

Categorical Variables and Contingency Tables: Description and Inference

STAT 526
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Reading:
Agresti Ch. 1, 2 and 3
Faraway Ch. 4

Univariate Binomial and Multinomial Measurements

Binomial Distribution

- Probability distribution:

- $Y_1, Y_2, \dots, Y_n \stackrel{iid}{\sim} \text{Bernoulli}(\pi)$

- $\sum_{i=1}^n Y_i \sim \text{Binomial}(n, \pi)$

- $p(y) = \binom{n}{y} \pi^y (1 - \pi)^{n-y}$

- $\mu = E(Y) = n\pi, \sigma^2 = \text{var}(Y) = n\pi(1 - \pi)$

- Log-likelihood:

- $L(\pi) = y \log(\pi) + (n - y) \log(1 - \pi)$

- Maximum Likelihood Estimator:

- $\hat{\pi} = y/n$

- $E(\hat{\pi}) = \pi, SE(\hat{\pi}) = \sqrt{\frac{\pi(1-\pi)}{n}}$

Large-sample tests for π

- For a known π_0 , test
 $H_0 : \pi = \pi_0$ vs $H_0 : \pi \neq \pi_0$

- Wald test:

$$z_W = \frac{\hat{\pi} - \pi_0}{SE} = \frac{\hat{\pi} - \pi_0}{\sqrt{\hat{\pi}(1 - \hat{\pi})/n}} \stackrel{H_0, approx}{\sim} \mathcal{N}(0, 1)$$

- Likelihood ratio Test:

$$z_L = 2(L_1 - L_0) = 2 \left(y \log \frac{\hat{\pi}}{\pi_0} + (n - y) \log \frac{1 - \hat{\pi}}{1 - \pi_0} \right) \stackrel{H_0, approx}{\sim} \chi_1^2$$

- Score Test:

$$z_S = \frac{\hat{\pi} - \pi_0}{SE_0} = \frac{\hat{\pi} - \pi_0}{\sqrt{\pi_0(1 - \pi_0)/n}} \stackrel{H_0, approx}{\sim} \mathcal{N}(0, 1)$$

Closer to $\mathcal{N}(0, 1)$ than Wald

Large-sample CI for π

- Based on the Wald test statistic:

$$\hat{\pi} \pm z_{\alpha/2} \sqrt{\frac{\hat{\pi}(1 - \hat{\pi})}{n}}$$

Performs poorly unless large n

- Based on the Score Test statistic:

$$\hat{\pi} \left(\frac{n}{n + z_{\alpha/2}^2} \right) + \frac{1}{2} \left(\frac{z_{\alpha/2}^2}{n + z_{\alpha/2}^2} \right) \pm$$

$$z_{\alpha/2} \sqrt{\frac{1}{n + z_{\alpha/2}^2} \left[\hat{\pi}(1 - \hat{\pi}) \left(\frac{n}{n + z_{\alpha/2}^2} \right) + \frac{1}{2} \cdot \frac{1}{2} \left(\frac{z_{\alpha/2}^2}{n + z_{\alpha/2}^2} \right) \right]}$$

Performs better than Wald

Multinomial Distribution

- Probability distribution:

- $(Y_{i1}, \dots, Y_{ic}) \sim \{Y_{ij} = 1 \text{ if in category } j, \text{ and } 0 \text{ otherwise} \}$

- $\sum_{i=1}^n Y_{ij} \sim \text{Multinomial}(\pi_1, \dots, \pi_c), n = \sum_{j=1}^c n_j$

- $p(n_1, n_2, \dots, n_{c-1}) = \left(\frac{n!}{n_1! n_2! \dots n_c!} \right) \pi_1^{n_1} \pi_2^{n_2} \dots \pi_c^{n_c}$

- $E(n_j) = n\pi_j$
 $var(n_j) = n\pi_j(1 - \pi_j), cov(n_j, n_k) = -n\pi_j\pi_k$

- Log-likelihood:

- $L(\pi) = \sum_{j=1}^c n_j \cdot \log \pi_j$

- Maximum Likelihood Estimator:

- $\hat{\pi}_j = n_j/n$

Large-Sample Test for (π_1, \dots, π_c)

- For known $(\pi_{10}, \pi_{20}, \dots, \pi_{c0})$, test
 $H_0 : \pi_j = \pi_{j0} \text{ vs } H_0 : \pi_j \neq \pi_{j0}$

