Hypothesis Testing - Categorical Data (part 2)

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Introduction

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 - Generally speaking, these approaches fell into one of three camps: simulation, exact probability calculations, Normal/Chi-square distributions

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- Previously, we discussed a variety of different hypothesis testing approaches for scenarios involving one categorical variable
 - Generally speaking, these approaches fell into one of three camps: simulation, exact probability calculations, Normal/Chi-square distributions
- ► This presentation will cover hypothesis testing approaches for scenarios involve *two categorical variables*
 - ► The most common example of this is a categorical response variable and a categorical explanatory variable that is used to distinguish treatment and control groups

Surgical Site Infections

- ► In the 1860's, surgeries often led to infections that resulted in death
- ► At the time, many experts believed the infections were due to "bad air"
 - Many hospitals had policies that required their wards open their windows at midday to air out

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- ► At the time, many experts believed the infections were due to "bad air"
 - Many hospitals had policies that required their wards open their windows at midday to air out
- ▶ It was customary for surgeons to move quickly from patient to patient with out any sort of special precautions
 - In fact, many took pride the accumulated stains on their surgical gowns as a measure of experience

Louis Pasteur

- In 1862, Louis Pasteur discovered that food spoilage was caused by the growth and proliferation of harmful micro-organisms
- ► Pasteur identified three methods for eliminating these micro-organisms: heat, filtration, and chemical disinfectants
 - The method of heating became known as pasteurization (named for Pasteur) and is widely applied to milk, beer, and many other food products

Joseph Lister

- Joseph Lister, a Professor of Surgery at the Glasgow Royal Infirmary, became aware of Pasteur's work and theorized that it might explain the infections that frequently occurred after surgery
 - ► How do you think Lister evaluated this theory?

Lister's Experiment

- ► Lister proposed a new protocol where surgeons were required to wash their hands, wear clean gloves, and disinfect their instruments with a carbolic acid solution
 - Lister randomly assigned 75 patients to receive either his new "sterile" surgery or the old standard of care
 - Lister than recorded how many of each group survived until their discharge from the hospital

	Died	Survived
Control	16	19
Sterile	6	34

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- Bias? Probably not, while double-blinding wasn't possible, it's unlikely there was biased measurement of the outcome (survival) or that this was a non-representative group of patients (sampling bias)
- 2) Confounding variables? No, we'd expect them to be balanced in the two groups due to random assignment
- 3) Random chance? ... We need to use a statistical test

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 - ▶ In words, what would your null model be for Lister's experiment?
- ▶ The null model is that the Lister's proposed sterilization procedure makes no difference
 - That is, equal proportions of the "Sterile" and "Control" groups are expected to die prior to discharge

$$H_0: p_1-p_2=0$$

 \triangleright Where p_1 denotes the proportion of deaths among the "Control" condition, and p_2 is the proportion of deaths among the "Sterile" condition



Simulating the Null Distribution

- ▶ If sterilization made no difference, the deaths observed in this study occurred at random in the two groups
 - ► Thus, under the null model, we can assume the overall death rate (estimated at 22 of 75, or 29%) applies to both groups
 - We can simulate the outcomes in both groups using sets of $n_1=35$ and $n_2=40$ "weighted coin-flips" with a 29% chance of death

Simulating the Null Distribution

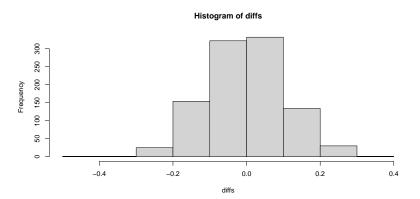
```
## Set seed (for replication purposes)
set.seed(123)

## Simulate deaths for the control group
control_deaths <- rbinom(1000, 35, .29)

## Simulate deaths for the sterile group
sterile_deaths <- rbinom(1000, 40, .29)</pre>
```

Null distribution

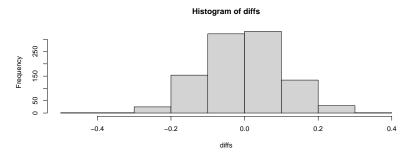
Calculate and graph the possible differences in proporte
diffs <- control_deaths/35 - sterile_deaths/40
hist(diffs)</pre>





Finding the *p*-value

▶ Recall that we observed $\hat{p}_1 = 16/35$ and $\hat{p}_2 = 6/40$, resulting in a sample difference in proportions of 0.307



▶ What would you estimate the two-sided *p*-value to be?



Finding the *p*-value

```
## Find the p-value
upper <- sum(diffs >= 0.307)/1000
2*upper
```

```
## [1] 0
```

Zero of 1000 simulations were as extreme as the actual experimental results. Thus, Lister's experiment provides *strong evidence* that the proposed sterilization protocol improves survival

In our study of confidence intervals, we learned that Central Limit Theorem (combined with some probability rules for independent random variables) leads to the following distributional result:

$$\hat{p}_1 - \hat{p}_2 \sim N\left(p_1 - p_2, \sqrt{\frac{p_1(1-p_1)}{n_1} + \frac{p_2(1-p_2)}{n_2}}\right)$$

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- $\hat{p}_1=6/40=0.15$ and $\hat{p}_2=16/35=0.46$ won't work, they don't satisfy the null hypothesis
- ▶ Instead, we must use a pooled proportion, or 22/75 = 0.29, in place of both p_1 and p_2



