

Model Based Statistics in Biology.

Part V. The Generalized Linear Model.

Chapter 18.3 Single Factor. Retrospective Analysis

ReCap. Part I (Chapters 1,2,3,4), Part II (Ch 5, 6, 7)

ReCap Part III (Ch 9, 10, 11), Part IV (Ch 13, 14)

18 Binomial Response Variables

18.1 Logistic Regression (Dose-Response)

18.2 Single Factor. Prospective Analysis

18.3 Single Factor. Retrospective Analysis

18.4 Single Random Factor.

18.5 Single Explanatory Variable. Ordinal Scale.

18.6 Two Categorical Explanatory Variables

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Ch18.xls

on chalk board

ReCap Part I (Chapters 1,2,3,4) Quantitative reasoning

ReCap Part II (Chapters 5,6,7) Hypothesis testing and estimation

ReCap (Ch 9, 10,11) The General Linear Model with a single explanatory variable.

ReCap (Ch 12,13,14,15) GLM with more than one explanatory variable

ReCap (Ch 16,17). Generalized Linear Model. Poisson response variables.

ReCap (Ch 18) We used logistic regression to quantify the intensity of natural selection (Kettlewell data). This is called a prospective analysis. It is a longitudinal analysis because we followed individuals through time.

Today Retrospective analysis of binomial response across two levels of a single factor.

Wrap-up.

We used logistic regression to quantify the risk of cancer in smokers. The data were cross-sectional, in which we contrast groups with different histories at a single point in time. This is called retrospective analysis.

Binomial response variables –Retrospective Analysis.

Odds Ratio and Relative Risk

Does smoking increase the risk of cancer ?

What seems obvious today was once not obvious. Lung cancer emerged suddenly as a major health issue in the US in the middle of the 20th century. Shrek *et al.* (1950 reported higher rates of lung cancer in smokers than non-smokers from a veteran's hospital in Chicago. At the time of the Schrek study there was no knowledge of the causal agent in cigarettes that cause cancer. As a result there was continuing debate, driven by economics and public health concerns. The debates about cigarette smoking, in the 1950s, were similar in many ways to subsequent public debates about the effect on community health of adding fluoride to public water supplies, or the debates in this century about the safety of oil and gas extraction by hydraulic fracturing.

The Shrek *et al.* study was cross-sectional, comparing case rates at one point in time. It thus falls short of the rigor of a longitudinal study, which follows cases over time. A rigorous study requires a longitudinal (*i.e.*, prospective) study, such as the before-after control-impact (BACI) study, the gold standard in environmental impact assessment. In a prospective study, individuals who smoke would be carefully matched with non-smokers with similar characteristics (age, health status, *etc.*). This cohort is then followed over many years, to obtain the proportion of smokers that develop lung cancer, for comparison to the proportion of non-smokers that develop lung cancer. The result is rigorous, costly, and won't produce results for decades.

In a landmark publication, Jerome Cornfield showed that the relative risk in a population could be estimated from a case-control study. The data for such a study consist of patients with similar characteristics, sorted into those with and without the disease, to estimate the risk for those exposed and not exposed to a suspected cause. We then use the odds ratio to estimate the relative risk in the population. This is a retrospective (cross-sectional) study. Unlike a prospective study, we do not begin with a known set of cases, then score them at a later time as having or not having an attribute (disease). Instead, we collect cases at a single point in time and assign them to categories in a 2 by 2 table: having or lacking the disease, and having or lacking exposure to the suspected cause. The result is a sample that is clearly far from representative of the population. Cornfield showed that the bias in the sample was large, and could be corrected. The results for lung cancer were clear, and set in motion research that established cigarette tars as the causal agent for lung cancer. Further, the Cornfield publication established the mathematical basis for using case-control samples to estimate risk in a population.

To illustrate retrospective analysis in its modern form we will use the data presented by Cornfield (1951), even though this publication never mentions the odds ratio and was published well before modern methods for retrospective analysis (Breslow and Day 1980). Cornfield used percentage data and numbers (N) from Shrek *et al.* (1950) who reported data for US Army veterans with cancer, 35 with lung cancer and 171 with other cancers. The veterans were in the 40-49 age group, taken from records of over 5000 veterans presenting with tumors at Veteran's Administration hospitals in Chicago, from 1940 to 1942. The veterans in this age cohort were mostly born between 1891 and 1902, and so were veterans of World War I. Cigarettes were issued as a ration to US troops in World War I (Goodman (2005)). They were used as barter in the front lines, and were one of few reliefs from the psychological stress of trench warfare.

