_0 true?

Hypothesis Testing

- When testing, use the null value for p, rather than the sample statistic
- If testing whether the proportion of people not taking HRT who develop invasive breast cancer is less than 0.1, the standard error would be

SE =
$$\sqrt{\frac{p_0(1-p_0)}{n}}$$
 = $\sqrt{\frac{0.1(1-0.1)}{8102}}$ = 0.0033

where p_0 is the hypothesized value for p.

Null Values

- Testing a difference in proportions: $H_0: p_1 = p_2$
- · Use the overall sample proportion from both groups (called the *pooled proportion*) to estimate the common value $p_1 = p_2 = p$:

$$\hat{p} = \frac{124 + 166}{8102 + 8506} = 0.017$$

· Note that this is in between the sample proportions for each group.

Standard Error Calculations

Substitute the pooled \hat{p} into

SE =
$$\sqrt{\frac{p_1(1-p_1)}{n_1} + \frac{p_2(1-p_2)}{n_2}}$$

to get

SE =
$$\sqrt{\frac{\hat{p}(1-\hat{p})}{n_1} + \frac{\hat{p}(1-\hat{p})}{n_2}}$$

Calculations give us the SE for the hypothesis test:

$$SE = \sqrt{\frac{0.017(1 - 0.017)}{8506} + \frac{0.017(1 - 0.017)}{8102}} = 0.002$$

Conclusion

- If there were no difference between HRT and placebo regarding invasive breast cancer, we would only see results this extreme about 2 out of 100 times.
- We have evidence that HRT increases risk of invasive breast cancer.

Follow-up: The trial was terminated because of these findings, and hormone replacement therapy is now no longer routinely recommended.

More Practice

- · Same trial, different variable of interest.
 - 8506 women were randomized to take HRT, 8102 were randomized to placebo.
 - 502 HRT and 458 placebo women developed any kind of cancer.

Does hormone replacement therapy cause increased risk of cancer in general?

Inference formulas for proportions

Parameter of Interest	Confidence Interval	Test of Significance
p	$\hat{p} \pm z^* \sqrt{\frac{\hat{p}(1-\hat{p})}{n}}$	$z = \frac{\hat{p} - p_0}{\sqrt{\frac{p_0(1 - p_0)}{n}}}$
$p_1 - p_2$	$(\hat{p}_1 - \hat{p}_2) \pm z^* \sqrt{\frac{\hat{p}_1(1 - \hat{p}_1)}{n_1} + \frac{\hat{p}_2(1 - \hat{p}_2)}{n_2}}$	$z = \frac{\hat{p}_1 - \hat{p}_2}{\sqrt{\frac{\hat{p}(1-\hat{p})}{n_1} + \frac{\hat{p}(1-\hat{p})}{n_2}}}$

Margin of Error and Sample Sizes

Margin of Error

$$\hat{p} \pm z^* \times \sqrt{\frac{\hat{p}(1-\hat{p})}{n}}$$

· Question: What is the margin of error?

Margin of Error for Choosing Sample Sizes

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

- You can choose your sample size in advance, depending on your desired margin of error.
- Given this formula for margin of error, solve for *n* to get.

$$n = \left(\frac{z^*}{\text{MOE}}\right)^2 \hat{p}(1 - \hat{p})$$

• But what to use for \hat{p} ?

Margin of Error

$$n = \left(\frac{z^*}{\text{MOE}}\right)^2 \hat{p}(1 - \hat{p})$$

- Neither p nor \hat{p} is known in advance. To be conservative, use $p^*=0.5$.
- For a 95% confidence interval, $z^* \approx 2$ so

$$n \approx \left(\frac{1}{\text{MOE}}\right)^2$$

Margin of Error Example

Suppose we want to estimate a proportion with a margin of error of 0.03 with 95% confidence.

How large a sample size do we need?