

# Inference on Two Sample Proportions in R

BIOST 514/517

Discussion - Week 6

# Proportions

Interest in examining a binary variable in a population

- ▶ Death
- ▶ Cancer relapse

```
psa <- read.table("../psa.txt",header=TRUE)

dat <- psa[!is.na(psa$grade),]

relapse24 <- dat$inrem=="no" & dat$obstime < 24
hiGrade <- dat$grade > 2
```

# Inference on Two Proportions

May want to compare proportions between two groups by their difference  $p_1 - p_2$

- ▶ Treatment versus control
- ▶ Low grade versus high grade tumor

## Difference in Sample Proportions

Use  $\hat{p}_1 - \hat{p}_2 = \frac{X_1}{n_1} - \frac{X_2}{n_2}$  to estimate  $p_1 - p_2$

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

# Testing the Difference in Proportions

Test the null hypothesis  $p_1 - p_2 = 0$  against  $p_1 - p_2 \neq 0$

- ▶ Under the null  $p_1 = p_2 = p$ , so the standard deviation is

$$\sqrt{p(1-p) \left( \frac{1}{n_1} + \frac{1}{n_2} \right)}$$

- ▶ We estimate  $p$  with  $\hat{p}$  the pooled sample proportion

$$\hat{p} = \frac{X_1 + X_2}{n_1 + n_2}$$

- ▶ Then we can use a  $Z$ -test with

$$Z = \frac{\hat{p}_1 - \hat{p}_2}{\sqrt{\hat{p}(1-\hat{p}) \left( \frac{1}{n_1} + \frac{1}{n_2} \right)}}$$

## Confidence Intervals for the Difference in Proportions

As in the one-sample case, we do not assume the null hypothesis for the standard deviation, and we use our best guess for  $p_1$  and  $p_2$

$$(\hat{p}_1 - \hat{p}_2) \pm z_{1-\alpha/2} \sqrt{\frac{\hat{p}_1(1 - \hat{p}_1)}{n_1} + \frac{\hat{p}_2(1 - \hat{p}_2)}{n_2}}$$

## prop.test for Difference in Proportions

Can still use `prop.test(x, n)`, with defaults

- ▶ Testing  $p_1 - p_2 = 0$  against  $p_1 - p_2 \neq 0$
- ▶ Continuity correction
- ▶ 95% confidence with argument `conf.level=0.95`

## prop.test for Difference in Proportions

```
diffGrade <- (xHigrade/nHigrade)-  
             (xLograde/nLograde)
```

We estimate the proportion experiencing relapse within 24 months is 0.04 higher in a group with high grade tumors than in a group with low grade tumors based on our data.



## prop.test for Difference in Proportions

```
diffTestCorrect <- prop.test(x=c(xLoggrade,xHigrade),  
                             n=c(nLoggrade,nHigrade))  
diffTestCorrect
```

```
##
```

```
## 2-sample test for equality of proportions with continuity
```

```
## correction
```

```
##
```

```
## data:  c(xLoggrade, xHigrade) out of c(nLoggrade, nHigrade)
```

```
## X-squared = 5.8362e-31, df = 1, p-value = 1
```

```
## alternative hypothesis: two.sided
```

```
## 95 percent confidence interval:
```

```
## -0.3847789  0.3097789
```

```
## sample estimates:
```

```
## prop 1 prop 2
```

```
## 0.4000 0.4375
```

## prop.test for Difference in Proportions

```
diffTestCorrect$conf.int
```

```
## [1] -0.3847789  0.3097789
```

```
## attr(,"conf.level")
```

```
## [1] 0.95
```

Our data are consistent with the proportion relapsing within 24 months in a high grade tumor group being 0.38 lower to 0.31 higher than in a low grade tumor group. Because 0 is in our confidence interval, we would not be surprised if the true proportions were similar between groups.

# Odds

A different way of looking at probabilities

$$o = \frac{p}{1-p} \implies \hat{o} = \frac{\hat{p}}{1-\hat{p}}$$

```
oLograde <- (xLograde/nLograde)/  
  ((nLograde-xLograde)/nLograde)  
oHigrade <- (xHigrade/nHigrade)/  
  ((nHigrade-xHigrade)/nHigrade)  
oLograde
```

```
## [1] 0.6666667
```

```
oHigrade
```

```
## [1] 0.7777778
```

## Odds Ratio

Compare the relative difference in odds instead of absolute difference in proportions

$$OR = \frac{o_1}{o_2}$$

```
orGrade <- oHigrade/oLograde  
orGrade
```

```
## [1] 1.166667
```

We estimate the odds of relapse within 24 months for a group with high grade tumors are 1.17 times the odds for a group with low grade tumors based on our data.

Alternatively, we can say that the odds of relapse within 24 months for a group with high grade tumors are  $(\widehat{OR} - 1) \cdot 100\% \approx 16.67\%$  higher than for a group with low grade tumors based on our data.

## Odds Ratio from a 2x2 Table

	$E = 0$	$E = 1$
$D = 0$	a	b
$D = 1$	c	d

$$\widehat{OR} = \frac{ad}{bc}$$

```
tabGrade <- table(relapse24,hiGrade)
tabGrade
```

```
##           hiGrade
## relapse24 FALSE TRUE
##      FALSE   15   9
##      TRUE    10   7
```

```
(15*7)/(9*10)
```

```
## [1] 1.166667
```

## Inference on the Odds Ratio

$$\log \widehat{OR} \sim N \left( \log OR, \sqrt{\frac{1}{a} + \frac{1}{b} + \frac{1}{c} + \frac{1}{d}} \right)$$

## 95% Confidence Interval for the Odds Ratio

$$\exp \left( \log \widehat{OR} \pm \sqrt{\frac{1}{a} + \frac{1}{b} + \frac{1}{c} + \frac{1}{d}} \right)$$

```
orSE <- sqrt((1/15)+(1/9)+(1/10)+(1/7))  
exp(log(orGrade) + c(-1,1)*qnorm(0.975)*orSE)
```

```
## [1] 0.3272565 4.1591563
```

## fisher.test for Inference on the Odds Ratio in R

`fisher.test` function includes CI for the odds ratio

- ▶ Default to 95% confidence interval with argument `conf.level=0.95`
- ▶ Estimates slightly different than “by hand”



## Confidence Interval for the Odds Ratio in R

```
infGrade <- fisher.test(tabGrade)
infGrade$conf.int
```

```
## [1] 0.2695186 4.9506212
## attr(,"conf.level")
## [1] 0.95
```

These data are consistent with the odds of relapse in 24 months in a high grade tumor group being 0.27 to 4.95 times those of a low grade tumor group. Because 1 is in our confidence interval, it would not be surprising if the true odds of relapse were similar between groups.