

Low Birth Weight

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Discussion of the Problem

The data contain information about infant mortality in 2003 and were obtained from the US National Center for Health Statistics. A random sample of 2,500+ observations is used in this example. This data are observational, in which case, meaningful inference is limited. The description below is for a causal inference example, which is beyond the scope of this course, given in SAS.

Our approach is to investigate this problem using the material given in the first part of Chapter 3 in the methods lecture notes.

The main variables in the analysis are as follows:

- The treatment variable is **Smoking**. It is an indicator of maternal smoking behavior, with values Yes and No.
- The outcome variable is **Death**. It is an indicator of infant death within one year of birth, with values Yes and No.
- The mediator variable is **LowBirthWgt**. It is an indicator of low birth weight (less than 2,500 grams), with values Yes and No.

The analysis also includes five confounding covariates:

- **AgeGroup** represents maternal ages of less than 20, between 20 and 35, and greater than 35, with values 1, 2, and 3, respectively.

- **Drinking** is an indicator of maternal drinking during pregnancy, with values Yes and No.
- **Married** is an indicator of marital status, with values Yes and No.
- **Race** is an indicator of race, with values Asian, Black, Hispanic, Native (native American), and White.
- **SomeCollege** is an indicator of whether the mother has 12 or more years of education, with values Yes and No.

R

Needed Packages

```
if(!require(FSA)){install.packages("FSA")}
if(!require(ggplot2)){install.packages("ggplot2")}
if (!require("mosaic")) install.packages("mosaic", dep=FALSE)
if (!require("nortest")) install.packages("nortest", dep=TRUE)
if (!require("epitools")) install.packages("epitools", dep=TRUE)
if (!require("prettyR")) install.packages("prettyR", dep=TRUE)
if (!require("rms")) install.packages("rms", dep=TRUE)
# add other as needed
```

Problem – Infant Birth Weight Data

Read data from SAS input file

```
# this data came from SASHELP.BWEIGHT
bw = read.csv('bwgt.csv', header = TRUE)
bw = data.frame(bw)
#summary(bw)
bw = transform(bw, AgeGroup.f = as.factor(AgeGroup))
bw = transform(bw, Race.f = as.factor(Race))
bw = transform(bw, Drinking.f = as.factor(Drinking))
bw = transform(bw, Death.f = as.factor(Death))
bw = transform(bw, Smoking.f = as.factor(Smoking))
bw = transform(bw, SomeCollege.f = as.factor(SomeCollege))
bw = transform(bw, LowBirthWgt.f = as.factor(LowBirthWgt))
```

```
tally(~ AgeGroup + Race.f, data=bw)
```

```
##           Race.f
## AgeGroup Asian Black Hispanic Native White
##      1      8    91      83      6   169
##      2    101   375     475     22  1337
##      3     36    52      66      4   264
```

```
tally(~ Race.f | AgeGroup.f, data=bw)
```

```
##           AgeGroup.f
## Race.f           1    2    3
## Asian           8  101   36
## Black          91  375   52
## Hispanic       83  475   66
## Native         6   22    4
## White        169 1337  264
```

```
library(mosaic)
mytab = tally(~ Race.f | AgeGroup.f, data=bw)
addmargins(mytab)
```

```
##           AgeGroup.f
## Race.f      1      2      3 Sum
##   Asian      8    101    36 145
##   Black     91    375    52 518
##   Hispanic  83    475    66 624
##   Native     6     22     4  32
##   White    169   1337   264 1770
##   Sum       357   2310   422 3089
```

```
prop.table(mytab, 1)
```

```
##           AgeGroup.f
## Race.f      1      2      3
##   Asian  0.05517241 0.69655172 0.24827586
##   Black  0.17567568 0.72393822 0.10038610
##   Hispanic 0.13301282 0.76121795 0.10576923
##   Native  0.18750000 0.68750000 0.12500000
##   White  0.09548023 0.75536723 0.14915254
```

```
library(epitools)
attach(bw)
mytab = tally(~ LowBirthWgt.f | Death.f, data=bw)
addmargins(mytab)
```

```
##           Death.f
## LowBirthWgt.f  No  Yes  Sum
##           No  2278  198 2476
##           Yes   205  408  613
##           Sum  2483  606 3089
```

```
prop.table(mytab, 1)
```

```
##           Death.f
## LowBirthWgt.f      No      Yes
##           No  0.92003231 0.07996769
##           Yes  0.33442088 0.66557912
```

```
riskratio(x=Smoking.f, y=Death.f)
```

```
## $data
##           Outcome
## Predictor    No Yes Total
##           156  41   197
##      No    1786 405  2191
##      Yes     541 160   701
##      Total 2483 606  3089
##
## $measure
##           risk ratio with 95% C.I.
## Predictor estimate      lower      upper
##           1.0000000      NA      NA
##      No  0.8881678 0.6670955 1.182502
##      Yes 1.0966911 0.8087967 1.487063
```

```
##
## $p.value
##          two-sided
## Predictor midp.exact fisher.exact chi.square
##          NA          NA          NA
##      No   0.4206715   0.4449380   0.4220298
##      Yes  0.5556210   0.6286442   0.5493629
##
## $correction
## [1] FALSE
##
## attr("method")
## [1] "Unconditional MLE & normal approximation (Wald) CI"
```

