



STAT 426

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All models are wrong, but some are useful.

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Chapter 1 Basic of Categorical Data

1.1 Variable Measurement

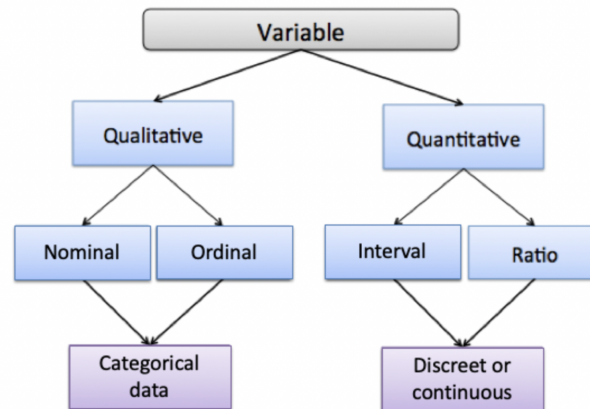


Figure 1.1: Variable Type

- a) Nominal: Categories do not have a natural order. Ex. blood type, gender.
- b) Ordinal: Categories have a natural order. Ex. low/middle/high education level
- c) Interval: There is a numerical distance (difference between two different values is meaningful) between any two values. Ex. blood pressure level, 100 blood pressure doesn't mean the double degree of 50 pressure.
- d) Ratio: An interval variable where ratios are valid (presence of absolute zero, i.e. zero is meaningful). Ex. weight, 4g is double degree of 2g, distance run by an athlete.

Levels of measurements

A variable's level of measurement determines the statistical methods to be used for its analysis.

Variables hierarchy: Ratio > Interval > Ordinal > Nominal

Statistical methods applied to variables at a lower level can be used with variables at a higher level, but the contrary is not true.

1.2 Statistical Inference for Categorical Data

There is a distribution $F(\beta)$ with p.d.f. (p.m.f.) $f(x | \beta)$, where β a generic unknown parameter and $\hat{\beta}$ the parameter estimate.

1.2.1 Maximum likelihood Estimation (MLE)

Given a set of observations $\vec{x} = (x_1, \dots, x_n)$, the likelihood function of these observations with parameter β is $l(\vec{x} | \beta)$. We want to find parameter $\hat{\beta}$ that maximizes the likelihood function,

$$\hat{\beta} = \arg \max_{\beta} l(\vec{x} | \beta)$$

which is also equivalent to maximizing the logarithm of the likelihood function $L(\vec{x} | \beta) = \log(l(\vec{x} | \beta))$,

$$\hat{\beta} = \arg \max_{\beta} L(\vec{x} | \beta)$$

Definition 1.1 (score function)

The score function is

$$u(\beta, \vec{x}) = \nabla_{\beta} L(\vec{x} | \beta) = \frac{\nabla_{\beta} l(\vec{x} | \beta)}{l(\vec{x} | \beta)}$$



Lemma 1.1 (mean of score function)

The mean of score function is 0,

$$\mathbb{E}_{\vec{x}} u(\beta, \vec{x}) = 0$$



Proof 1.1

$$\begin{aligned} \mathbb{E}_{\vec{x}} u(\beta, \vec{x}) &= \int_{\vec{x}} l(\vec{x} | \beta) \frac{\nabla_{\beta} l(\vec{x} | \beta)}{l(\vec{x} | \beta)} d\vec{x} \\ &= \int_{\vec{x}} \nabla_{\beta} l(\vec{x} | \beta) d\vec{x} \\ &= \nabla_{\beta} \left(\int_{\vec{x}} l(\vec{x} | \beta) d\vec{x} \right) \\ &= \nabla_{\beta} 1 = 0 \end{aligned}$$

Lemma 1.2 (variance of score function)

The variance of the score function is

$$\text{Var}_{\vec{x}}(u(\beta, \vec{x})) = \mathbb{E}_{\vec{x}} (u(\beta, \vec{x}) u(\beta, \vec{x})^T)$$



Proof 1.2

Prove by the zero mean.

