### Solution to HW9

### Problem 6.16

(a) Denote Y = 1, 2, 3, 4 for 4 categories of the cholesterol level, x = 1/0 for treatment and control, mid-point  $y_0 = 1.7, 3.75, 4.5, 5.3$  for baseline cholesterol level. We consider the following cumulative logit model for  $\tau_j(x, y_0) = P(Y \le j)$ :

$$logit(\tau_i(x, y_0) = \alpha_i + \beta_1 x + \beta_2 y_0, j = 1, 2, 3.$$

The SAS program and part of the output:

```
data prob6_16;
  input 1d10 c1-c4 t1-t4;
  datalines;
1.7 18 8 0 0 21 4 2 0
3.75 16 30 13 2 17 25 6 0
4.5 0 14 28 7 11 35 36 6
5.3 0 2 15 22 1 5 14 12
data prob6_16; set prob6_16;
     array temp1 {4} c1-c4;
     array temp2 {4} t1-t4;
     do trt=0 to 1;
  do y=1 to 4;
          if trt=0 then count=temp1(y);
           else
             count=temp2(y);
          output;
        end;
     end;
     drop c1-c4 t1-t4;
run;
title "Problem 6.16(a)";
proc logistic;
  freq count;
  model y = trt ldl0 / aggregate scale=none;
```

Come Took for the December 2 Odda Assumption

Score Test for the Proportional Odds Assumption

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Chi-Square	DF	Pr > ChiSq
10.3917	4	0.0343

### Deviance and Pearson Goodness-of-Fit Statistics

Criterion	Value	DF	Value/DF	Pr > ChiSq
Deviance	37.0189	19	1.9484	0.0079
Pearson	41.4759	19	2.1829	0.0021

# Analysis of Maximum Likelihood Estimates

Parameter	DF	Estimate	Standard Error	Wald Chi-Square	Pr > ChiSq
Intercept 1	1	4.3071	0.5389	63.8825	<.0001
Intercept 2		6.5405	0.5927	121.7646	<.0001
Intercept 2 Intercept 3	1	8.7275	0.6493	180.6736	<.0001
trt	1	0.7767	0.2086	13.8669	0.0002
1d10	1	-1.6029	0.1368	137.3653	<.0001

### Odds Ratio Estimates

Effect	Point Estimate	95% Wald Confidence Limits		
trt 1d10	2.174	1.445	3.272	

Based on the output, we know that the treatment has a significant effect in reducing cholesterol level: the odds that patients receiving the treatment have better (lower) cholesterol level is 2.17 times the odds that patients receiving the control have better cholesterol level given that they had the same baseline cholesterol level (P-value=0.0002).

(b) Denote  $D_1, D_2, D_2$  three dummy variables for the first three categories for the baseline cholesterol level. Consider the following cumulative logit model for  $\tau_j(x, D's) = P(Y \leq j)$ :

$$logit(\tau_i(x, D's) = \alpha_i + \beta_1 x + \beta_2 D_1 + \beta_3 + D_2 \beta_4 D_3, \quad j = 1, 2, 3.$$

The SAS program and part of the output:

```
title "Problem 6.16(b)";
proc logistic;
  class ld10 / param=ref;
  freq count;
  model y = trt ld10 / aggregate scale=none;
run;
```

\*

Score Test for the Proportional Odds Assumption

Chi-Square	DF	Pr > ChiSq	
0.8827	8	0.9989	

Deviance and Pearson Goodness-of-Fit Statistics

Criterion	Value	DF	Value/DF	Pr > ChiSq
Deviance	14.4679	17	0.8511	0.6338
Pearson	11.6904	17	0.6877	0.8185

#### Analysis of Maximum Likelihood Estimates

Parameter		DF	Estimate	Standard Error	Wald Chi-Square	Pr > ChiSq
Intercept Intercept		1	-4.9700 -2.6791	0.3721 0.3158	178.4240 71.9755	<.0001 <.0001
Intercept		1	-0.2062	0.2516	0.6716	0.4125
trt 1d10	1.7	1 1	0.7924 5.6437	0.2097 0.4661	14.2765 146.6434	0.0002 <.0001
ld10 ld10	3.75 4.5	1 1	3.7689 1.9467	0.3603 0.3150	109.4417 38.2025	<.0001 <.0001

## Odds Ratio Estimates

Effect	Point Estimate	95% Wald Confidence Limits	}
trt	2.209	1.464 3.33	32

Based on the output, we know that the treatment has a significant effect in reducing cholesterol level: the odds that patients receiving the treatment have better cholesterol level is 2.21

times the odds that patients receiving the control have better cholesterol level given that they had the same baseline cholesterol level (P-value=0.0002).