SAS

Code 1

```
/*
The Sashelp.BirthWgt data set contains 100,000 random observations about
infant mortality in 2003 from the US National Center for Health Statistics.
Each observation records infant death within one year of birth, birth weight,
maternal smoking and drinking behavior, and other background characteristics
of the mother.
*/

title "Sashelp.bweight --- Infant Birth Weight";
data birthwgt; set sashelp.birthwgt;
run;

proc contents data=birthwgt varnum;
ods select position;
run;

title "The First Five Observations Out of 100,000";
proc print data=birthwgt(obs=10);
run;
```

Sashelp.bweight — Infant Birth Weight

The CONTENTS Procedure

Variables in Creation Order			
#	Variable	Type	Len
1	LowBirthWgt	Char	3
2	Married	Char	3
3	AgeGroup	Num	8
4	Race	Char	9
5	Drinking	Char	3
6	Death	Char	3
7	Smoking	Char	3
8	SomeCollege	Char	3

The First Five Observations Out of 100,000

<i>Obs</i>	<i>LowBirthWgt</i>	<i>Married</i>	<i>AgeGroup</i>	<i>Race</i>	<i>Drinking</i>	<i>Death</i>	<i>Smoking</i>	<i>SomeCollege</i>
1	No	No	3	Asian	No	No	No	Yes
2	No	No	2	White	No	No	No	No
3	Yes	Yes	2	Native	No	Yes	No	No
4	No	No	2	White	No	No	No	No
5	No	No	2	White	No	No	No	Yes
6	No	No	2	White	No	No	No	
7	No	No	2	Asian	No	No	No	Yes
8	No	No	3	White	No	No	No	Yes
9	No	Yes	1	Black	No	No	No	No
10	No	No	2	Native	No	No	No	Yes

Code 2

I have changed 'Yes' responses to 'Affirm' as SAS orders the variables in the tables using an alphabetical ordering. This new order allows one to have a better interpretation of results.

```

*Create a new smaller data set;
title 'New Sample of Size 2,500';
proc surveyselect data=birthwgt out=new2 method=srs n=2500
    seed=2021;
run;

/* I needed more death records than the srs gave me */
data new; set birthwgt; if death = 'Yes';
run;

/*merge the two files into one */
data new_bwgt; set new new2;
run;

data new_bwgt; set new_bwgt;
if LowBirthWgt = 'Yes' then LowBirthWgt = 'Affirm';
if Death = 'Yes' then Death = 'Affirm';
if Smoking = 'Yes' then Smoking = 'Affirm';
if Drinking = 'Yes' then Drinking = 'Affirm';

title 'Test for Association between Low Birth Weight and Smoking';
proc freq data=new_bwgt;* order=freq;
tables smoking*LowBirthWgt/norow nopercnt chisq relrisk riskdiff;
run;

title 'Test for Association between Low Birth Weight and drinking';
proc freq data=new_bwgt;* order=freq;
tables drinking*LowBirthWgt/norow nopercnt chisq relrisk riskdiff;
run;

```

New Sample of Size 2,500

The SURVEYSELECT Procedure

<i>Selection Method</i>	Simple Random Sampling
-------------------------	------------------------

<i>Input Data Set</i>	BIRTHWGT
<i>Random Number Seed</i>	2021
<i>Sample Size</i>	2500
<i>Selection Probability</i>	0.025
<i>Sampling Weight</i>	40
<i>Output Data Set</i>	NEW2

Test for Association between Low Birth Weight and Smoking

The FREQ Procedure

<i>Table of Smoking by LowBirthWgt</i>			
<i>Smoking</i>	<i>LowBirthWgt</i>		
	<i>Aff</i>	<i>No</i>	<i>Total</i>
<i>Aff</i>	155 26.96	546 23.56	701
<i>No</i>	420 73.04	1771 76.44	2191
<i>Total</i>	575	2317	2892
<i>Frequency Missing = 197</i>			

Note	Statistics for Table of Smoking by LowBirthWgt
------	--

In the following table there is not a significant association at the .05 level between Low Birth Weight and Smoking. This is seen in the chi-square statistic and the relative risk and odds ratio.

<i>Statistic</i>	<i>DF</i>	<i>Value</i>	<i>Prob</i>
<i>Chi-Square</i>	1	2.8856	0.0894
<i>Likelihood Ratio Chi-Square</i>	1	2.8341	0.0923
<i>Continuity Adj. Chi-Square</i>	1	2.7038	0.1001
<i>Mantel-Haenszel Chi-Square</i>	1	2.8846	0.0894
<i>Phi Coefficient</i>		0.0316	
<i>Contingency Coefficient</i>		0.0316	
<i>Cramer's V</i>		0.0316	

<i>Fisher's Exact Test</i>	
<i>Cell (1,1) Frequency (F)</i>	155
<i>Left-sided Pr ≤ F</i>	0.9593
<i>Right-sided Pr ≥ F</i>	0.0510
<i>Table Probability (P)</i>	0.0102
<i>Two-sided Pr ≤ P</i>	0.0921

