

17: Odds Ratios from Case-Control Studies

Independent Samples

The prior chapter used cohort data to quantify exposure–disease relation with a risk ratio. This chapter uses data from case-control studies to quantify exposure-disease relations with odds ratios. We will discuss the sampling basis of case-control studies in lecture, along with some of its advantages and disadvantages. For details, please see pp. 208– 212 in *Epidemiology Kept Simple* (Gerstman, 2003).

The general idea is to select all cases that occur in the population. From the same source population, select a simple random sample of non-cases (controls). The cross-tabulate the data in the usual fashion:

Exposure variable	Response variable		Total
	+	–	
+	a_1	b_1	n_1
–	a_2	b_2	n_2
Total	m_1	m_2	N

Although we cannot calculate incidences or prevalences from these data (the sample was conditioned on the outcome), we can still determine the relation between the exposure and the disease by considering the following the odds of exposure in cases is $o_1 = A_1 / A_0$, the odds of exposure in controls is $o_0 = B_1 / B_0$ and the odds ratio (denoted OR or ψ) is

$$\psi = \frac{A_1 B_2}{A_2 B_1}$$

which is merely the cross-product ratio (i.e., multiply the diagonals cells and make them into a ratio). It can be shown that the odds ratio from a case-control study is stochastically equivalent to a relative incidence (i.e., relative risk) when the study is free of biases.

Let ψ (or OR) denote the odds ratio parameter and ψ^{\wedge} (or OR^{\wedge}) represent the odds ratio estimate (i.e., sample statistic). The **confidence interval for ψ** is

$$e^{\ln \hat{OR} \pm z \cdot SE_{\ln \hat{OR}}}$$

where e is the base on the natural logarithms ($e \approx 2.71828\dots$), z is a Standard Normal deviate corresponding to the desired level of confidence ($z = 1.645$ for 90% confidence, z

= 1.96 for 95% confidence, and $z = 2.576$ for 99% confidence), and

$$SE_{\ln \hat{\psi}} = \sqrt{\frac{1}{a_1} + \frac{1}{a_2} + \frac{1}{b_1} + \frac{1}{b_2}}.$$

A test of $H_0: \psi = 1$ is calculated with a chi-square statistic or Fisher's test, depending on the size of the sample (see prior chapter).

Example: Alcohol and esophageal cancer. Data from a case-control study of 200 esophageal cancer cases and 775 community-based controls are shown below.¹ Detailed dietary data were obtained by interview. This example addresses the relation between alcohol consumption (dichotomized at 80 grams per day) and esophageal cancer. Data are:

Alcohol g/day	Esophageal cancer		Total
	+	-	
+	96	109	205
-	104	666	770
Total	200	775	975

The odds ratio = $(96)(666)/(109)(104) = 5.6401 = 5.64$, suggesting esophageal cancer is 5.64 times as frequent in the exposed group in the source population.

To calculate confidence intervals, note that $\ln(\hat{\psi}) = \ln(5.640) = 1.7299$ (by calculator) and

standard error $SE_{\ln \hat{\psi}} = \sqrt{\frac{1}{96} + \frac{1}{104} + \frac{1}{109} + \frac{1}{666}} = 0.1752$. The 95%

confidence interval for the $\psi = e^{1.7299 \pm (1.96)(0.1752)} = e^{1.7299 \pm 0.3433} = e^{1.3866, 2.0732} = 4.00$ to 7.95 . The 90% confidence interval for the $\psi = e^{1.7299 \pm (1.645)(0.1752)} = e^{1.7299 \pm 0.2882} = e^{1.4417, 2.0181} = 4.23$ to 7.52 .

The P -value for testing $H_0: \psi = 1$ can be derived by chi-square test. $X^2_{\text{stat}} = 110.26$ and $X^2_{\text{stat, cont-corrected}} = 108.22$. Both have 1 df and $P \approx 0.00000$.

Results may be confirmed with SPSS (individual records), WinPepi or EpiCalc2000 (cross-tabulated data).

As always, the primary threats in practice are systematic errors (bias), not random, errors (imprecision).

Matched samples

A matched design may be used by both cohort and case-control studies to help control for extraneous factors. For cohort data, matched-pairs may be displayed as follows:

Exposed pair-member	Non-exposed pair-member		Total
	Case	Non-case	
Case	t	u	n_1
Non-case	v	w	n_2
Total	m_1	m_2	N

For case-control data, matched-pairs may be displayed as follows:

Case pair-member	Control pair-member		Total
	Exposed	Non-exposed	
Exposed	t	u	n_1
Non-exposed	v	w	n_2
Total	m_1	m_2	N

Note that data represent the numbers of pairs (not individuals).

Cells t and w in this table contain the number of **concordant pairs** in the sample. Concordant pairs are the same with respect to exposure. Cells u and v contain **discordant pairs**. *Discordant* pairs differ with respect to exposure. There are N pairs total, but we are interested in only the $(u + v)$ discordant pairs.

The **odds ratio** for these data is:

$$\hat{\psi} = \frac{u}{v}$$

The **confidence interval** for ψ is

$$e^{\ln \hat{\psi} \pm z \cdot SE_{\ln \hat{\psi}}}$$

where e is the base on the natural logarithms ($e \approx 2.71828\dots$), z is a Standard Normal deviate corresponding to the desired level of confidence ($z = 1.645$ for 90% confidence, $z = 1.96$ for 95% confidence, and $z = 2.576$ for 99% confidence), and $SE_{\ln \hat{\psi}} = \sqrt{\frac{1}{u} + \frac{1}{v}}$.