- Pearson test:

$$X^2 = \sum_{j=1}^c \frac{(O_j - E_{j0})^2}{E_{j0}} = \sum_{j=1}^c \frac{(n_j - n\pi_{j0})^2}{n\pi_{j0}} \stackrel{H_0, \text{ approx}}{\sim} \chi_{c-1}^2$$

E.g. in genetics: test theories of trait inheritance

- Likelihood Ratio test:

$$G^2 = 2(L_1 - L_0) = 2 \sum_{j=1}^c \log\left(\frac{n_j}{n\pi_{j0}}\right) \stackrel{H_0, \text{ approx}}{\sim} \chi_{c-1}^2$$

- Asymptotically equivalent when H_0 is true.
- For $n/c < 5$, X^2 converges faster

Poisson Distribution

- Probability distribution:
 - Y - number of events in a fixed interval of space/time
 - $Y \sim \text{Poisson}(\mu)$
 - $p(y) = \frac{e^{-\mu} \mu^y}{y!}$, $y = 0, 1, \dots$; $E(Y) = \text{var}(Y) = \mu$
 - $Y_1, Y_2, \dots, Y_c \stackrel{\text{ind}}{\sim} \text{Poisson}(\mu_i)$, $\sum_{i=1}^c Y_i \sim \text{Poisson}(\sum_{i=1}^c \mu_i)$
- c indep. Poisson r.v. | total \sim Multinomial

$$\begin{aligned}
 & P(Y_1 = n_1, \dots, Y_c = n_c \mid \sum_i Y_i = n) \\
 &= \frac{P(Y_1 = n_1, \dots, Y_c = n_c)}{P(\sum_i Y_i = n)} \\
 &= \frac{\prod_i [\exp(-\mu_i) \mu_i^{n_i} / n_i!]}{\exp(-\sum_i \mu_i) (\sum_i \mu_i)^n / n!} = \frac{n!}{\prod_i n_i!} \prod_i \pi_i^{n_i}, \quad \pi_i = \frac{\mu_i}{\sum_i \mu_i}
 \end{aligned}$$

2-Way Contingency Tables

Contingency Tables

- *Contingency Table = Classification Table:* frequency of outcomes
- *Two-Way Table:* frequency outcomes of two categorical variables
- *$I \times J$ table:* a table with I rows and J columns.
- Contingency tables can arise from several sampling schemes
 - Inference depends on the sampling scheme
- Example:

Smoking	Lung Cancer		Total
	Cases	Controls	
Yes	688	650	1338
No	21	59	80
Total	709	709	1418

Joint Distribution and Independence

- Underlying probability distribution of X (smoking) and Y (cancer)
- Joint distribution:
 - π_{ij} , probability of cell (i, j)
- Marginal distribution:
 - $\pi_{i+} = \sum_{j=1}^J \pi_{ij}$, probability of row i
 - $\pi_{+j} = \sum_{i=1}^I \pi_{ij}$, probability of column j
- Conditional distribution:
 - $\pi_{j|i} = \pi_{ij}/\pi_{i+}$, distribution of j given i
- Independence:
 - $\pi_{ij} = \pi_{i+}\pi_{+j}$ for all i and j

Multinomial Sampling

- The total sample size n is fixed, but the row and column totals are not
- X and Y are treated equally
 - $P(X = i, Y = j) = \pi_{ij}$, $i = 1, \dots, I$; $j = 1, \dots, J$
 - describe associations with joint distributions.
 - back to the case of the Multinomial distribution
- Likelihood and log-likelihood:

$$\text{Likelihood} = \frac{n!}{n_{11}! \dots n_{IJ}!} \prod_i^I \prod_{j=1}^J \pi_{ij}^{n_{ij}}$$

$$L = \sum_{i=1}^I \sum_{j=1}^J n_{ij} \log(\pi_{ij}) + \text{constant}$$

Multinomial Sampling: Testing for Independence

- Hypotheses:

- H_0 : reduced model $\pi_{ij} = \pi_{i+}\pi_{+j}$, for all i and j
- H_a : full model $\pi_{ij} \neq \pi_{i+}\pi_{+j}$, for some i and j

- Pearson χ^2 test:

- $X^2 = \sum_{i=1}^I \sum_{j=1}^J \frac{(O_{ij} - E_{ij})^2}{E_{ij}} \stackrel{H_0, \text{ approx.}}{\sim} \chi_{(I-1)(J-1)}^2$

- $O_{ij} = n_{ij}$, $E_{ij} = n\hat{\pi}_{i+}\hat{\pi}_{+j} = n_{i+}n_{+j}/n$

- $Df = (I - 1)(J - 1) = (IJ - 1) - (I - 1) - (J - 1)$

- Likelihood Ratio test:

- Full model: $\hat{\pi}_{ij} = n_{ij}/n_{++}$

- Reduced model: $\hat{\pi}_{i+} = n_{i+}/n_{++}$; $\hat{\pi}_{+j} = n_{+j}/n_{++}$.

$$\begin{aligned} G^2 &= 2(L_1 - L_0) \\ &= 2 \sum_{i=1}^I \sum_{j=1}^J n_{ij} \log \frac{n_{ij}n_{++}}{n_{i+}n_{+j}} \stackrel{H_0, \text{ approx.}}{\sim} \chi_{(I-1)(J-1)}^2 \end{aligned}$$

Independent (or Product) Multinomial Sampling

- The row totals n_{i+} , $i = 1, \dots, I$, are fixed
 - E.g., X is an explanatory variable, and response Y occurs separately at each setting of X .
 - View categorical response as function of categorical predictor
 - Describe associations in terms of conditional distributions

$$P(Y = j|X = i) = \pi_{j|i}, \quad i = 1, \dots, I; \quad j = 1, \dots, J$$

- For a fixed i , $\{n_{ij}, j = 1, \dots, J\}$ follow a multinomial distribution

$$f(n_{i1} \dots, n_{iJ}) = \frac{n_{i+}!}{n_{i1}! \dots n_{iJ}!} \prod_{j=1}^J \pi_{j|i}^{n_{ij}}$$

Compare Proportions

- Independent Multinomial Sampling
- $H_0 : \pi_1 = \pi_2$ vs $H_a : \pi_1 \neq \pi_2$
- ML estimate of the difference:
 - $\hat{\pi}_1 - \hat{\pi}_2 = \frac{y_1}{n_1} - \frac{y_2}{n_2}$
 - $SE(\hat{\pi}_1 - \hat{\pi}_2) = \left[\frac{\pi_1(1-\pi_1)}{n_1} + \frac{\pi_2(1-\pi_2)}{n_2} \right]^{1/2}$
- Wald Confidence Interval:
 - $\hat{\pi}_1 - \hat{\pi}_2 \pm z_{\alpha/2} \widehat{SE}(\hat{\pi}_1 - \hat{\pi}_2)$
 - Replace π with $\hat{\pi}$ to estimate SE
- Usually too narrow
- Better methods (e.g. delta method) exist

Testing for Independence of Rows and Columns

- Independent Multinomial Sampling
- Independence in this context is often called *homogeneity* of the conditional distributions
- X and Y are independent
 $\iff \pi_{j|1} = \dots = \pi_{j|I}$, for all j
- Can interpret the independence in terms of product of marginal probabilities
- $\pi_{ij} = \pi_{i+}\pi_{+j}$ for all i and j
 $\iff \pi_{j|1} = \dots = \pi_{j|I}$ for all j

“ \Rightarrow ” $\pi_{j|i} = \pi_{ij}/\pi_{i+} = (\pi_{i+}\pi_{+j})/\pi_{i+} = \pi_{+j}$

“ \Leftarrow ” Let $\pi_{j|i} = a_j$, then $\pi_{+j} = \sum_{i=1}^I \pi_{ij} = \sum_{i=1}^I \pi_{i+}a_j = a_j$
 $\implies \pi_{ij} = \pi_{i+}\pi_{+j}$

Testing for Independence of Rows and Columns

- Test the homogeneity of conditional distributions

Row	Column			Total
	1	...	J	
1	π_{11} $(\pi_{1 1})$...	π_{1J} $(\pi_{J 1})$	π_{1+}
\vdots	\vdots	\vdots	\vdots	\vdots
I	π_{I1} $(\pi_{1 I})$...	π_{IJ} $(\pi_{J I})$	π_{I+}
Total	π_{+1}	...	π_{+J}	π_{++}