<i>Column 1 Risk Estimates</i>						
	<i>Risk</i>	<i>ASE</i>	<i>95% Confidence Limits</i>		<i>Exact 95% Confidence Limits</i>	
<i>Row 1</i>	0.2211	0.0157	0.1904	0.2518	0.1909	0.2537
<i>Row 2</i>	0.1917	0.0084	0.1752	0.2082	0.1754	0.2088
<i>Total</i>	0.1988	0.0074	0.1843	0.2134	0.1844	0.2138
<i>Difference</i>	0.0294	0.0178	−0.0054	0.0643		
<i>Difference is (Row 1 - Row 2)</i>						

<i>Column 2 Risk Estimates</i>						
	<i>Risk</i>	<i>ASE</i>	<i>95% Confidence Limits</i>		<i>Exact 95% Confidence Limits</i>	
<i>Row 1</i>	0.7789	0.0157	0.7482	0.8096	0.7463	0.8091
<i>Row 2</i>	0.8083	0.0084	0.7918	0.8248	0.7912	0.8246
<i>Total</i>	0.8012	0.0074	0.7866	0.8157	0.7862	0.8156
<i>Difference</i>	−0.0294	0.0178	−0.0643	0.0054		
<i>Difference is (Row 1 - Row 2)</i>						

<i>Odds Ratio and Relative Risks</i>			
<i>Statistic</i>	<i>Value</i>	<i>95% Confidence Limits</i>	
<i>Odds Ratio</i>	1.1970	0.9725	1.4734
<i>Relative Risk (Column 1)</i>	1.1535	0.9796	1.3582
<i>Relative Risk (Column 2)</i>	0.9636	0.9218	1.0074

Note	Sample Size = 2892 Frequency Missing = 197
------	---

Test for Association between Low Birth Weight and drinking

In the following table there is not a significant association at the .05 level between Low Birth Weight and Drinking. This is seen in the chi-square statistic and the relative risk and odds ratio.

The FREQ Procedure

Table of Drinking by LowBirthWgt			
Drinking	LowBirthWgt		
	Aff	No	Total
Aff	74 12.87	325 14.03	399
No	501 87.13	1992 85.97	2493
Total	575	2317	2892
Frequency Missing = 197			

Note	Statistics for Table of Drinking by LowBirthWgt
------	---

Statistic	DF	Value	Prob
Chi-Square	1	0.5187	0.4714
Likelihood Ratio Chi-Square	1	0.5263	0.4682
Continuity Adj. Chi-Square	1	0.4260	0.5140
Mantel-Haenszel Chi-Square	1	0.5185	0.4715
Phi Coefficient		−0.0134	
Contingency Coefficient		0.0134	
Cramer's V		−0.0134	

Fisher's Exact Test	
Cell (1,1) Frequency (F)	74
Left-sided Pr ≤ F	0.2588
Right-sided Pr ≥ F	0.7835
Table Probability (P)	0.0423
Two-sided Pr ≤ P	0.4998

Column 1 Risk Estimates						
	Risk	ASE	95% Confidence Limits		Exact 95% Confidence Limits	
Row 1	0.1855	0.0195	0.1473	0.2236	0.1485	0.2271
Row 2	0.2010	0.0080	0.1852	0.2167	0.1854	0.2172
Total	0.1988	0.0074	0.1843	0.2134	0.1844	0.2138
Difference	−0.0155	0.0210	−0.0568	0.0258		
Difference is (Row 1 - Row 2)						

<i>Column 2 Risk Estimates</i>						
	<i>Risk</i>	<i>ASE</i>	<i>95% Confidence Limits</i>		<i>Exact 95% Confidence Limits</i>	
<i>Row 1</i>	0.8145	0.0195	0.7764	0.8527	0.7729	0.8515
<i>Row 2</i>	0.7990	0.0080	0.7833	0.8148	0.7828	0.8146
<i>Total</i>	0.8012	0.0074	0.7866	0.8157	0.7862	0.8156
<i>Difference</i>	0.0155	0.0210	−0.0258	0.0568		
<i>Difference is (Row 1 - Row 2)</i>						

<i>Odds Ratio and Relative Risks</i>			
<i>Statistic</i>	<i>Value</i>	<i>95% Confidence Limits</i>	
<i>Odds Ratio</i>	0.9053	0.6906	1.1869
<i>Relative Risk (Column 1)</i>	0.9229	0.7406	1.1500
<i>Relative Risk (Column 2)</i>	1.0194	0.9689	1.0725

Note	Sample Size = 2892 Frequency Missing = 197
------	---

Code 3

```

title 'Test for Association between Low Birth Weight and Smoking';
title2 'Controlling for Death';
proc freq data=new_bwgt;* order=freq;
tables death*smoking*LowBirthWgt /nopercent norow chisq cmh;
run;

title 'Test for Association between Low Birth Weight and Drinking';
title2 'Controlling for Death';
proc freq data=new_bwgt;* order=freq;
tables death*drinking*LowBirthWgt /nopercent norow chisq cmh;
run;