When the number of discordant pairs $(u + v)$ is 10 or greater, you can **test** of $H_0: \psi = 1$ with McNemar's chi-square. The regular and continuity-correct McNemar's chi-squares are shown below:

$$X_{\text{McN}}^2 = \frac{(u - v)^2}{u + v}$$

$$X_{\text{McN,cc}}^2 = \frac{(|u - v| - 1)^2}{u + v}$$

McNemar's chi-square statistics have 1 df.

Because of the relation between the chi-square distributions and z distributions, the above formulas can be re-expressed:

$$z_{\text{stat,McN}} = \sqrt{\frac{(u - v)^2}{u + v}}$$

$$z_{\text{stat,McNcc}} = \sqrt{\frac{(|u - v| - 1)^2}{u + v}}$$

With small samples, let the number of positive discordant pairs (u) be the numerator of a proportion and let the total number of discordant pairs ($u + v$) represent the denominator of a proportion. Then test, $H_0: p = 1/2$ with an exact binomial test (see Chapter 16 in the new biostat-text for details).

Example in cohort mode. *Smoking and mortality in identical twins.* When smoking was first suspected as a cause of disease, Sir Ronald Fisher (the world's greatest statistician, and a smoker), offered the *constitution hypothesis* as an explanation for the observed association. This hypothesis suggested that people genetically disposed to lung cancer were more likely to smoke. In other words, the relation between smoking and disease was *confounded* by constitutional factors. The constitutional hypothesis was put to the ultimate test by a study in which 22 smoking-discordant monozygotic twins were studied to see which twin first succumbed to death.² In this study, the smoking-twin died first in 17 of the pairs (i.e., $u = 17$, $u + v = 22$, so $v = 5$). Analyses derive:

- $\hat{OR} = \frac{u}{v} = \frac{17}{5} = 3.40$. The smoking twin was 3.4 as likely to die first.
- In testing, $H_0: OR = 1$, $z_{\text{stat,McN}} = \sqrt{\frac{(u - v)^2}{u + v}} = \sqrt{\frac{(17 - 5)^2}{17 + 5}} = 2.56$; $P = 0.010$. With continuity correction, $z_{\text{stat,McNcc}} = \sqrt{\frac{(|u - v| - 1)^2}{u + v}} = \sqrt{\frac{(|17 - 5| - 1)^2}{17 + 5}} = 2.35$; $P = 0.019$), providing "significant" support for the constitutional hypothesis.

Example in case-control mode. *Fruits, vegetables, and adenomatous polyps.* A case-control study used matched-pairs to study the risk of adenomatous polyps of the colon in relation to diet. Cases and controls had undergone sigmoidoscopic screening. Controls were matched to cases on time of screening, clinic, age, and sex. One of the study's analyses considered the effects of low fruit and vegetable consumption on colon polyp risk. There were 45 pairs in which the case but not the control reported low fruit/veggie consumption. There were 24 pairs in which the control but not the case reported low fruit/veggie consumption.³

- $\hat{OR} = \frac{u}{v} = \frac{45}{24} = 1.88$. Low fruit/veggie consumption is associated with an 88% increase in risk.
- $\ln(\hat{OR}) = 0.6286$ and $SE_{\ln \hat{\psi}} = \sqrt{\frac{1}{u} + \frac{1}{v}} = \sqrt{\frac{1}{45} + \frac{1}{24}} = 0.2528$
- 95% confidence interval for $OR = e^{0.6286 \pm (1.96)(0.2528)} = e^{0.6286 \pm 0.4959} = \psi = e^{(0.1331, 1.1241)} = (1.14, 3.07)$
- In testing, $H_0: OR = 1$, $z_{\text{stat,McN}} = \sqrt{\frac{(u-v)^2}{u+v}} = \sqrt{\frac{(45-24)^2}{45+24}} = 2.53$; $P = 0.011$.

With continuity correction, $z_{\text{stat,McNcc}} = \sqrt{\frac{(|u-v|-1)^2}{u+v}} = \sqrt{\frac{(|45-24|-1)^2}{45+24}} = 2.41$; $P = 0.016$.

References

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- ¹ Tuyns, A. J., Pequignot, G., & Jensen, O. M. (1977). [Esophageal cancer in Ille-et-Vilaine in relation to levels of alcohol and tobacco consumption. Risks are multiplying]. *Bulletin du Cancer*, 64(1), 45-60.
- ² Kaprio, J., & Koskenvuo, M. (1989). Twins, smoking and mortality: a 12-year prospective study of smoking-discordant twin pairs. *Social Science & Medicine*, 29(9), 1083-1089.
- ³ Witte, J. S., Longnecker, M. P., Bird, C. L., Lee, E. R., Frankl, H. D., & Haile, R. W. (1996). Relation of vegetable, fruit, and grain consumption to colorectal adenomatous polyps. *American Journal of Epidemiology*, 144(11), 1015-1025. Summary of frequencies reported in Rothman & Greenland, 1998, p. 287.