Definition 1.2 (Fisher information)

The (Fisher) information is

$$\iota(\beta) = -\mathbb{E}_{\vec{x}} [\nabla_{\beta}^2 L(\vec{x} | \beta)]$$

**Lemma 1.3**

The Fisher information is equal to the variance of score function.

$$\text{Var}_{\vec{x}}(u(\beta, \vec{x})) = \mathbb{E}_{\vec{x}}(u(\beta, \vec{x})u(\beta, \vec{x})^T) = -\mathbb{E}_{\vec{x}}[\nabla_{\beta}^2 L(\vec{x} | \beta)] = \iota(\beta)$$

**Proof 1.3**

$$\mathbb{E}_{\vec{x}}[\nabla_{\beta}^2 L(\vec{x} | \beta)] = \mathbb{E}_{\vec{x}}\left(\frac{\partial \frac{\nabla_{\beta} l(\vec{x} | \beta)}{l(\vec{x} | \beta)}}{\partial \beta}\right) = \mathbb{E}_{\vec{x}}\left(\frac{\nabla_{\beta}^2 l(\vec{x} | \beta)}{l(\vec{x} | \beta)} - \frac{\nabla_{\beta} l(\vec{x} | \beta) \nabla_{\beta} l(\vec{x} | \beta)^T}{(l(\vec{x} | \beta))^2}\right)$$

where $\mathbb{E}_{\vec{x}}\left(\frac{\nabla_{\beta}^2 l(\vec{x} | \beta)}{l(\vec{x} | \beta)}\right) = \int_{\vec{x}} l(\vec{x} | \beta) \frac{\nabla_{\beta}^2 l(\vec{x} | \beta)}{l(\vec{x} | \beta)} d\vec{x} = \int_{\vec{x}} \nabla_{\beta}^2 l(\vec{x} | \beta) d\vec{x} = \nabla_{\beta}^2 \int_{\vec{x}} l(\vec{x} | \beta) d\vec{x} = \nabla_{\beta}^2 1 = 0$

Hence,

$$\mathbb{E}_{\vec{x}}[\nabla_{\beta}^2 L(\vec{x} | \beta)] = -\mathbb{E}_{\vec{x}}\left(\frac{\nabla_{\beta} l(\vec{x} | \beta) \nabla_{\beta} l(\vec{x} | \beta)^T}{(l(\vec{x} | \beta))^2}\right) = -\mathbb{E}_{\vec{x}}(u(\beta, \vec{x})u(\beta, \vec{x})^T)$$

Proposition 1.1

When the sample x is made up of i.i.d. observations, the covariance matrix of the maximum likelihood estimator $\hat{\beta}$ is approximately equal to the inverse of the information matrix.

$$\text{Cov}(\hat{\beta}) \approx (\iota(\beta))^{-1}$$



Hence, the covariance matrix can be estimated as $(\iota(\hat{\beta}))^{-1}$. Similarly, SE is estimated by $\sqrt{(\iota(\hat{\beta}))^{-1}}$.

1.2.2 Likelihood Inference (Wald, Likelihood-Ratio, Score)

We want to test

$$H_0 : \beta = \beta_0 \quad H_{\alpha} : \beta \neq \beta_0$$

or form a confidence interval (CI) for β .

Definition 1.3 (Wald Test)

The Wald statistic:

$$z_W = \frac{\hat{\beta} - \beta_0}{SE} = \frac{\hat{\beta} - \beta_0}{\sqrt{(\iota(\hat{\beta}))^{-1}}}$$

where $SE = \sqrt{(\iota(\hat{\beta}))^{-1}}$.

Usually, as $n \rightarrow \infty$, $z_W \xrightarrow{d} N(0, 1)$ under $H_0 : \beta = \beta_0$.

(1) We reject the H_0 if $|z_W| \geq z_{\frac{\alpha}{2}}$ for a two-sided level α test.