- Consider the new notation:

$$\pi_j(x) = P(Y = j|X = x)$$
- Although the interpretation is different, use the same Pearson X^2 test and the LR test

Test for Independence: Odds Ratio

- Odds Ratio:

$$\begin{aligned}\theta &= \frac{\pi_{11}/\pi_{12}}{\pi_{21}/\pi_{22}} = \frac{\pi_{11}\pi_{22}}{\pi_{12}\pi_{21}} \\ &= \frac{P(Y=1|X=1)/P(Y=2|X=1)}{P(Y=1|X=2)/P(Y=2|X=2)} \\ &= \frac{P(X=1|Y=1)/P(X=2|Y=1)}{P(X=1|Y=2)/P(X=2|Y=2)}\end{aligned}$$

- Equally valid for prospective (conditional on X), retrospective (conditional on Y) and cross-sectional (multinomial) sampling designs

- MLE: $\hat{\theta} = \frac{n_{11}/n_{12}}{n_{21}/n_{22}} = \frac{n_{11}n_{22}}{n_{12}n_{21}}$

- When some $n_{ij} = 0$, $\hat{\theta}$ is not a good estimator. Is improved by adding 0.5 to each cell count:

$$\tilde{\theta} = \frac{(n_{11} + 0.5)(n_{22} + 0.5)}{(n_{12} + 0.5)(n_{21} + 0.5)}$$

Test for Independence: Odds Ratio

- X and Y are independent

$$\iff \theta = \frac{\pi_{11}/\pi_{12}}{\pi_{21}/\pi_{22}} = \frac{\pi_{11}\pi_{22}}{\pi_{12}\pi_{21}} = 1$$

- to check, substitute $\pi_{ij} = \pi_{i+}\pi_{+j}$ in the formula above

- Asymptotically, $\log \hat{\theta} \sim N(\log(\theta), \hat{\sigma}^2)$, where

$$\hat{\sigma}^2 = \frac{1}{n_{11}} + \frac{1}{n_{12}} + \frac{1}{n_{21}} + \frac{1}{n_{22}}$$

- Large-sample CI for $\log \theta$:

$$\log \hat{\theta} \pm z_{\alpha/2} \widehat{SE}(\log \hat{\theta}) = [L, U]$$

- Large-sample CI for $\theta : [e^L, e^U]$.

- Usually too wide

Poisson Sampling

- Observe a process over a period of time, and observe the number of occurrences
 - No fixed quantities
 - Poisson sampling assumes each $Y_{ij} \stackrel{ind}{\sim} \text{Poisson}(\pi_{ij})$

- Denote Y_{ij} the count of cell (i, j)

$$\sum_{i=1}^I \sum_{j=1}^J Y_{ij} \sim \text{Poisson} \left(\sum_{i=1}^I \sum_{j=1}^J \pi_{ij} \right)$$

- Hypothesis of independence of X and Y has the form $\log(\pi_{ij}) = \lambda + \alpha_i + \beta_j$
 - This is the log-linear model of independence for two-way contingency tables
 - Under independence, $\log(\mu_{ij})$ is an additive function of a row effect α_i and a column effect β_j .
 - Since we don't have a replicate table, the model with the interaction is saturated

Poisson Sampling

- An additive model

$$\log \pi_{ij} = \mu + \alpha_i + \beta_j$$

implies the independence of the margins

$$\begin{aligned}\pi_{ij} &= \frac{E(count)}{\text{sum of all } E(count)} \\ &= \frac{e^{\mu + \alpha_i + \beta_j}}{e^{\mu}(\sum_i e^{\alpha_i})(\sum_j e^{\beta_j})} = \pi_{i+}\pi_{+j},\end{aligned}$$

where

$$\begin{aligned}\pi_{i+} &= e^{\alpha_i} / \sum_i e^{\alpha_i} = \sum_j \pi_{ij}, \\ \pi_{+j} &= e^{\beta_j} / \sum_j e^{\beta_j} = \sum_i \pi_{ij}.\end{aligned}$$

- Test for independence: Pearson X^2 or LR test as before (more on this later)

Hypergeometric Sampling

- Both row and column margins are fixed.
- When X and Y are independent, given the row and column margins, follows hypergeometric distribution

$$\frac{\left(\prod_{i=1}^I n_{i+}!\right) \left(\prod_{j=1}^J n_{+j}!\right)}{n_{++}! \prod_{i=1}^I \prod_{j=1}^J n_{ij}!}$$

— the distribution is parameter free

- For a 2×2 table

$$P(n_{11} = k) = \frac{\binom{n_{1+}}{k} \binom{n_{2+}}{n_{+1} - k}}{\binom{n_{++}}{n_{+1}}},$$

$$\max(0, n_{1+} + n_{+1} - n) \leq k \leq \min(n_{1+}, n_{+1})$$

— *Fisher's exact test*: p -value = total probability of all outcomes more extreme than the one observed.