References

- Breslow, N.E. and Day, N.E. (1980). *Statistical Methods in Cancer Research. Volume I – The Analysis of Case-Control Studies*. Lyon: International Agency for Research on Cancer.
- Cornfield, J. (1951). A method of estimating comparative rates from clinical data. Application to cancer of the lung, breast and cervix. *Journal of the National Cancer Institute* 11: 1269–1275.
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- Schrek, R. Baker, L.A., Ballard, G.P., Dolgoff, S. 1950. Tobacco smoking as an etiologic factor in disease. I. Cancer. *Cancer Research* 10:49–58.
- Cook, T.D. 2002. Up with Odds Ratios! A case for odds ratios when outcomes are common. *Academic Emergency Medicine* 9:1430-4.
Argues that risk ratios can be easily misinterpreted and odds ratios may be more appropriate even with frequent outcomes.
- McNutt LA, Wu C, Xue X, Hafner JP. 2003 Estimating the relative risk in cohort studies and clinical trials of common outcomes. *Am J Epidemiol* 157(10):940-3.
These authors point out that when the outcome event is common (incidence of 10% or more), it is often more desirable to estimate an RR since there is an increasing differential between the RR and OR with increasing incidence rates, and there is a tendency for some to interpret ORs as if they are RRs.

Retrospective Analysis. Odds Ratios and Relative Risk.

Here are the percentages and cohort numbers (N) from Schrek *et al*, used by Cornfield (1951).

	Lung Tumors		
	Present	Absent	Total
heavy smokers	27	99	126
light smokers	8	72	80

For heavy smokers the odds are
 $27/99 = 0.27$ to 1

For light smokers the odds are $8/72 = 0.11$ to 1

The odds ratio is 2.45

The heavy smokers are more than twice as likely to have tumors as light smokers.

Retrospective Analysis. Odds Ratios and Relative Risk.

The odds ratio is also known as the cross-product ratio.

The cross product ratio for counts in a 2 X 2 table is defined as follows.

$$CPR = \frac{a}{b} \cdot \left(\frac{c}{d} \right)^{-1}$$

		Prospective Study	
		<u>Disease (Cancer)</u>	
		Present	Absent
<u>Exposure</u>	smokers	a	b
	non-smokers	c	d

Where the risk in the population is small, $1-a$ is large, the ratio $(1-a)/(1-c)$ approaches a value of 1, and the CPR is an estimate of the relative risk where the incidence rate of lung cancer is $155/999845 = 10^{-4}$ (Cornfield 1951). However, for the sample the incidence rate is $35/206 = 17\%$, higher than the recommended level of 10% for using the prevalence ratio to estimate the relative risk.

Here are the calculations for the Shrek *et al.* data.

	Lung Tumors			Odds	OR	Prevalence	Ratio
	Present	Absent	Total				
heavy smokers	27	99	126	0.27	2.45	0.21	2.14
light smokers	8	72	80	0.11		0.10	
Sum	35	171	206				
Tumor Odds	3.38	1.38					
OR	2.45			CPR	2.45		

The prevalence ratio is 2.14, slightly less than the odds ratio 2.45.

How good is the evidence?

Can the observed increase in odds be dismissed as mere chance?

We use the generic recipe for the Generalized Linear Model.

1. Model and data equations.

Verbal. Risk of tumors in male smokers age 40-49 increases relative to non-smokers.

Graphical Not useful. We have only 4 numbers.

Response variable: Odds of lung tumors

Explanatory variable: heavy smoking vs light smoking.

Note that Shrek *et al* reported percent smokers in two groups, those with and those without lung tumors.

Identifying the response and explanatory variable in a two-way table depends on the science. Do we expect the odds of lung tumors to depend on smoking? Or is it vice versa: smoking depends on lung tumors? Stating the alternatives, makes it clear that our interest is in the former.

Write formal model

Distribution $N_{\text{tumors}} \sim \text{Binomial}(N, \pi)$

Link Odds = e^η

$$\eta = \beta_{\text{ref}} + \beta_{\text{Smoke}} \cdot \text{Smoke}$$

$$e^{\beta_{\text{ref}}} = \text{Tumor odds, control group}$$

$$e^{\beta_{\text{Smoke}}} = \text{Odds ratio, case (heavy) vs control (light smoker)}$$

$$e^{\beta_{\text{ref}} + \beta_{\text{Smoke}} \cdot \text{Smoke}} = \text{Tumor odds, case group (heavy smokers)}$$

Place data in model format for generalized linear model routine.