(2) The $(1 - \alpha)100\%$ Wald (confidence) interval is

$$\{\beta_0 : |z_W| = \frac{|\hat{\beta} - \beta_0|}{SE} < z_{\frac{\alpha}{2}}\} = (\hat{\beta} - z_{\frac{\alpha}{2}} SE, \hat{\beta} + z_{\frac{\alpha}{2}} SE)$$

(3) The Wald test also has a chi-squared form, using

$$z_W^2 = \frac{(\hat{\beta} - \beta_0)^2}{(\iota(\hat{\beta}))^{-1}} \sim \chi_1^2 \quad (\text{under } H_0)$$



Definition 1.4 (Likelihood Ratio Test)

Let

$$\Lambda = \frac{l(\vec{x} | \beta_0)}{l(\vec{x} | \hat{\beta})}$$

where $l(\vec{x} | \hat{\beta}) = \max_{\beta} l(\vec{x} | \beta)$, so the ratio $\Lambda \in [0, 1]$.

The **likelihood-ratio test (LRT) chi-squared statistic**:

$$-2 \ln \Lambda = -2 \left(L(\beta_0) - L(\hat{\beta}) \right)$$

It has an approximate χ_1^2 distribution under $H_0 : \beta = \beta_0$, and otherwise tends to be larger.

(1) Thus, reject H_0 if

$$-2 \ln \Lambda \geq \chi_1^2(\alpha)$$

(2) The $(1 - \alpha)100\%$ likelihood-ratio (confidence) interval is

$$\{\beta_0 : -2 \ln \Lambda = -2 \left(L(\beta_0) - L(\hat{\beta}) \right) < \chi_1^2(\alpha)\}$$

Unlike Wald, this interval is not degenerate. (i.e., For general case, the interval does not have an explicit form.)



Definition 1.5 (Score Test)

The **score statistic**:

$$z_S = \frac{u(\beta_0)}{\sqrt{\iota(\beta_0)}}$$

As $n \rightarrow \infty$, $z_S \xrightarrow{d} N(0, 1)$ under $H_0 : \beta = \beta_0$. Otherwise, it tends to be further from zero.

(1) Thus, reject H_0 if $|z_S| \geq z_{\frac{\alpha}{2}}$ for a two-sided level α test.

(2) The $(1 - \alpha)100\%$ score (confidence) interval is

$$\{\beta_0 : |z_S| = \frac{|u(\beta_0)|}{\sqrt{\iota(\beta_0)}} < z_{\frac{\alpha}{2}}\}$$

Unlike Wald, it is not degenerate for some distributions.

(3) *There is also a chi-squared form:*

$$z_S^2 = \frac{u(\beta_0)^2}{\iota(\beta_0)} \sim \chi_1^2 \quad (\text{under } H_0)$$



We can also use P-value to measure the probability of the statistic is more extreme under the H_0 . We can reject H_0 if the P-value is $\leq \alpha$.

All three kinds tend to be “asymptotically equivalent” as $n \rightarrow \infty$. For smaller n , the likelihood-ratio and score methods are preferred.

Chapter 2 Association in Contingency Tables

2.1 Association in Two-Way Contingency Tables

Consider joint observations of two categorical variables: X with I categories, Y with J categories.

We can summarize data in an $I \times J$ **contingency table**:

		Y		
		1	...	J
X	1			
	\vdots			
	I			

Each **cell** contains a count n_{ij} .

2.1.1 Distribution

If both X and Y are random, let

$$\pi_{ij} = P(X \text{ in row } i, Y \text{ in col } j)$$

be the **joint** distribution of X and Y .

The **marginal** distribution of X is defined by

$$\pi_{i+} = P(X \text{ in row } i)$$

and similarly for Y :

$$\pi_{+j} = P(Y \text{ in col } j)$$

The **conditional** distribution of Y given that X is in row i is defined by

$$\pi_{j|i} = P(Y \text{ in col } j \mid X \text{ in row } i) = \frac{\pi_{ij}}{\pi_{i+}}$$

2.1.2 Descriptive Statistics

Let n_{ij} = count in row i and col j and $n = \sum_i \sum_j n_{ij}$.

