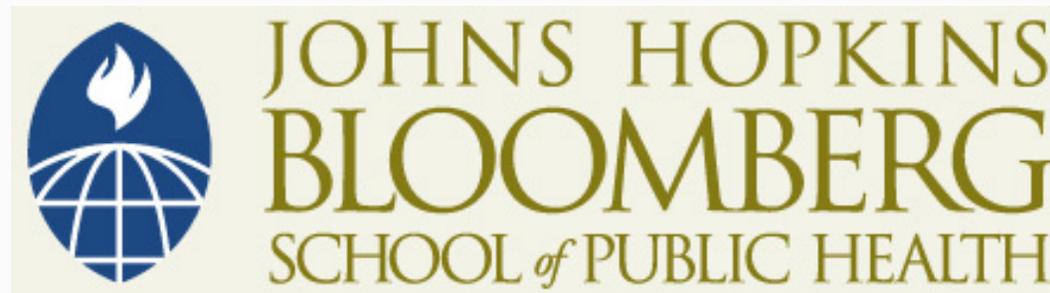


This work is licensed under a [Creative Commons Attribution-NonCommercial-ShareAlike License](https://creativecommons.org/licenses/by-nc-sa/4.0/). Your use of this material constitutes acceptance of that license and the conditions of use of materials on this site.



Copyright 2009, The Johns Hopkins University and John McGready. All rights reserved. Use of these materials permitted only in accordance with license rights granted. Materials provided "AS IS"; no representations or warranties provided. User assumes all responsibility for use, and all liability related thereto, and must independently review all materials for accuracy and efficacy. May contain materials owned by others. User is responsible for obtaining permissions for use from third parties as needed.



JOHNS HOPKINS
BLOOMBERG
SCHOOL *of* PUBLIC HEALTH

Section E

Fisher's Exact Test

Recall: HIV Transmission/AZT Example

- Recall 2X2 (contingency) table

		Drug Group		
		AZT	Placebo	
HIV Transmission	Yes	13	40	53
	No	167	143	310
		180	183	363

Hypothesis Testing Problem: AZT and HIV Transmission

- Testing equality of two population proportions using data from two samples
 - $H_o: p_1 = p_2$ $H_o: p_1 - p_2 = 0$
 - $H_a: p_1 \neq p_2$ $H_A: p_1 - p_2 \neq 0$
 - In the context of the 2x2 table, this is testing whether there is a relationship between the rows (HIV status) and columns (treatment type)

Statistical Test Procedures

- (Pearson's) Chi-Square Test (χ^2)/Two-sample z-test
 - Both based on central limit theorem “kicking in”
 - Both results are “approximate,” but are excellent approximations if sample sizes are large
 - These do not perform so well in smaller samples

Statistical Test Procedures

- Fisher's Exact Test
 - Calculations are difficult
 - Always appropriate to test equality of two proportions
 - Computers are usually used
 - Exact p-value (no approximations)—no minimum sample size requirements

Fisher's Exact Test: HIV Transmission/AZT

- Rationale
 - Suppose H_0 is true—no association between AZT and maternal/infant HIV transmission
 - Imagine putting 53 red balls (the infected) and 310 blue balls (non-infected) in a jar
 - Shake it up

Fisher's Exact Test

- Now choose 180 balls (that's AZT group)
 - The remaining balls are the placebo group
- We calculate the probability you get 13 or fewer red balls among the 180
 - That is the one-sided p-value
- The two-sided p-value is just about (but not exactly) twice the one-sided
- p-value
 - It accounts for the probability of getting either extremely few red balls or a lot of red balls in the AZT group
 - The p-value is the probability of obtaining a result as or more extreme (more imbalance) than you did by chance alone

Using Stata: AZT/HIV Example

- Results from *csi* command, with *exact* option

```
. csi 13 40 167 143, exact
```

	Exposed	Unexposed	Total
Cases	13	40	53
Noncases	167	143	310
Total	180	183	363
Risk	.0722222	.2185792	.1460055
	Point estimate		[95% Conf. Interval]
Risk difference	-.146357		-.2171766 -.0755374
Risk ratio	.3304167		.1829884 .5966235
Prev. frac. ex.	.6695833		.4033765 .8170116
Prev. frac. pop	.3320248		