```

Test for Association between Low Birth Weight and Smoking

Controlling for Death

The FREQ Procedure

Table 1 of Smoking by LowBirthWgt			
Controlling for Death=Aff			
Smoking	LowBirthWgt		
	Aff	No	Total
Aff	102 26.63	58 31.87	160
No	281 73.37	124 68.13	405
Total	383	182	565
Frequency Missing = 41			

Note	Statistics for Table 1 of Smoking by LowBirthWgt Controlling for Death=Aff
------	---

Statistic	DF	Value	Prob
Chi-Square	1	1.6664	0.1967
Likelihood Ratio Chi-Square	1	1.6467	0.1994
Continuity Adj. Chi-Square	1	1.4185	0.2337
Mantel-Haenszel Chi-Square	1	1.6635	0.1971
Phi Coefficient		−0.0543	
Contingency Coefficient		0.0542	
Cramer's V		−0.0543	

Fisher's Exact Test	
Cell (1,1) Frequency (F)	102
Left-sided Pr ≤ F	0.1172
Right-sided Pr ≥ F	0.9172

<i>Fisher's Exact Test</i>	
<i>Table Probability (P)</i>	0.0344
<i>Two-sided Pr <= P</i>	0.2304

Note	Sample Size = 565 Frequency Missing = 41
------	---

<i>Table 2 of Smoking by LowBirthWgt</i>			
<i>Controlling for Death=No</i>			
<i>Smoking</i>	<i>LowBirthWgt</i>		
	<i>Aff</i>	<i>No</i>	<i>Total</i>
<i>Aff</i>	53 27.60	488 22.86	541
<i>No</i>	139 72.40	1647 77.14	1786
<i>Total</i>	192	2135	2327
<i>Frequency Missing = 156</i>			

Note	Statistics for Table 2 of Smoking by LowBirthWgt Controlling for Death=No
------	--

<i>Statistic</i>	<i>DF</i>	<i>Value</i>	<i>Prob</i>
<i>Chi-Square</i>	1	2.2246	0.1358
<i>Likelihood Ratio Chi-Square</i>	1	2.1450	0.1430
<i>Continuity Adj. Chi-Square</i>	1	1.9666	0.1608
<i>Mantel-Haenszel Chi-Square</i>	1	2.2237	0.1359
<i>Phi Coefficient</i>		0.0309	
<i>Contingency Coefficient</i>		0.0309	
<i>Cramer's V</i>		0.0309	

<i>Fisher's Exact Test</i>	
<i>Cell (1,1) Frequency (F)</i>	53
<i>Left-sided Pr <= F</i>	0.9410
<i>Right-sided Pr >= F</i>	0.0821
<i>Table Probability (P)</i>	0.0231
<i>Two-sided Pr <= P</i>	0.1532

Note	Sample Size = 2327 Frequency Missing = 156
------	---

Test for Association between Low Birth Weight and Smoking

Controlling for Death

The FREQ Procedure

Note	Summary Statistics for Smoking by LowBirthWgt Controlling for Death
------	--

<i>Cochran-Mantel-Haenszel Statistics (Based on Table Scores)</i>				
<i>Statistic</i>	<i>Alternative Hypothesis</i>	<i>DF</i>	<i>Value</i>	<i>Prob</i>
1	Nonzero Correlation	1	0.0640	0.8003
2	Row Mean Scores Differ	1	0.0640	0.8003
3	General Association	1	0.0640	0.8003

<i>Common Odds Ratio and Relative Risks</i>				
<i>Statistic</i>	<i>Method</i>	<i>Value</i>	<i>95% Confidence Limits</i>	
<i>Odds Ratio</i>	Mantel–Haenszel	1.0328	0.8004	1.3327
	Logit	1.0374	0.8066	1.3341
<i>Relative Risk (Column 1)</i>	Mantel–Haenszel	1.0170	0.8909	1.1610
	Logit	0.9675	0.8563	1.0932
<i>Relative Risk (Column 2)</i>	Mantel–Haenszel	0.9954	0.9599	1.0323
	Logit	0.9809	0.9513	1.0114

<i>Breslow-Day Test for Homogeneity of Odds Ratios</i>	
<i>Chi-Square</i>	3.8091
<i>DF</i>	1
<i>Pr > ChiSq</i>	0.0510

Note	Sample Size = 2892 Frequency Missing = 197
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Test for Association between Low Birth Weight and Drinking

Controlling for Death

The FREQ Procedure

Table 1 of Drinking by LowBirthWgt			
Controlling for Death=Aff			
Drinking	LowBirthWgt		
	Aff	No	Total
Aff	45 11.75	25 13.74	70
No	338 88.25	157 86.26	495
Total	383	182	565
Frequency Missing = 41			

Note	Statistics for Table 1 of Drinking by LowBirthWgt Controlling for Death=Aff
------	--

Statistic	DF	Value	Prob
Chi-Square	1	0.4487	0.5029
Likelihood Ratio Chi-Square	1	0.4419	0.5062
Continuity Adj. Chi-Square	1	0.2843	0.5939
Mantel-Haenszel Chi-Square	1	0.4479	0.5033
Phi Coefficient		-0.0282	
Contingency Coefficient		0.0282	
Cramer's V		-0.0282	