[1] 0.00326354

```
## Calculate the Z-value
pool <- 22/75
se <- sqrt(pool*(1-pool)/35 + pool*(1-pool)/40)
z <- ((0.46 - .15) - 0)/se

## Find the p-value
upper <- pnorm(z, lower.tail = FALSE)
2*upper</pre>
```



Chi-Squared Test for Differences in Proportions

A third way to approach hypothesis testing in this scenario involves observed vs. expected frequencies, recall we observed:

	Died	Survived
Control	16	19
Sterile	6	34

If the sterilization procedure made no difference, we'd expect 29% of each group to have died, leading to the following table of expected counts:

	Died	Survived
Control	35*.29 = 10.2	35*.71 = 24.9
Sterile	40*.29 = 11.6	40*.71 = 28.4

Chi-Squared Test for Differences in Proportions

Comparing these observed and expected frequencies, we can set up a Chi-squared test:

$$X^{2} = \frac{(16-10.2)^{2}}{10.2} + \frac{(19-24.9)^{2}}{24.9} + \frac{(6-11.6)^{2}}{11.6} + \frac{(34-28.4)^{2}}{28.4} = 8.5$$

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In a 2x2 table with fixed margins, if you know one of the four cells you can calculate the rest, meaning df = 1:

```
## p-value from Chi-squared df =1
pchisq(8.5, df = 1, lower.tail = FALSE)
```

```
## [1] 0.003551465
```

```
## Using chisq.test
tab <- data.frame(Died = c(16,6), Survived = c(19,34))
chisq.test(tab, correct = FALSE)$p.value</pre>
```

```
## [1] 0.003560924
```



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- Chi-squared testing in this context is known as testing for association or testing for independence
- ► Chi-squared tests can be performed on sample data in any two-way frequency table, not just 2x2 tables
 - For these tests, df = (I 1) * (J 1), where I is the number of rows and J is the number of columns in the table
- We calculated expected counts using a pooled proportion that collapsed the table's rows
 - We could have done something similar using the table's columns and arrived at an identical result
 - Alternatively, many places teach the formula $E_{ij} = \frac{\text{Observed}_{ij}}{\text{Row Total}_i * \text{Column Total}_j}$



Comments on Chi-Squared Tests (continued)

- ► As a final comment, recall the Chi-squared distribution shares a connection with the standard normal distribution and Z-test
 - ► This makes the test a large-sample result which is only accurate when every cell has an expected frequency of at least 5
- ► Fortunately, an exact test for independence in two-way tables exists, which we can use in these small-sample situations

Fisher's Exact Test

- ► Two-way frequency tables are used so frequently that Ronald Fisher derived an exact test for association in them
- We will not get into the details, but the gist is that if you fix the row and column totals, the remaining numbers that are free to vary will follow a hypergeometric distribution under the null hypothesis of independence

```
## Using fisher.test
tab <- data.frame(Died = c(16,6), Survived = c(19,34))
fisher.test(tab)$p.value</pre>
```

```
## [1] 0.005018047
```

Statistical vs. Clinical Significance

- ► The χ^2 test for independence, and Fisher's exact test can both be used to evaluate the strength of an association that exists between two categorical variables
 - ▶ The lower the *p*-value, the more strongly the variables are associated (That is, the more incompatible the data are with the variables being independent)

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- ▶ These methods do not tell us anything about the nature of the association
 - We could report the sample difference in proportions (accompanied by a confidence interval), but this summary measure has a major shortcoming
- Consider the proportions of smokers and non-smokers that develop lung cancer in a 10-year period
 - ► These proportions are estimated at 0.00438 and 0.00045 respectively, or a difference of 0.0039 (far less than 1%)



Odds Ratios

- ► A commonly reported measure of association describing the relationship between two categorical variables is the **odds ratio**
 - ► The odds of an event is the ratio of how often it happens to how often it doesn't happen
 - ▶ If a team has a 75% probability of winning a game, the odds of winning are 3, which is often spoken as "3 to 1"

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- In our smoking example, the odds of a smoker developing lung cancer are $\frac{0.00438}{1-0.00438} = 0.00440$
 - Similarly, the odds of a non-smoker developing lung cancer are $\frac{0.00045}{1-0.00045} = 0.00045$

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 - Similarly, the odds of a non-smoker developing lung cancer are $\frac{0.00045}{1-0.00045} = 0.00045$
- ► Thus, the *odds ratio* is $\frac{0.00440}{0.00045} = 9.8$
 - ► We say that the odds of a smoker developing lung cancer are 9.8 times those of a non-smoker developing lung cancer



Confidence Interval for an Odds Ratio in R.

```
## 95% CI for an OR (Lister's Experiment)
tab <- data.frame(Died = c(16,6), Survived = c(19,34))
fisher.test(tab, conf.int = TRUE,
            conf.level = .95)$conf.int[1:2]
```

```
## [1] 1.437621 17.166416
```

Thus, we can conclude with 95% the odds of death in the Control group are between 1.4 and 17.2 times higher than the odds of death in the Sterile group

Summary

- ► This presentation covered several hypothesis testing approaches for evaluating relationships between two categorical variables
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 - Fisher's exact test
- ▶ In general, Fisher's exact test is the most robust of these methods, but it is very computationally intensive for large tables
 - ightharpoonup Because of this, χ^2 tests tend to be most widely used in practice
- ▶ We also introduced the **odds ratio** as a measure of association that can be used to gauge clinical vs. statistical significance