The **margins** of the table:

$$n_{i+} = \sum_j n_{ij}, \quad n_{+j} = \sum_i n_{ij}$$

Natural Estimation

1. Natural estimate of π_{ij} : $\hat{\pi}_{ij} = \frac{n_{ij}}{n}$
2. Similarly marginals: $\hat{\pi}_{i+} = \sum_j \hat{\pi}_{ij} = \frac{n_{i+}}{n}$; $\hat{\pi}_{+j} = \sum_i \hat{\pi}_{ij} = \frac{n_{+j}}{n}$
3. And conditionals: $\hat{\pi}_{j|i} = \frac{\hat{\pi}_{ij}}{\hat{\pi}_{i+}} = \frac{n_{ij}}{n_{i+}}$

2.1.3 Sampling Models (Examples)

Possible joint distributions for counts in $I \times J$ table:

1. Poisson (random total): $Y_{ij} = \text{count in cell } (i, j)$,

$$Y_{ij} \sim \text{Poisson}(\mu_{ij})$$

and the Y_{ij} s are independent.

2. Multinomial (fixed total n): $N_{ij} = \text{count in cell } (i, j)$,

$$\{N_{ij}\} \sim \text{multinomial}(n, \{\pi_{ij}\})$$

3. Independent Multinomial: Assume n_{i+} (row totals n_i) are fixed,

$$\left. \begin{aligned} \{N_{1j}\}_{j=1}^J &\sim \text{multinomial}(n_1, \{\pi_{j|1}\}_{j=1}^J) \\ &\vdots \\ \{N_{Ij}\}_{j=1}^J &\sim \text{multinomial}(n_I, \{\pi_{j|I}\}_{j=1}^J) \end{aligned} \right\}$$

(When $J = 2$, this is independent binomial sampling, for which we may just write π_i for $\{\pi_{1|i}, \pi_{2|i}\}$.)

2.1.4 Independent / Homogeneity

Definition 2.1 (independent)

If both X and Y are random, they are **independent** if

$$\pi_{ij} = \pi_{i+}\pi_{+j}, \forall i, j$$

which implies $\pi_{j|i} = \frac{\pi_{i+}\pi_{+j}}{\pi_{i+}} = \pi_{+j}, \forall i, j$. That is, $\pi_{j|i}$ doesn't depend on i and is the same as the marginal distribution of Y . (Intuitively, knowing X tells nothing about Y .)



Definition 2.2 (homogeneity)

Even if X is not really random, the condition that $\pi_{j|i} = \pi_{+j}, \forall i, j$ is called **homogeneity**. This might still be relevant in a situation where X is deliberately chosen and Y is observed as a response.



2.1.5 Measuring Inhomogeneity

Homogeneity is the condition $\pi_1 = \pi_2$. We can measure inhomogeneity by three different measures:

n_{11}	n_{12}
n_{21}	n_{22}

Y_1	$n_1 - Y_1$
Y_2	$n_2 - Y_2$

where $Y_i \sim \text{indep. binomial}(n_i, \pi_i)$. This regards row totals as fixed.

1. difference of proportions:

$$\pi_1 - \pi_2$$

The estimation is

$$\hat{\pi}_1 - \hat{\pi}_2 = \frac{y_1}{n_1} - \frac{y_2}{n_2}$$

The approx $(1 - \alpha)100\%$ confidence interval is:

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

(Problematic if π_1 and π_2 are near 0 or 1.)

2. relative risk:

$$RR = \frac{\pi_1}{\pi_2}$$

The estimation is

$$r = \frac{\hat{\pi}_1}{\hat{\pi}_2} = \frac{y_1/n_1}{y_2/n_2}$$

The approx $(1 - \alpha)100\%$ confidence interval of $\ln RR$ is:

$$\ln r \pm z_{\frac{\alpha}{2}} \sqrt{\frac{1 - \hat{\pi}_1}{y_1} + \frac{1 - \hat{\pi}_2}{y_2}}$$

3. odds ratio:

$$\theta = \frac{\pi_1/(1 - \pi_1)}{\pi_2/(1 - \pi_2)}$$

When $\theta = 1$, we can say there is no association.