Fisher's Exact Test	
Cell (1,1) Frequency (F)	45
Left-sided Pr ≤ F	0.2940
Right-sided Pr ≥ F	0.7912
Table Probability (P)	0.0852
Two-sided Pr ≤ P	0.4976

Note	Sample Size = 565 Frequency Missing = 41
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<i>Table 2 of Drinking by LowBirthWgt</i>			
<i>Controlling for Death=No</i>			
<i>Drinking</i>	<i>LowBirthWgt</i>		
	<i>Aff</i>	<i>No</i>	<i>Total</i>
<i>Aff</i>	29 15.10	300 14.05	329
<i>No</i>	163 84.90	1835 85.95	1998
<i>Total</i>	192	2135	2327
<i>Frequency Missing = 156</i>			

Note	Statistics for Table 2 of Drinking by LowBirthWgt Controlling for Death=No
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<i>Statistic</i>	<i>DF</i>	<i>Value</i>	<i>Prob</i>
<i>Chi-Square</i>	1	0.1608	0.6884
<i>Likelihood Ratio Chi-Square</i>	1	0.1581	0.6909
<i>Continuity Adj. Chi-Square</i>	1	0.0858	0.7696
<i>Mantel-Haenszel Chi-Square</i>	1	0.1607	0.6885
<i>Phi Coefficient</i>		0.0083	
<i>Contingency Coefficient</i>		0.0083	
<i>Cramer's V</i>		0.0083	

<i>Fisher's Exact Test</i>	
<i>Cell (1,1) Frequency (F)</i>	29
<i>Left-sided Pr <= F</i>	0.7002
<i>Right-sided Pr >= F</i>	0.3773
<i>Table Probability (P)</i>	0.0775
<i>Two-sided Pr <= P</i>	0.6660

Note	Sample Size = 2327 Frequency Missing = 156
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Test for Association between Low Birth Weight and Drinking

Controlling for Death

The FREQ Procedure

Note	Summary Statistics for Drinking by LowBirthWgt Controlling for Death
------	---

<i>Cochran-Mantel-Haenszel Statistics (Based on Table Scores)</i>				
<i>Statistic</i>	<i>Alternative Hypothesis</i>	<i>DF</i>	<i>Value</i>	<i>Prob</i>
1	Nonzero Correlation	1	0.0102	0.9194
2	Row Mean Scores Differ	1	0.0102	0.9194
3	General Association	1	0.0102	0.9194

<i>Common Odds Ratio and Relative Risks</i>				
<i>Statistic</i>	<i>Method</i>	<i>Value</i>	<i>95% Confidence Limits</i>	
<i>Odds Ratio</i>	Mantel-Haenszel	0.9834	0.7080	1.3659
	Logit	0.9836	0.7110	1.3609
<i>Relative Risk (Column 1)</i>	Mantel-Haenszel	0.9908	0.8261	1.1883
	Logit	0.9668	0.8191	1.1413
<i>Relative Risk (Column 2)</i>	Mantel-Haenszel	1.0021	0.9611	1.0449
	Logit	0.9942	0.9592	1.0305

<i>Breslow-Day Test for Homogeneity of Odds Ratios</i>	
<i>Chi-Square</i>	0.5994
<i>DF</i>	1
<i>Pr > ChiSq</i>	0.4388

Note	Sample Size = 2892 Frequency Missing = 197
------	---

Code 4

```

title 'Test for Association between Low Birth Weight and Death';
title2 '';
proc freq data=new_bwgt;* order=freq;
tables LowBirthWgt*death/norow nopercent chisq relrisk riskdiff;
run;

ods latex close;

```

Test for Association between Low Birth Weight and Death

In the following table there is a significant association at the .05 level between Death and Low Birth Weight. This is seen in the chi-square statistic and the relative risk and odds ratio.

The FREQ Procedure

<i>Table of LowBirthWgt by Death</i>			
<i>LowBirthWgt</i>	<i>Death</i>		
	<i>Aff</i>	<i>No</i>	<i>Total</i>
<i>Aff</i>	408 67.33	205 8.26	613
<i>No</i>	198 32.67	2278 91.74	2476
<i>Total</i>	606	2483	3089

Note	Statistics for Table of LowBirthWgt by Death
------	--

<i>Statistic</i>	<i>DF</i>	<i>Value</i>	<i>Prob</i>
<i>Chi-Square</i>	1	1068.5596	<.0001
<i>Likelihood Ratio Chi-Square</i>	1	897.1241	<.0001
<i>Continuity Adj. Chi-Square</i>	1	1064.8493	<.0001
<i>Mantel-Haenszel Chi-Square</i>	1	1068.2137	<.0001
<i>Phi Coefficient</i>		0.5882	
<i>Contingency Coefficient</i>		0.5070	
<i>Cramer's V</i>		0.5882	

<i>Fisher's Exact Test</i>	
<i>Cell (1,1) Frequency (F)</i>	408
<i>Left-sided Pr <= F</i>	1.0000
<i>Right-sided Pr >= F</i>	<.0001
<i>Table Probability (P)</i>	<.0001
<i>Two-sided Pr <= P</i>	<.0001