— Takes discrete values for small samples

Case study: Agresti p.80

```
#-----read the data-----
X <- data.frame(y=c(178, 138, 108, 570, 648,
  442, 138, 252, 252),
  belief=rep(c("1-Fundam", "2-Moder", "3-Liber"), 3),
  degree=rep(c("1-<HS", "2-HS", "3-BS/grad"), 1, each=3)
)
```

```
#----- a table of observed values (ov)-----
ov <- xtabs(y ~ degree+belief, data=X)
> ov
```

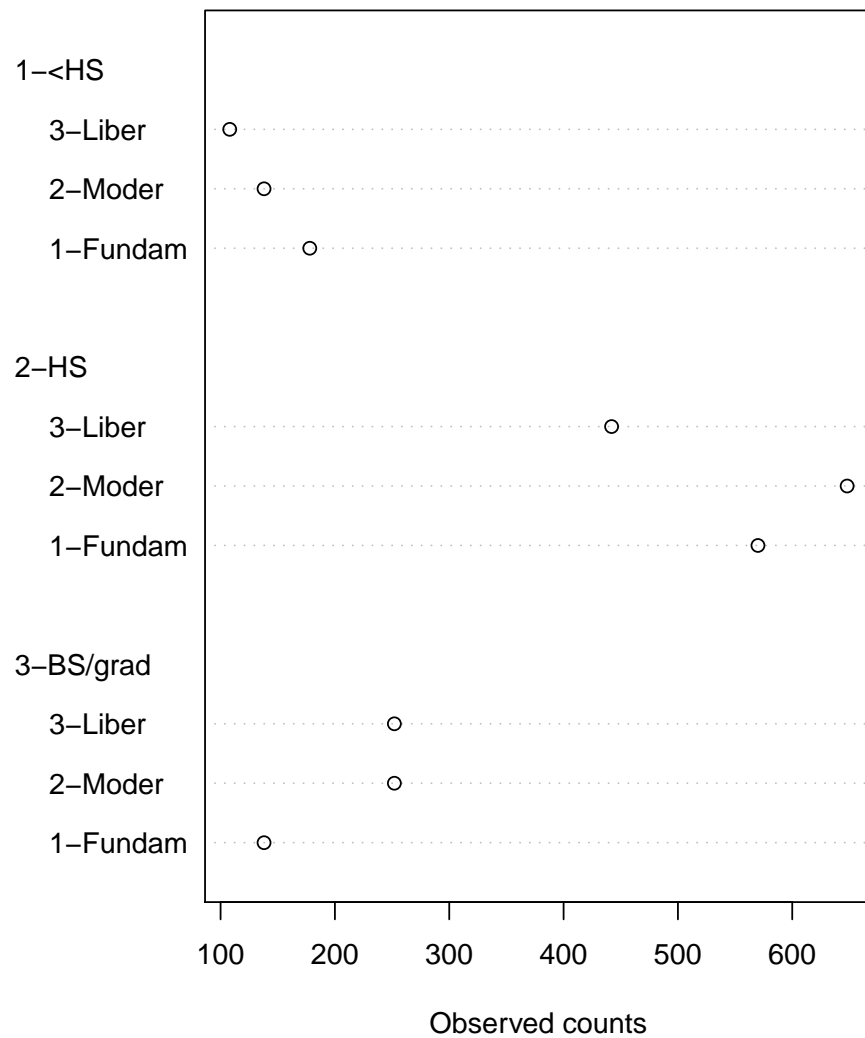
	belief		
degree	1-Fundam	2-Moder	3-Liber
1-<HS	178	138	108
2-HS	570	648	442
3-BS/grad	138	252	252

```
#-----export the table into latex-----
# export the table into latex
library(xtable)
xtable(ov)
```

```
\begin{table}[ht]
\begin{center}
\begin{tabular}{rrrr}
\hline
& 1-Fundam & 2-Moder & 3-Liber \\
\hline
1-<$HS & 178.00 & 138.00 & 108.00 \\
2-HS & 570.00 & 648.00 & 442.00 \\
....
```