(c) SAS program and output for the CMH test:

```
title "Problem 6.16(c)";
proc freq;
weight count;
tables ldl0*trt*y / cmh;
run;
```

Summary Statistics for trt by y
Controlling for 1d10

Cochran-Mantel-Haenszel Statistics (Based on Table Scores)

Statistic	Alternative Hypothesis	DF	Value	Prob
1	Nonzero Correlation	1	15.4001	<.0001
2	Row Mean Scores Differ	1	15.4001	<.0001
3	General Association	3	15.4562	0.0015

Using score (1,2,3,4) for ending cholesterol level, the CMH test for conditional independence between the treatment and ending cholesterol level given the baseline cholesterol level produces  $\chi^2 = 15.4001$ , df = 1 and P - value < 0.0001. Treating the ending cholesterol level as nominal categorical variable, the CMH test for conditional independence between the treatment and ending cholesterol level given the baseline cholesterol level produces  $\chi^2 = 15.4562$ , df = 3 and P - value < 0.0015. Both tests indicate that we should reject the conditional independence between the treatment and ending cholesterol level giving baseline cholesterol level.

### Problem 8.1

The McNemar test statistic is

$$\chi^2 = \frac{(16 - 37)^2}{16 + 37} = 8.32, \text{ P - value} = P(\chi_1^2 \ge 8.32) = 0.004.$$

Therefore, we reject the null hypothesis that the diabetes probabilities between MI cases and MI controls are the same. From the table, we know that MI cases have higher diabetes probability than MI controls.

### Problem 8.2

(a) Here we can use the McNemar test to test this null hypothesis since we have matched data.

The McNemar test statistic is

$$\chi^2 = \frac{(125-2)^2}{125+2} = 119$$
,  $P - \text{value} = P(\chi_1^2 \ge 119) = 0$ .

Therefore, we reject the null hypothesis that the population proportions answering "yes" were the same for heaven and hell (almost at any level).

(b) The difference of sample proportions answering "yes" for heaven and hell is:

$$p_1 - p_2 = (833 + 125)/1120 - (833 + 2)/1120 = 0.855 - 0.746 = 0.11.$$

The estimated variance (and SE) of the above difference:

$$\widehat{\text{var}}(p_1 - p_2) = \frac{(125 + 2) - (125 - 2)^2 / 1120)}{1120^2} = 9 \times 10^{-5}, \ \widehat{\text{SE}}(p_1 - p_2) = 0.009.$$

So a 90% CI for the proportion difference is

$$0.11 \pm 1.645 \times 0.009 = [0.095, 0.125].$$

## Problem 8.3

(a) Denote  $Y_{ij}$  the indicator variable for "yes" for either heaven or hell, x the indicator for heaven (1 for heaven, 0 for hell). The marginal model that will produce marginal odds-ratio is:

$$logit{P(Y_{ij} = 1|x)} = \alpha + x\beta.$$

The correlation among the repeated observations within the same subject can be taken into account using GEE. The SAS program and part of the output is:

```
data prob8_3;
  input heaven hell count;
  datalines;
  1 1 833
  1 0 125
  0 1 2
  0 0 160
;

title "recover individual data";
data newdata; set prob8_3;
  retain id;
  if _n_=1 then id=0;

  do i=1 to count;
  id = id+1;
   do x=0 to 1;
    if x=0 then
       y=heaven;
    output;
  end;
end;
end;
```

```
run;
title "Problem 8.3(a)";
proc genmod data=newdata descending;
  class id;
  model y = x / dist=bin link=logit;
  repeated subject=id / type=un;
run:
```

\*

### Analysis Of GEE Parameter Estimates Empirical Standard Error Estimates

Parameter	Estimate	Standard Error	95% Con: Lim	Z 1	Pr >  Z
Intercept x		0.0686 0.0621			

The odds-ratio estimate is  $e^{0.7023} = 2.02$ , indicating that the population odds of believing heaven is 2.02 times the population odds of believing hell. The P-value also indicates that these two population odds are not the same.

(b) The conditional model that will produce subject-specific odds-ratio is

$$logit{P(Y_{ij} = 1 | \alpha_i, x)} = \alpha_i + x\beta.$$

We use conditional logistic regression to fit the above model. The SAS program and part of the output is:

```
title "Problem 8.3(b)";
proc logistic data=newdata descending;
  model y = x;
  strata id;
run:
```

\*

Analysis of Conditional Maximum Likelihood Estimates

Parameter	DF	Estimate	Standard Error	Wald Chi-Square	Pr > ChiSq
x	1	4.1352	0.7127	33.6606	<.0001
		Odds I	Ratio Estima	tes	
	Effec	Poin t Estima		95% Wald idence Limits	
	х	62.50	00 15.4	59 252.677	

The subject-specific odds-ratio estimate is  $e^{4.1352} = 62.5$ , indicating for each subject, the odds of believing heaven is 62.5 times the odds of believing hell. The P-value also indicates that these two subject-specific odds are not the same.