<i>Column 1 Risk Estimates</i>						
	<i>Risk</i>	<i>ASE</i>	<i>95% Confidence Limits</i>		<i>Exact 95% Confidence Limits</i>	
<i>Row 1</i>	0.6656	0.0191	0.6282	0.7029	0.6267	0.7029
<i>Row 2</i>	0.0800	0.0055	0.0693	0.0907	0.0696	0.0914
<i>Total</i>	0.1962	0.0071	0.1822	0.2102	0.1823	0.2106
<i>Difference</i>	0.5856	0.0198	0.5468	0.6245		
<i>Difference is (Row 1 - Row 2)</i>						

<i>Column 2 Risk Estimates</i>						
	<i>Risk</i>	<i>ASE</i>	<i>95% Confidence Limits</i>		<i>Exact 95% Confidence Limits</i>	
<i>Row 1</i>	0.3344	0.0191	0.2971	0.3718	0.2971	0.3733
<i>Row 2</i>	0.9200	0.0055	0.9093	0.9307	0.9086	0.9304
<i>Total</i>	0.8038	0.0071	0.7898	0.8178	0.7894	0.8177
<i>Difference</i>	-0.5856	0.0198	-0.6245	-0.5468		
<i>Difference is (Row 1 - Row 2)</i>						

Note all the confidence intervals do not contain one, indicating a strong association between infant birth weight and survival.

<i>Odds Ratio and Relative Risks</i>			
<i>Statistic</i>	<i>Value</i>	<i>95% Confidence Limits</i>	
<i>Odds Ratio</i>	22.8979	18.3410	28.5869
<i>Relative Risk (Column 1)</i>	8.3231	7.2003	9.6210
<i>Relative Risk (Column 2)</i>	0.3635	0.3249	0.4067

Note	Sample Size = 3089
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Loglinear Models for Contingency Tables

I do not normally cover this material until the spring semester. It is covered in greater detail in our graduate course on Categorical Models. Yet, this example is ideally suited for this approach. Agresti covers this material in Chapter 8 and SDK covers the material in chapter 16.

Two-Way Tables

Loglinear Models for the 2×2 Table

Suppose that one has the 2×2 table given by

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

With cell probabilities given by

X	Y=1	Y=2	Total
1	π_{11}	π_{12}	π_{1+}
2	π_{21}	π_{22}	π_{2+}
Total	π_{+1}	π_{+2}	1

One of the foundational issues in analysis of these tables is to test for independence of the two random variables X and Y . Independence implies that,

$$\frac{\pi_{11}}{\pi_{+1}} = \frac{\pi_{12}}{\pi_{+2}} = \pi_{1+}$$

and

$$\pi_{11} = \pi_{1+}\pi_{+1}.$$

From which it follows that

$$\pi_{ij} = \pi_{i+}\pi_{+j} \quad i, j = 1, 2.$$

If X and Y are independent then it follows that the odds ratio, ψ , is

$$\psi = \frac{\pi_{11}\pi_{22}}{\pi_{12}\pi_{21}} = 1.$$

Taking the log of both sides leads to,

$$\log \psi = \log \pi_{11} + \log \pi_{22} - \log \pi_{12} - \log \pi_{21} = 0.$$

Now consider the log transformation of the expected counts for the ij^{th} cell given by m_{ij} . For which one has,

$$\log (m_{ij}) = \mu + \lambda_i^X + \lambda_j^Y + \lambda_{ij}^{XY}$$

for $i, j = 1, 2$ where $m_{ij} = n\pi_{ij}$. This equation is called the **saturated loglinear model** for the 2×2 table.

Since there are $1 + 2 + 2 + 4 = 9$ parameters in this model and only four observations (cell frequencies) it is necessary to define the following constraints on the model,

$$\sum_i \lambda_i^X = 0 \quad \sum_j \lambda_j^Y = 0 \quad \sum_i \lambda_{ij}^{XY} = \sum_j \lambda_{ij}^{XY} = 0.$$

The loglinear model expected cell counts can be written as,

X	Y=1	Y=2
1	$\exp(\mu + \lambda_1^X + \lambda_1^Y + \lambda_{11}^{XY})$	$\exp(\mu + \lambda_1^X - \lambda_1^Y - \lambda_{11}^{XY})$
2	$\exp(\mu - \lambda_1^X + \lambda_1^Y - \lambda_{11}^{XY})$	$\exp(\mu - \lambda_1^X - \lambda_1^Y + \lambda_{11}^{XY})$

Using the above loglinear model, the odds ratio can be written as

$$\psi = \frac{m_{11}m_{22}}{m_{12}m_{21}}$$

so that

$$\log \psi = \log m_{11} + \log m_{22} - \log m_{12} - \log m_{21} = 4\lambda_{11}^{XY}.$$

In which case, the hypothesis of independence of X and Y is equivalent to $H_0 : \lambda_{11}^{XY} = 0$. Thus, the loglinear model when the assumption of independence holds becomes,

$$\log(m_{ij}) = \mu + \lambda_i^X + \lambda_j^Y \quad i, j = 1, 2.$$

$R \times C$ Tables

Agresti considers the model given by,

$$\mu_{ij} = \mu\alpha_i\beta_j.$$

From which one has the *saturated model*

$$\log \mu_{ij} = \lambda + \lambda_i^X + \lambda_j^Y + \lambda_{ij}^{XY} \quad (1)$$

and the *independent model*,

$$\log \mu_{ij} = \lambda + \lambda_i^X + \lambda_j^Y \quad (2)$$

for $i = 1, 2, \dots, R, j = 1, 2, \dots, C$.