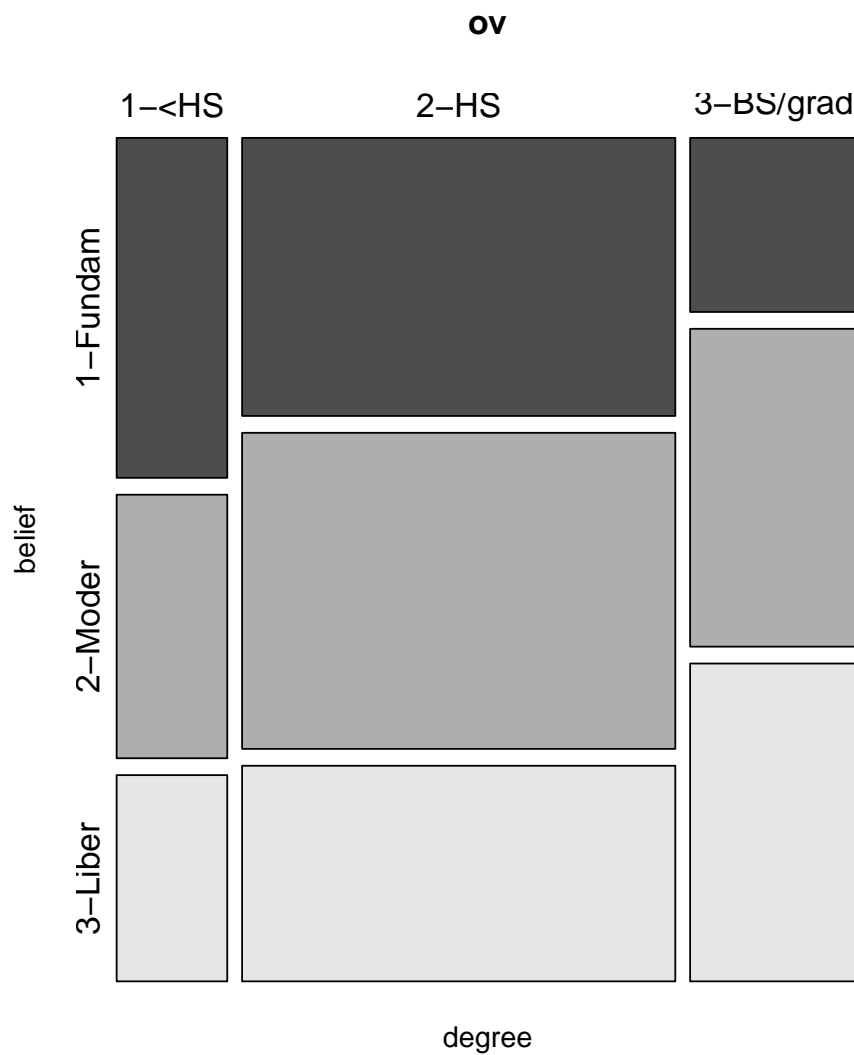
Data visualization

```
#-----dotchart-----  
dotchart(t(ov), xlab="Observed counts")
```



Data visualization

```
#-----mosaic plot-----  
mosaicplot(ov, color=TRUE)
```



2 x 2 table: Compare Proportions

- Independent multinomial sampling:
restrictions on the rows
 - compare proportions of columns, given rows
 - also implements the Pearson X^2 test with Yates correction for small samples (from each O-E, subtract 0.5 if positive, and add 0.5 if negative)

```
> prop.test(ov[1:2,1:2])
2-sample test for equality of proportions
with continuity correction

data:  ov[1:2, 1:2]
X-squared = 8.7451, df = 1, p-value = 0.003104
alternative hypothesis: two.sided
95 percent confidence interval:
 0.03187153 0.15875016
sample estimates:
   prop 1    prop 2 
0.5632911 0.4679803 

#-----double-check the proportions-----
> 178/(178+138)
[1] 0.5632911
> 570/(570+648)
[1] 0.4679803
```