The **odds** for a probability π is $\Omega = \frac{\pi}{1 - \pi}$. Note $\pi = \frac{\Omega}{1 + \Omega}$.

(In the multinomial model: $\theta = \frac{\pi_{11}\pi_{22}}{\pi_{12}\pi_{21}}$ ("cross-product ratio"); in Poisson model: $\theta = \frac{\mu_{11}\mu_{22}}{\mu_{12}\mu_{21}}$)

The usual (unrestricted) estimates

$$\hat{\theta} = \frac{n_{11}n_{22}}{n_{12}n_{21}}$$

The approx $(1 - \alpha)100\%$ confidence interval for $\ln \theta$ is

$$\ln \hat{\theta} \pm z_{\frac{\alpha}{2}} \sqrt{\frac{1}{n_{11}} + \frac{1}{n_{12}} + \frac{1}{n_{21}} + \frac{1}{n_{22}}}$$

Useful properties of odds ratio:

- (1) Interchanging rows (or cols) changes θ to $\frac{1}{\theta}$.
- (2) Interchanging X and Y doesn't change θ .
- (3) Multiplying a row (or col) by a factor doesn't change $\hat{\theta}$.
- (4) Relationship to relative risk: $\theta = RR \cdot \frac{1-\pi_2}{1-\pi_1}$. (θ and RR are similar if both π_1 and π_2 are small.)

2.1.6 Delta Method

It is easy to obtain approximate CI for a mean based on a sample mean by using the Central Limit Theorem and a consistent estimate of standard error.

But the log Odds Ratio and log Relative Risk are transformed means. How were their CI's derived? And why take logs?

Suppose a statistic T_n and parameter θ such that

$$\sqrt{n}(T_n - \theta) \xrightarrow[n \rightarrow \infty]{d} N(0, \sigma^2)$$

(e.g. T_n might be a sample mean from a sample of size n with population mean θ and variance σ^2)

We want a CI for $g(\theta)$, for some smooth g .

The Taylor expand at T_n is

$$g(\theta) \approx g(T_n) + g'(T_n)(\theta - T_n)$$

So,

$$\sqrt{n}(g(T_n) - g(\theta)) \approx g'(T_n)\sqrt{n}(T_n - \theta) \xrightarrow[n \rightarrow \infty]{d} N(0, (g'(T_n))^2 \sigma^2)$$

(This is useful only if $g'(T_n) \neq 0$) Hence, when n is large

$$\sqrt{n} \frac{g(T_n) - g(\theta)}{|g'(T_n)|\sigma} \sim N(0, 1)$$

which suggests this approximate CI for $g(\theta)$:

$$g(T_n) \pm z_{\frac{\alpha}{2}} \frac{|g'(T_n)|\sigma}{\sqrt{n}}$$

2.1.7 Testing Independence: X^2 and G^2 Test

Let $\mu_{ij} = \mathbb{E}(N_{ij}) = n\pi_{ij}$. Under $H_0 : \pi_{ij} = \pi_{i+}\pi_{+j}, \forall i, j$

$$\mu_{ij} = n\pi_{ij} = n\pi_{i+}\pi_{+j}$$

Under H_0 , can show the MLEs are

$$\hat{\mu}_{ij} = n\hat{\pi}_{i+}\hat{\pi}_{+j} = n \left(\frac{n_{i+}}{n} \right) \left(\frac{n_{+j}}{n} \right) = \frac{n_{i+}n_{+j}}{n}$$

(assuming no empty rows or cols)

Residuals:

1. Raw: $n_{ij} - \hat{\mu}_{ij}$
2. Pearson: $e_{ij} = \frac{n_{ij} - \hat{\mu}_{ij}}{\sqrt{\hat{\mu}_{ij}}}$. $X^2 = \sum_i \sum_j e_{ij}^2$.
3. Standardized: $r_{ij} = \frac{n_{ij} - \hat{\mu}_{ij}}{\sqrt{\hat{\mu}_{ij}(1 - \hat{\pi}_{i+})(1 - \hat{\pi}_{+j})}}$

Usage: Look for Pearson or standardized residuals with absolute value *exceeding 2 or 3*. These suggest the reason for significant dependence.