The data can be listed by patient. In this example there would be 206 patients, each scored as LungTumor Y/N, and Smoke Light/Heavy.

Alternatively, the data in the twoway table is listed in 3 columns.

Ntmr	Ntot	Smoker	Column N	number of patients in each group	
27	126	Heavy	Column Ntmr		
8	80	Light	Column Smoker		

2. Execute analysis.

Code the model statement in a statistical package. Here is the logistic regression code in Minitab.

```
MTB > BLogistic 'Ntmr' 'Ntot' = 'Smoker';
SUBC> ST;
SUBC> Logit;
SUBC> Brief 2.                                            Minitab commands
```

Here is the model statement in a generalized linear model routine (SAS)

```
Proc Genmod; Classes Smoker;
  Model Ntmr/Ntot = Smoker/
    Link=logit dist=binomial type1 type3;           SAS command file
```

Here is the model statement in a generalized linear model routine (R)

```
glm(Ntmr/Ntot ~ Smoker,
     family = binomial(link = logit), data = Cornfield,
     weights = cases)                                     R code
```

For R/S+ we can recode the response variable $Ntumor$ to a proportion $ptumor$. Note the coding of exposure as 1 (heavy smoking)= and zero (light smoking).

Exposure			
Smoking	ptumor	cases	In(odds)
1	0.2143	126	-2.20
0	0.1000	80	-1.30

```
Call: glm(formula = ptumor ~ Smoking,
          family = binomial(link = logit), data = Cornfield,
          weights = cases)
```

Code generated by S+

The binomial approach based on 206 cases (0 or 1) with 205 degrees of freedom appears to be far better than the approach based on only 1 degree of freedom for the aggregated approach. However, we will be testing the improvement in fit due to only 1 parameter, which drops the degrees of freedom either from 205 to 204 df, or from 1 to 0 df. The improvement in fit will be the same for both approaches.

Residuals will be zero for the aggregated approach because this is a saturated model (two parameters and two observations).

Generalized linear model routines require a statement of:

- the error structure (binomial in this case)
- the link function (logit in this case).
- the structural model (explanatory variables).

2. Execute analysis.

Here are the parameter estimates.

	Value	Std. Error	t value	exp(Value)
(Intercept)	-2.20	0.3727	-5.90	0.111
Smoking	0.90	0.4313	2.08	2.455

The intercept is the logarithm of the odds for one of the cases. In this example the intercept is the log odds for light smokers because we listed light smoking as a lower value (zero) than heavy smoking (value of 1). The parameter estimate is $Odds = \exp(-2.20) = 0.111$. The smoking coefficient is the logarithm of the odds ratio for heavy smokers relative to light smokers. The parameter estimate is $OR = \exp(0.9) = 2.455$

3. Use parameter estimates to calculate residuals, evaluate model.

Residuals are all zero, so we can't use them for evaluation. We assume that the $35+171 = 206$ trials were independent events. That is, developing a lung tumor does not depend on the chance of another participant in the study developing a lung tumor.

4. What is the evidence?

From the ANODEV table we obtain the improvement in fit.

From this we calculate the LR.

$$LR = e^{4.814/2} = 11.1$$

	Df	Deviance	Resid df	Residual Deviance
NULL			1	4.814
Exposure	1	4.814	0	0

There is at best weak evidence ($20 > LR > 10$) for greater odds of tumors in heavy than light smokers.

5. Decide on mode of inference. Is hypothesis testing appropriate?

We will use confidence limits to evaluate uncertainty on the evidence reported by Cornfield (1951) for the Schrek *et al.* (1950) data.

10. Analysis of parameters of biological interest.

Odds for light smokers

$$e^{\beta_0} = e^{-2.2} = 0.111$$

Odds ratio for heavy relative to light smokers $e^{\beta_{Smoke}} = e^{0.9} = 2.455$

Confidence limits for β_{Smoke} are $\hat{\beta}_{Smoke} \pm 1.96 \cdot sterr$

The standard error on the estimate of $\hat{\beta}_{Smoke}$ was 0.4313.

The confidence interval is from 0.0526 to 1.74

The confidence interval for the *OR* is $\exp(0.0526)$ to $\exp(1.74)$,

i.e. from 1.05 to 5.1

We can exclude the null hypothesis (*OR* = 1) and we can exclude odds ratios greater than 5 times higher for heavy smokers.