2 x 2 table: Hypergeometric

- Sampling conditional on both margins
 - Hypergeometric test
 - compare distributions of counts within the 4 cells
 - H_0 is specified in terms of $OR=1$
 - produces CI for the OR

```
> fisher.test(ov[1:2,1:2])
```

Fisher's Exact Test for Count Data

```
data:  ov[1:2, 1:2]
p-value = 0.002961
alternative hypothesis: true odds ratio is not equal to 1
95 percent confidence interval:
 1.134415 1.897137
sample estimates:
odds ratio
 1.465974
```

I x J table: Pearson χ^2

- (Independent) multinomial sampling restrictions on a margin, or on the total
 - H_0 in terms of independence of rows and columns

```
> summary(ov)
```

```
Call: xtabs(formula = y ~ degree + belief, data = X)
Number of cases in table: 2726
Number of factors: 2
Test for independence of all factors:
Chisq = 69.16, df = 4, p-value = 3.42e-14
```

- Pearson residuals
 - $e_{ij} = \frac{n_{ij} - \hat{\mu}_{ij}}{\hat{\mu}_{ij}^{1/2}}$
 - divide residual by $\widehat{SE}(n_{ij})$ in Poisson sampling

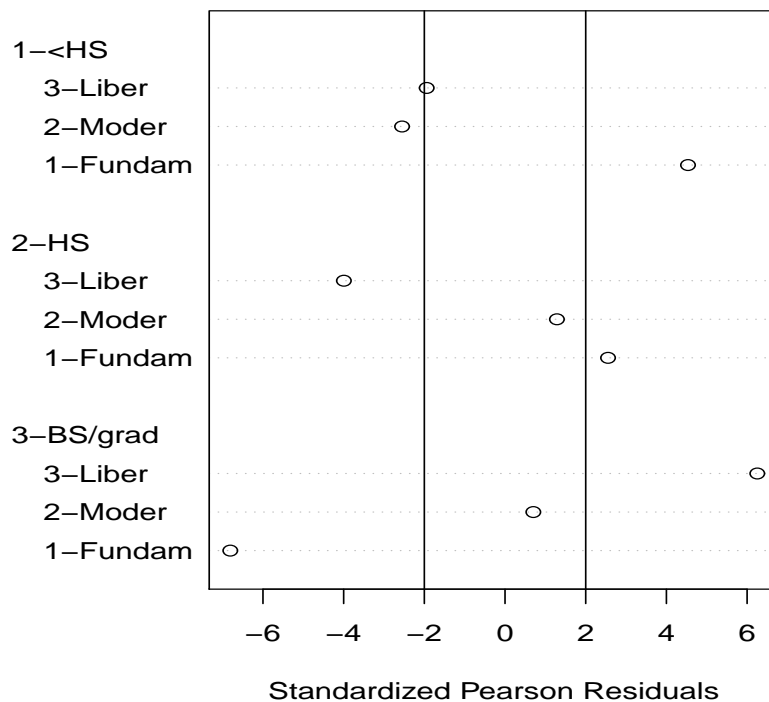
- Standardized Pearson residuals
 - $e_{ij} = \frac{n_{ij} - \hat{\mu}_{ij}}{\sqrt{\hat{\mu}_{ij} \cdot (1 - p_{i+}) (1 - p_{+j})}}$
 - divide residual by $\widehat{SE}(residual)$ in Poisson sampling

Visualizing the association

```
# --Compute Pearson and standardized Pearson residuals ---
e <- apply(ov, 1, sum) %*% t(apply(ov, 2, sum)) / sum(ov)
pearsonResid <- (ov - e)/sqrt(e)

pRow <- 1-apply(ov, 1, sum) / sum(ov)
pCol <- 1-apply(ov, 2, sum) / sum(ov)
standPearsonResid <- pearsonResid/ sqrt(pRow %*% t(pCol) )

dotchart( t(standPearsonResid) )
abline(v=c(-2,2))
```