Remark: Under independence, both Pearson and standardized residuals are asymp. normal, but only standardized has asymp. variance equal to 1.

Definition 2.3 (X^2 Test: Pearson χ^2 Test (Score Test))

$$X^2 = \sum_{ij} \frac{(n_{ij} - \hat{\mu}_{ij})^2}{\hat{\mu}_{ij}} \underset{H_0}{\sim} \chi^2_{(I-1)(J-1)}$$

Note:

$$\begin{aligned} (I-1)(J-1) &= (IJ-1) - ((I-1) + (J-1)) \\ &= \text{total \# params.} - \# \text{ params. under } H_0 \end{aligned}$$

Reject H_0 if

$$X^2 > \chi^2_{(I-1)(J-1)}(\alpha)$$

(or use P-value)



Definition 2.4 (G^2 Test: Likelihood Ratio χ^2 Test)

$$G^2 = 2 \sum_{ij} n_{ij} \ln \frac{n_{ij}}{\hat{\mu}_{ij}} \underset{H_0}{\sim} \chi^2_{(I-1)(J-1)}$$

Reject H_0 if

$$G^2 > \chi^2_{(I-1)(J-1)}(\alpha)$$

(or use P-value)

(Convention: $0 \ln 0 = 0$)



Comparison:

1. X^2 and G^2 are asymptotically equivalent under H_0
2. The X^2 tends to be better.

Example 2.1 Testing independence is equivalent to testing homogeneity in the indep. binomial model:

$$H_0 : \pi_1 = \pi_2$$

Can show

$$X^2 = z^2$$

where

$$z = \frac{\hat{\pi}_1 - \hat{\pi}_2}{\sqrt{\hat{\pi}(1-\hat{\pi})(1/n_1 + 1/n_2)}} \quad \hat{\pi} = \frac{y_1 + y_2}{n_1 + n_2}$$

Remark: The X^2 and G^2 tests are not necessarily compatible with the Wald CIs. For example,

$$\text{reject } H_0 \nleftrightarrow \text{odds ratio } \theta = \frac{\pi_1/(1-\pi_1)}{\pi_2/(1-\pi_2)} = 1 \text{ not in Wald CI}$$

2.1.8 Testing Independence: Fisher's Exact Test

When cell counts are small, the X^2 and G^2 independence tests are not recommended: The χ^2 approximations are poor. In this section we introduce a *Fisher's Exact Test*.

Consider a 2×2 table with row and col totals fixed:

	Y		
X	N_{11}	N_{12}	n_{1+}
	N_{21}	N_{22}	n_{2+}
	n_{+1}	n_{+2}	n

Note: Any cell count, say N_{11} , determines the whole table.

Can show that, under H_0 : independence, N_{11} is (conditionally) hypergeometric:

$$P_{H_0}(N_{11} = t) = \frac{\binom{n_{1+}}{t} \binom{n_{2+}}{n_{+1} - t}}{\binom{n}{n_{+1}}}$$

In terms of odds ratio $\theta = \frac{\pi_1/(1-\pi_1)}{\pi_2/(1-\pi_2)}$, independence is

$$H_0 : \theta = 1$$

Possible alternatives:

$$H_\alpha : \theta > 1 \quad \Rightarrow \quad N_{11} \text{ tends larger}$$

$$H_\alpha : \theta < 1 \quad \Rightarrow \quad N_{11} \text{ tends smaller}$$

$$H_\alpha : \theta \neq 1 \quad \Rightarrow \quad N_{11} \text{ tends larger or smaller}$$

For $H_\alpha : \theta > 1$, the (one-sided) p -value is $P_{H_0}(N_{11} \geq t_0)$, where $t_0 = n_{11}$ is the observed value of N_{11} .

Remarks: Could use mid p -values instead; Implemented in R function `fisher.test()`; Can be extended to $I \times J$ tables (with some computational difficulty).