1-sided Fisher's exact P = 0.0001
2-sided Fisher's exact P = 0.0001

Small Sample Application

- Sixty-five pregnant women, all who were classified as having a high risk of pregnancy induced hypertension, were recruited to participate in a study of the effects of aspirin on hypertension*
- The women were randomized to receive either 100 mg of aspirin daily, or a placebo during the third trimester of pregnancy

Notes: *Schiff, E. et al. The use of aspirin to prevent pregnancy-induced hypertension and lower the ratio of thromboxane A2 to prostacyclin in relatively high risk pregnancies. *New England Journal of Medicine*, 321, 6.

Display Data in a 2x2 Table

- Results

		Group		
		Aspirin	Placebo	
Hypertension	Yes	4	11	15
	No	30	20	50
		34	31	65

Display Data in a 2x2 Table

- Sample proportion of subjects with hypertension

$$\hat{p}_{aspirin} = \frac{4}{34} = .12$$

$$\hat{p}_{placebo} = \frac{11}{31} = .35$$

Smaller Sample

- In this example . . . (just FYI)

$$n_{aspirin} * \hat{p}_{aspirin} * (1 - \hat{p}_{aspirin}) = 34 * .12 * .88 = 3.6$$

$$n_{placebo} * \hat{p}_{placebo} * (1 - \hat{p}_{placebo}) = 31 * .35 * .65 = 7.1$$

Fishers Exact

- Results from *csi* command, with *exact* option

```
. csi 4 11 30 20, exact
```

	Exposed	Unexposed	Total
Cases	4	11	15
Noncases	30	20	50
Total	34	31	65
Risk	.1176471	.3548387	.2307692
	Point estimate		[95% Conf. Interval]
Risk difference	-.2371917		-.4374335 -.0369498
Risk ratio	.3315508		.1176925 .9340096
Prev. frac. ex.	.6684492		.0659904 .8823075
Prev. frac. pop	.3496503		

1-sided Fisher's exact P = 0.0236
2-sided Fisher's exact P = 0.0378

Chi Square

- Results from *csi* command, without *exact* option

```
. csi 4 11 30 20
```

	Exposed	Unexposed	Total
Cases	4	11	15
Noncases	30	20	50
Total	34	31	65
Risk	.1176471	.3548387	.2307692
	Point estimate		[95% Conf. Interval]
Risk difference	-.2371917		-.4374335 -.0369498
Risk ratio	.3315508		.1176925 .9340096
Prev. frac. ex.	.6684492		.0659904 .8823075
Prev. frac. pop	.3496503		

chi2(1) = 5.14 Pr>chi2 = 0.0234

Fishers Exact

- 95% CI: not quite correct in smaller samples, but “good enough”

```
. csi 4 11 30 20, exact
```

	Exposed	Unexposed	Total
Cases	4	11	15
Noncases	30	20	50
Total	34	31	65
Risk	.1176471	.3548387	.2307692
	Point estimate	[95% Conf. Interval]	
Risk difference	-.2371917	-.4374335	-.0369498
Risk ratio	.3315508	.1176925	.9340096
Prev. frac. ex.	.6684492	.0659904	.8823075
Prev. frac. pop	.3496503		

```
1-sided Fisher's exact P = 0.0236
2-sided Fisher's exact P = 0.0378
```

Comparing Proportions between Independent Populations

- To get a p-value for testing:
 - $H_0: p_1 = p_2$
 - $H_A: p_1 \neq p_2$
- Two sample z-test or chi-squared test (give same p-value): work better in “bigger” samples and will match results of Fishers Exact Test
- Fisher’s exact test: always appropriate

Comparing Proportions between Independent Populations

- To create a 95% confidence interval for the difference in two proportions:

$$\hat{p}_1 - \hat{p}_2 \pm 2SE(\hat{p}_1 - \hat{p}_2)$$

- Fine for “bigger samples,” can be used as a “guideline” in smaller samples
- Not quite correct in “smaller samples” but will give you a good sense of width/range of CI