Ordered Categories

- Ordered categories have more info
- Assign scores to categories
 - Rows: $(u_1 \leq, \dots \leq u_I)$, e.g. $(1, \dots, I)$
 - Cols: $(v_1 \leq, \dots \leq v_J)$, e.g. $(1, \dots, J)$
 - $H_0 : \text{cor}(u, v) = 0$ vs $H_a : \text{cor}(u, v) \neq 0$
 - produces CI for the OR
- Study the linear trend

$$r = \frac{\sum_{i=1}^I \sum_{j=1}^J (u_i - \bar{u})(v_j - \bar{v})n_{ij}}{\sqrt{\left[\sum_{i=1}^I \sum_{j=1}^J (u_i - \bar{u})^2 n_{ij} \right] \cdot \left[\sum_{i=1}^I \sum_{j=1}^J (v_i - \bar{v})^2 n_{ij} \right]}}$$

$$- \bar{u} = \sum_{i=1}^I \sum_{j=1}^J u_i n_{ij} / n; \quad \bar{v} = \sum_{i=1}^I \sum_{j=1}^J v_i n_{ij} / n;$$

$$- M^2 = (n - 1)r^2 \stackrel{H_0}{\sim} \chi_1^2$$

Case Study: Ordered Categories

```
#-----existing implementation-----  
> library(coin)  
> lbl_test(as.table(ov))
```

Asymptotic Linear-by-Linear Association Test

```
data:  belief (ordered) by degree (1-<HS < 2-HS < 3-BS/grad)  
chi-squared = 56.0849, df = 1, p-value = 6.939e-14
```

```
#-----manually-----  
u <- as.vector(scale(1:3, center=sum(c(1:3)*ov)/sum(ov),  
  scale=FALSE))  
v <- as.vector(scale(1:3, center=sum(t(ov)*c(1:3))/sum(ov),  
  scale=FALSE))  
  
r <- sum(u%*%t(v)*ov) / sqrt(sum(u^2*ov) * sum(t(ov) * v^2))  
  
M2 <- (sum(ov) - 1) * r^2  
> 1-pchisq(M2, 1, lower=TRUE)  
[1] 6.938894e-14
```

2x2 pairs: Matched Pairs

- Repeated measurements on same subjects
 - ask the same people the same question twice
 - goal: compare proportions
 - absence of association cannot be interpreted as independence
- Example (Agresti Ch. 10.1)
 - Approval of the President's performance, one month apart, for a same sample of Americans.

	Approve	Disapprove
Approve	794.00	150.00
Disapprove	86.00	570.00

- H_0 : Marginal homogeneity. $\pi_{1+} = \pi_{+1}$
 - $\delta = \pi_{1+} - \pi_{+1} = (\pi_{11} + \pi_{12}) - (\pi_{11} + \pi_{21}) = \pi_{12} - \pi_{21}$
 - Equivalent to testing table symmetry

Large-sample test and CI

- CI

- $\hat{\delta} = p_{+1} - p_{1+} = p_{2+} - p_{+2}$
- $var(\hat{\delta}) = [\pi_{1+}(1 - \pi_{+1}) + \pi_{+1}(1 - \pi_{+1}) - 2(\pi_{11}\pi_{22} - \pi_{12}\pi_{21})] / n$
- smaller variance than in independent samples, therefore a more efficient design
- $\widehat{var}(\hat{\delta}) = [(p_{12} + p_{21}) - (p_{12} - p_{21})^2] / n$
- CI: $\hat{\delta} \pm z_{\alpha/2} \widehat{SE}(\hat{\delta})$

- Wald Test

- $z = \frac{\hat{\delta}}{\widehat{SE}(\delta)} = \frac{n_{21} - n_{12}}{(n_{21} + n_{12})^{1/2}}$
- $z^2 \stackrel{H_0}{\sim} \chi_1^2$ (called McNemar test)
- Only depends on counts outside of the diagonal

President Approval Example

```
#-----Read the data-----  
Performance <-  
matrix(c(794, 86, 150, 570),  
       nrow = 2, dimnames =  
       list("1st Survey" = c("Approve", "Disapprove"),  
            "2nd Survey" = c("Approve", "Disapprove"))  
       )
```

```
> Performance  
      2nd Survey  
1st Survey Approve Disapprove  
Approve      794      150  
Disapprove    86      570
```

```
#-----Test-----  
> mcnemar.test(Performance)
```

McNemar's Chi-squared test with continuity correction

data: Performance

McNemar's chi-squared = 16.8178, df = 1, p-value = 4.115e-05

- significant change (in fact, drop) in approval ratings