In a decision-theoretic context we can reject the null hypothesis (*OR* = 1) at a 5% limit on Type I error.

G = 4.814

p = 0.029 from a χ^2 distribution with a single degree of freedom.

We have adequate certainty (p = 0.029) on weak evidence (LR = 11) for greater risk of lung cancer in this sample of veterans. A larger sample is needed to strengthen the evidence.

Cornfield's 1951 publication showed how to infer from the sample to a larger population, all males age 40-49 in the American midwest in 1940-42. At that time, the risk of developing lung cancer for this population was 155 in a million.

The sample (veterans with cancer of all types) is biased relative to the population. To correct the bias, Cornfield used the population disease risk (155 per million) to recompute the relative risk for the population.

Subsequent publications in the medical and health sciences routinely list characteristics of the sample (age, gender, etc) as a guide to the relevant population. They rarely report the disease risk in the population (as Cornfield did) or the exposure risk. Cornfield corrected for bias in the disease risk, but could not correct for exposure risk in the sample, compared to the population. A higher exposure risk would be expected for the sample (WW I veterans) given free access to a highly addictive substance (nicotine in cigarettes) in a war zone.

Population Risk Calculations.

Because lung cancer risk in the population was small (155 in a million) Cornfield argued that the relative risk in the sample could be used to estimate the relative risk in the population. Here are Cornfield's recalculated proportions for lung tumor presence/absence in light and heavy smokers in the population, after applying the disease risk of 155 per million for the same age cohort in the population. Odds and odds ratios, which Cornfield did not use, have been added.

	Lung tumors		Exposure			
	Present	Absent	Odds	OR	Risk	RelRisk
heavy smokers	0.00011935	0.579910	2.06E-04	2.42	1.E-04	3.35
light smokers	0.00003565	0.419935	8.49E-05		4.E-05	
population	0.00015500	0.999845	1.55E-04			
Disease Odds		3.35	1.38			
Disease OR		2.42				
Disease Risk	0.00011935	0.579910				
RelRisk	0.00020581					

Veterans smoke more than non-veterans and thus the sample is not representative of the population with respect to exposure. A correction similar to that for disease risk in the population could also be applied, to improve the inference from the case-control sample to the population.

Binomial response variables –Retrospective Analysis. Multiple categories.

Cornfield considered only two categories, light and heavy smoking.

What is the risk relative to non-smokers?

Does number of cigarettes/day increase risk ?

Here are data from a cross-sectional (case-control) study by Zang and Wynder (1992 *Cancer* 70: 69-76) who report frequency of tumors in 2225 subjects at 5 levels of cigarette smoking. cf Sokal and Rohlf 1995, Exercise 17.20

	Lung Cancer (males)			%	odds of cancer	odds ratio
	Present	absent	total			
non-smokers	15	822	837	1.79%	0.018 : 1	
1-10 cig	36	136	172	20.9%	0.265 : 1	14.51
11-20 cig	133	328	461	28.9%	0.405 : 1	22.22
21-40 cig	226	311	537	42.1%	0.727 : 1	39.82
>41 cig	127	91	218	58.3%	1.396 : 1	76.48

The odds appear to increase substantially, depending on level of smoking.

Here is the result for a classical goodness of fit test. The null hypothesis is the expected proportion: 537 with cancer / 2225 subjects = 0.24.

f	$= \hat{p} \cdot N_i$	+	residual	$2 \ln L = 2f \ln(f / \hat{p} \cdot N_i)$
15	$= 0.24 \cdot 837$	-	187	-78
36	$= 0.24 \cdot 172$	-	6	-10
133	$= 0.24 \cdot 461$	+	22	47
226	$= 0.24 \cdot 537$	+	96	251
127	$= 0.24 \cdot 518$	+	74	224

$$G^2 = -2 \sum f \ln(f / \hat{p}N_i) = 434$$

Here is the ANODEV table for the Zang and Wynder data.

Source	LR Statistics For Type 1 Analysis			Pr > ChiSq
	Deviance	DF	Chi-Square	
Intercept	551.2222			
Smoke	0.0000	4	551.22	<.0001

SAS output file

The goodness of fit of the null model to the data is $G^2 = 551.2$

The fit of the alternative model to the data is perfect $G^2 = 0.0$

The improvement is $\Delta G^2 = 551.2$

Do you get the same result?

Compare the odds for smokers and non-smokers, as an odds ratio.

Calculate the confidence limits on this odds ratio, and interpret.