Three-Way Tables

Types of Independence

- The **saturated model** has loglinear form,

$$\log \mu_{ijk} = \lambda + \lambda_i^X + \lambda_j^Y + \lambda_k^Z + \lambda_{ij}^{XY} + \lambda_{ik}^{XZ} + \lambda_{jk}^{YZ} + \lambda_{ijk}^{XYZ}$$

- **Mutual Independence** X , Y , and Z are mutually independent when

$$\pi_{ijk} = \pi_{i++}\pi_{+j+}\pi_{++k}$$

for all i, j and k . Mutual independence implies that the expected frequencies $\{\mu_{ijk}\}$ have the loglinear form given by,

$$\log \mu_{ijk} = \lambda + \lambda_i^X + \lambda_j^Y + \lambda_k^Z,$$

in which case

$$\lambda_{ij}^{XY} = \lambda_{ik}^{XZ} = \lambda_{jk}^{YZ} = \lambda_{ijk}^{XYZ} = 0.$$

- **Joint Independence** Variable Y is jointly independent of X and Z when

$$\pi_{ijk} = \pi_{i+k}\pi_{+j+}$$

for all i, j and k . The loglinear form can be written as,

$$\log \mu_{ijk} = \lambda + \lambda_i^X + \lambda_j^Y + \lambda_k^Z + \lambda_{ik}^{XZ},$$

in which case

$$\lambda_{ij}^{XY} = \lambda_{jk}^{YZ} = \lambda_{ijk}^{XYZ} = 0.$$

Mutual independence implies joint independence of any one variable from the others.

- **Conditional Independence** X and Y are conditionally independent, given Z when

$$\pi_{ij|k} = \pi_{i+|k}\pi_{+j|k}$$

for all i, j and k . This holds for joint probabilities over the entire table, hence,

$$\pi_{ijk} = \pi_{i+k}\pi_{+jk}/\pi_{++k}.$$

The loglinear form can be written as,

$$\log \mu_{ijk} = \lambda + \lambda_i^X + \lambda_j^Y + \lambda_k^Z + \lambda_{ik}^{XZ} + \lambda_{jk}^{YZ},$$

in which case

$$\lambda_{ij}^{XY} = \lambda_{ijk}^{XYZ} = 0.$$

SAS

Code 5

```
title 'Three Way Model';
title2 'Death, Low Birth Weight and Smoking';
proc catmod data=new_bwgt;
  model death*smoking*lowbirthwgt=_response_
    / noparm pred=freq;
  loglin death|smoking|lowbirthwgt @ 2;
run;

title2 'Final Model';
proc catmod data=new_bwgt;
  model death*smoking*lowbirthwgt=_response_
    / noparm pred=freq;
  loglin death|lowbirthwgt death|smoking;
run;
```

Three Way Model

Death, Low Birth Weight and Smoking

The CATMOD Procedure

<i>Data Summary</i>			
<i>Response</i>	Death*Smoking*LowBirthWg	<i>Response Levels</i>	8
<i>Weight Variable</i>	None	<i>Populations</i>	1
<i>Data Set</i>	NEW_BWGT	<i>Total Frequency</i>	2892
<i>Frequency Missing</i>	197	<i>Observations</i>	2892

<i>Population Profiles</i>	
<i>Sample</i>	<i>Sample Size</i>
1	2892

<i>Response Profiles</i>			
<i>Response</i>	<i>Death</i>	<i>Smoking</i>	<i>LowBirthWgt</i>
1	Aff	Aff	Aff
2	Aff	Aff	No
3	Aff	No	Aff
4	Aff	No	No
5	No	Aff	Aff
6	No	Aff	No
7	No	No	Aff
8	No	No	No

<i>Maximum Likelihood Analysis</i>
Maximum likelihood computations converged.

<i>Maximum Likelihood Analysis of Variance</i>			
<i>Source</i>	<i>DF</i>	<i>Chi-Square</i>	<i>Pr > ChiSq</i>
<i>Death</i>	1	154.35	<.0001
<i>Smoking</i>	1	369.69	<.0001
<i>Death*Smoking</i>	1	3.46	0.0627
<i>LowBirthWgt</i>	1	152.09	<.0001
<i>Death*LowBirthWgt</i>	1	718.75	<.0001
<i>Smoking*LowBirthWgt</i>	1	0.06	0.8002
<i>Likelihood Ratio</i>	1	3.73	0.0535