2.2 Conditional Association in Three-Way Tables

Add a third categorical variable Z .

Example 2.2 Is a drug more effective at curing a disease among younger patients than among older? X = drug or placebo; Y = disease cured or not; Z = age group (young, old).

2.2.1 Conditional Association

Z may be called a **stratification variable**. We are interested in the distribution of (X, Y) *conditional* on Z .

Definition 2.5 (partial table)

Each Z category defines a **partial table** for X and Y .



Example 2.3 When $Z = 1, 2$ and X, Y are binary ($2 \times 2 \times 2$ table):

$$\begin{array}{c|cc} & \text{Y} & \\ \hline Z = 1 : X & \begin{array}{c|c} n_{111} & n_{121} \\ \hline n_{211} & n_{221} \end{array} & \end{array} \quad \begin{array}{c|cc} & \text{Y} & \\ \hline Z = 2 : X & \begin{array}{c|c} n_{112} & n_{122} \\ \hline n_{212} & n_{222} \end{array} & \end{array}$$

These represent **conditional associations**.

Definition 2.6 (marginal table)

The *marginal table* sums the partial tables:


$$\begin{array}{c|cc} & \multicolumn{2}{c}{Y} \\ X & n_{11+} & n_{12+} \\ \hline & n_{21+} & n_{22+} \end{array}$$

This represents the **marginal association** (ignoring Z).

In general, let $\mu_{ijk} = \text{expected count in row } i, \text{ col } j, \text{ table } k$.

The conditional odds ratios,

$$\theta_{XY(k)} = \frac{\mu_{11k}\mu_{22k}}{\mu_{12k}\mu_{21k}}$$

which are estimated by

$$\hat{\theta}_{XY(k)} = \frac{n_{11k}n_{22k}}{n_{12k}n_{21k}}$$

The marginal odds ratio

$$\theta_{XY} = \frac{\mu_{11+}\mu_{22+}}{\mu_{12+}\mu_{21+}}$$

is estimated from the marginal table.

2.2.2 Simpson's Paradox

Some counter-intuitive but possible situations:

1. There are conditional associations ($\theta_{XY(k)} \neq 1$) but no marginal association ($\theta_{XY} = 1$)
2. There is a marginal association ($\theta_{XY} \neq 1$) but no conditional associations ($\theta_{XY(k)} = 1$)
3. **Simpson's paradox:** The conditional associations are in the opposite direction from the marginal, e.g.
 $\theta_{XY(k)} > 1, \theta_{XY} < 1$

	Full Population, $N = 52$			Men (M), $N = 20$			Women ($\neg M$), $N = 32$		
	Success (S)	Failure ($\neg S$)	Success Rate	Success	Failure	Success Rate	Success	Failure	Success Rate
Treatment (T)	20	20	50%	8	5	$\approx 61\%$	12	15	$\approx 44\%$
Control ($\neg T$)	6	6	50%	4	3	$\approx 57\%$	2	3	$\approx 40\%$

TABLE 1: Simpson's Paradox: the type of association at the population level (positive, negative, independent) changes at the level of subpopulations. Numbers taken from Simpson's original example (1951).

Figure 2.1: Simpson's paradox

2.2.3 Conditional Independence, Marginal Independence

Definition 2.7 (conditionally independent given Z , marginal independent)

We also call X and Y are **conditionally independent given $Z = k$** if $\theta_{XY(k)} = 1$. If this is true for all k , X and Y are **conditionally independent given Z** . Not the same to " X and Y are **marginal independent** if $\theta_{XY} = 1$ ".



Proposition 2.1

For multinomial sampling, can show that conditional independence is

$$\pi_{ijk} = \frac{\pi_{i+k}\pi_{+jk}}{\pi_{++k}}, \quad \forall i, j, k$$



2.2.4 Homogeneous Association

Definition 2.8

Let Z have K categories. X and Y have **homogeneous association** over Z if

$$\theta_{XY(1)} = \theta_{XY(2)} = \cdots = \theta_{XY(K)}$$

(Conditional independence is a special case.)

