$ORIGIN \equiv 1$

2 X 2 Contingency Test of Association

Contingency tests consider data from categorical (also called nominal) variables - variables in which observations may be placed in classes, but the classes themselves need not have numerical or ordinal significance. When comparing two categorical variables it is customary to construct a **contingency table** showing which observations may be simultaneously classified according to the classes. From the contingency table, tests of association (or alternatively tests of independence) may be performed. Here we look at the 2X2 case in which there are only 2 classes for each of two variables.

Data Structure:

Zar 2010 pp. 497-499 describes three different interpretations of a 2X2 contingency table depending on what's determined in advance by the researcher:

1) *No fixed margins* - row and column totals are not determined in advance.

2) One fixed margin - row or column totals determined in advance but not both.

3) *Both margins fixed* - both row and column totals are determined in advance

Disagreements exist over whether or not to correct the test statistic accordingly. Zar suggests use of Yate's correction shown below in the case of 3) only. Other correction methods are also available.

Assumptions:

Observed values X₁, X₂, X₃, ... X_{n1} are a random sample
 Observed values Y₁, Y₂, Y₃, ... Y_{n2} are a random sample.

Model:

Let Probabilities: $-P_i = P(X=i)$ $-P_j = P(Y=j)$ $-P_{ii} = (X=i, Y=j)$

Continger		
1,1	1,2	X=1
2,1	2,2	X=2
Y=1	Y=2	Grand Total

Zar Example 23.2

$\mathbf{O} \coloneqq \begin{pmatrix} 6 & 12 \\ 28 & 24 \end{pmatrix}$	< observed		boys	girls	total
		left-handed	6	12	18
GM := 70 < grand total	< grand total	right-handed	28	24	52
		total	34	36	70

Hypotheses:

$H_0: P_{ij} = (P_i)(P_j)$	< That is, variables X & Y are independent!
$\mathbf{H}_{1}:\mathbf{P}_{ij} \diamondsuit (\mathbf{P}_{i})(\mathbf{P}_{j})$	< Two sided test

Criterion for Normal Approximation:

- IF expected values in each cell $E_{ij} \ge 5$ THEN Normal Approximation may be used OTHERWISE use an Exact Test such as Fisher's Exact Test

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Normal Approximation:

Construct Expected for each cell:

- Tabulate O_{ii} for each cell
- Calculate Observed Row and Column Totals
- Calculate Expected for each cell

$$E := \begin{pmatrix} \frac{18 \cdot 34}{70} & \frac{18 \cdot 36}{70} \\ \frac{52 \cdot 34}{70} & \frac{52 \cdot 36}{70} \end{pmatrix} \qquad E = \begin{pmatrix} 8.7429 & 9.2571 \\ 25.2571 & 26.7429 \end{pmatrix}$$

^ multiply row total by column total and divide by total

Test Statistic:

$$i := 1..2 \qquad j := 1..2$$

$$\chi := \sum_{i} \sum_{j} \frac{\left(O_{i,j} - E_{i,j}\right)^{2}}{E_{i,j}} \qquad \chi = 2.2524 \qquad < \text{Standard statistic}$$

$$\chi_{Y} := \sum_{i} \sum_{j} \frac{\left(\left|O_{i,j} - E_{i,j}\right| - \frac{1}{2}\right)^{2}}{E_{i,j}} \qquad \chi_{Y} = 1.5061 \qquad < \text{Yate's corrected statistic}$$

Sampling Distribution:

If Assumptions hold, H_0 is true and assuming Normal Approximation, then $\chi_Y \sim \chi^2_{(1)}$

Critical Value of the Test:

 $\alpha := 0.05$ < **Probability of Type I error must be explicitly set**

 $CV := qchisq(1 - \alpha, 1)$ CV = 3.8415

Decision Rule:

IF $\chi > CV$ THEN REJECT H₀, OTHERWISE ACCEPT H₀ IF $\chi_Y > CV$ THEN REJECT H₀, OTHERWISE ACCEPT H₀

Probability Value:

$$P := (1 - pchisq(\chi, 1)) \qquad P = 0.1334 \qquad < Standard probability$$
$$P_{Y} := (1 - pchisq(\chi_{Y}, 1)) \qquad P_{Y} = 0.2197 \qquad < with Yate's correction$$

Prototype in R:

#2X2 CONTINGENCY TABLES ZAR=read.table("c:/DATA/Biostatistics/ZarEX23.2R.txt") ZAR attach(ZAR) Y=count A=handedness B=sex

#CROSS-TABULATION IN R: X=xtabs(Y~A+B) X

#USING CHI-SQUARE STATISTIC: chisq.test(X,correct=F) #USING YATE'S CORRECTED STATISTIC: chisq.test(X,correct=T)

#DATA KEYED INTO MATRIX BY ROW #USING CONCATENATION FUNCTION c() AND MATRIX FUNCTION matrix():

X=matrix(c(6,12,28,24),nrow=2,byrow=T)

Χ

Pearson's Chi-squared test

data: X X-squared = 2.2524, df = 1, p-value = 0.1334

Pearson's Chi-squared test with Yates' continuity correction

> X

B A I

boys girls

left 6 12

right 28 24

data: X X-squared = 1.5061, df = 1, p-value = 0.2197