<i>Maximum Likelihood Predicted Values for Response Functions</i>					
<i>Function Number</i>	<i>Observed</i>		<i>Predicted</i>		<i>Residual</i>
	<i>Function</i>	<i>Standard Error</i>	<i>Function</i>	<i>Standard Error</i>	
1	−2.78174	0.102035	−2.70818	0.091798	−0.07355
2	−3.34627	0.133598	−3.47632	0.123381	0.130054
3	−1.76836	0.064544	−1.79023	0.064295	0.021877
4	−2.58643	0.093122	−2.52479	0.08532	−0.06164
5	−3.43642	0.139553	−3.58015	0.127129	0.143728
6	−1.2164	0.05154	−1.19711	0.050321	−0.01929
7	−2.47224	0.088326	−2.41662	0.081669	−0.05562

<i>Maximum Likelihood Predicted Values for Frequencies</i>							
<i>Death</i>	<i>Smoking</i>	<i>LowBirthWgt</i>	<i>Observed</i>		<i>Predicted</i>		<i>Residual</i>
			<i>Frequency</i>	<i>Standard Error</i>	<i>Frequency</i>	<i>Standard Error</i>	
<i>Aff</i>	<i>Aff</i>	<i>Aff</i>	102	9.919803	109.2989	9.548808	−7.2989
<i>Aff</i>	<i>Aff</i>	<i>No</i>	58	7.539018	50.7011	5.985303	7.298899
<i>Aff</i>	<i>No</i>	<i>Aff</i>	281	15.92786	273.7011	15.29083	7.298899
<i>Aff</i>	<i>No</i>	<i>No</i>	124	10.89418	131.2989	10.55222	−7.2989
<i>No</i>	<i>Aff</i>	<i>Aff</i>	53	7.213092	45.7011	5.566915	7.298899
<i>No</i>	<i>Aff</i>	<i>No</i>	488	20.14086	495.2989	19.91188	−7.2989
<i>No</i>	<i>No</i>	<i>Aff</i>	139	11.50301	146.2989	11.1763	−7.2989
<i>No</i>	<i>No</i>	<i>No</i>	1647	26.62762	1639.701	26.38253	7.298899

Three Way Model

Final Model

The CATMOD Procedure

Data Summary			
Response	Death*Smoking*LowBirthWgt	Response Levels	8
Weight Variable	None	Populations	1
Data Set	NEW_BWGT	Total Frequency	2892
Frequency Missing	197	Observations	2892

Population Profiles	
Sample	Sample Size
1	2892

Response Profiles			
Response	Death	Smoking	LowBirthWgt
1	Aff	Aff	Aff
2	Aff	Aff	No
3	Aff	No	Aff
4	Aff	No	No
5	No	Aff	Aff
6	No	Aff	No
7	No	No	Aff
8	No	No	No

Maximum Likelihood Analysis
Maximum likelihood computations converged.

Maximum Likelihood Analysis of Variance			
Source	DF	Chi-Square	Pr > ChiSq
Death	1	167.50	<.0001
LowBirthWgt	1	201.07	<.0001
Death*LowBirthWgt	1	721.21	<.0001
Smoking	1	405.06	<.0001
Death*Smoking	1	6.34	0.0118
Likelihood Ratio	2	3.79	0.1502

All the above terms are significant, including the association between smoking and death!

<i>Maximum Likelihood Predicted Values for Response Functions</i>					
<i>Function Number</i>	<i>Observed</i>		<i>Predicted</i>		<i>Residual</i>
	<i>Function</i>	<i>Standard Error</i>	<i>Function</i>	<i>Standard Error</i>	
1	−2.78174	0.102035	−2.71524	0.08769	−0.0665
2	−3.34627	0.133598	−3.45927	0.102826	0.112998
3	−1.76836	0.064544	−1.78652	0.06252	0.018168
4	−2.58643	0.093122	−2.53055	0.082415	−0.05588
5	−3.43642	0.139553	−3.60304	0.089917	0.166622
6	−1.2164	0.05154	−1.19431	0.049075	−0.02208
7	−2.47224	0.088326	−2.40873	0.075344	−0.06351

<i>Maximum Likelihood Predicted Values for Frequencies</i>							
<i>Death</i>	<i>Smoking</i>	<i>LowBirthWgt</i>	<i>Observed</i>		<i>Predicted</i>		<i>Residual</i>
			<i>Frequency</i>	<i>Standard Error</i>	<i>Frequency</i>	<i>Standard Error</i>	
<i>Aff</i>	<i>Aff</i>	<i>Aff</i>	102	9.919803	108.4602	8.907781	−6.46018
<i>Aff</i>	<i>Aff</i>	<i>No</i>	58	7.539018	51.53982	5.057426	6.460177
<i>Aff</i>	<i>No</i>	<i>Aff</i>	281	15.92786	274.5398	14.94761	6.460177
<i>Aff</i>	<i>No</i>	<i>No</i>	124	10.89418	130.4602	9.976526	−6.46018
<i>No</i>	<i>Aff</i>	<i>Aff</i>	53	7.213092	44.63773	3.537735	8.362269
<i>No</i>	<i>Aff</i>	<i>No</i>	488	20.14086	496.3623	19.48686	−8.36227
<i>No</i>	<i>No</i>	<i>Aff</i>	139	11.50301	147.3623	10.41251	−8.36227
<i>No</i>	<i>No</i>	<i>No</i>	1647	26.62762	1638.638	26.05253	8.362269