Homework #8

Stat 545 (GLM & Categorical Data Analysis)

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1. Problem 8.1

(a) The model is fit by R function *clogit*.

```
> library('survival')
> vote <- c(rep(c(1,1),175),rep(c(1,0),16),rep(c(0,1),54),rep(c(0,0),188))
> year <- rep(c(0,1),433)</pre>
> id <- ceiling(1:866/2)</pre>
> fit <- clogit(vote~year + strata(id))</pre>
coxph(formula = Surv(rep(1, 866L), vote) ~ year + strata(id),
    method = "exact")
  n= 866, number of events= 420
       coef exp(coef) se(coef)
                                   z Pr(>|z|)
year 1.2164 3.3750 0.2846 4.273 1.92e-05 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
     exp(coef) exp(-coef) lower .95 upper .95
                   0.2963
         3.375
                              1.932
Rsquare= 0.025 (max possible= 0.106 )
Likelihood ratio test= 21.78 on 1 df,
                                         p=3.051e-06
                   = 18.26 on 1 df,
                                        p=1.924e-05
Score (logrank) test = 20.63 on 1 df, p=5.576e-06
```

Hence $\hat{\beta} = 1.216$ and $\exp(\hat{\beta}) = 3.375 = 54/16$. β describes the conditional association between the 2004 vote and 2008 vote for each fixed individual. The exponential of its MLE is equal to the ratio of the off-diagonal counts in the table. $\exp(\beta)$ may also be called the true odds ratio for each individual.

Since R reports a p-value 1.9×10^{-5} for $\hat{\beta}$, there is strong evidence that β is actually bigger than 0. Voter preference for Democrats has increased.

- (b) α_i represents a fixed subject-specific effect for each individual. A larger α_i means the individual has a stronger tendency to vote for Democrats in both 2004 and 2008.
- (c) The MLE of β for population averaged effect model is simply the log odds ratio.

$$\hat{\beta}_2 = \log \frac{229 \times 242}{191 \times 204} = 0.352$$

The MLE of β for subject-specific model is given above, $\hat{\beta}_1 = 1.216$.

They are not same and $\hat{\beta}_2 < \hat{\beta}_1$. This is expected. As is illustrated in Figure 13.1 of the textbook, when two individuals have very different α_i , their probability curves $P(Y_i = 1)$ v.s. x_i (in this problem, $x_i \in [0, 1]$) are spaced far apart. The marginal model tries to fit a curve that is averaged over all the individuals' curves and thus it has a shallower slope. By the approximation formula of Zeger et al. (1988),

$$\hat{\beta}_1 \approx \hat{\beta}_2 (1 + 0.346\sigma^2)^{-1/2}$$
 (1)

where σ^2 is the variance of α_i if a random effect model is assumed. In our problem, σ^2 is big because most people didn't change their side, which implies many people have a very large α_i while many others have a very small α_i . Therefore, by (1), $\hat{\beta}_1$ should be much larger.

(d) The McNemar's test is done in R with no continuity correction.

```
> marg.table <- matrix(c(175,16,54,188),2,2,byrow=T)
> mcnemar.test(marg.table,correct=F)

McNemar's Chi-squared test

data: marg.table
McNemar's chi-squared = 20.629, df = 1, p-value = 5.576e-06
```

The p-value is similar to that of conditional logistic regression. This is expected. McNemar test is actually the score test. Asymptotically, it is equivalent to Wald test and likelihood ratio test. Therefore, the p-value of the conditional logistic regression, which is usually computed by Wald test, is often very close to the p-value of McNemar test, especially when the sample size is large.

(e) No. Neither of them would change. This is because both McNemar's test and the conditional likelihood only depend on the off-diagonal counts. For McNemar's test, recall that the p-value is evaluated under the null distribution and the diagonal counts only influence the inferences under the alternative, e.g., how much heterogeneity there exists. For conditional logistic regression, observe that the MLE of β given the whole table is the same as the MLE given only the off-diagonal cells. This is because, for any $\beta \in \mathbb{R}$, the likelihood of the diagonal cells can always take any value in \mathbb{R} by choosing the appropriate values of α_i .

```
> vote <- c(rep(c(1,1),100),rep(c(1,0),16),rep(c(0,1),54),rep(c(0,0),263))
> fit <- clogit(vote~year + strata(id))
> summary(fit)
Call:
coxph(formula = Surv(rep(1, 866L), vote) ~ year + strata(id),
   method = "exact")
 n= 866, number of events= 270
      coef exp(coef) se(coef)
                                 z Pr(>|z|)
year 1.2164 3.3750 0.2846 4.273 1.92e-05 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
    exp(coef) exp(-coef) lower .95 upper .95
        3.375
                  0.2963
                             1.932
Rsquare= 0.025 (max possible= 0.106 )
Likelihood ratio test= 21.78 on 1 df,
                                       p=3.051e-06
                 = 18.26 on 1 df,
Wald test
                                        p=1.924e-05
Score (logrank) test = 20.63 on 1 df, p=5.576e-06
> marg.table <- matrix(c(100,16,54,263),2,2,byrow=T)</pre>
> mcnemar.test(marg.table,correct=F)
McNemar's Chi-squared test
data: marg.table
McNemar's chi-squared = 20.629, df = 1, p-value = 5.576e-06
```

2. Problem 8.2

(a) The 'response' column in the data provided is treated as Y. Logistic regression is fitted in R. The fitted model is

```
logit(P(Y_t = 1)) = -0.125 + 0.149I(t = 1) + 0.0520I(t = 2) + 0.00358X
                                                                                       (2)
          > dat <- as.data.frame(read.table('attitude.csv',header=T,sep=','))</pre>
          > fit <- glm(response~gender+dummy1+dummy2,data=dat,family='binomial')
          > summary(fit)
          Call:
          glm(formula = response ~ gender + dummy1 + dummy2, family = "binomial",
             data = dat)
          Deviance Residuals:
           Min
                  1Q Median
                                 30
          -1.189 -1.148 -1.125 1.207
                                      1.231
          Coefficients:
                     Estimate Std. Error z value Pr(>|z|)
          0.003582 0.054138 0.066 0.9472
          gender
                     0.149347 0.065825 2.269
                                                0.0233 *
          dummv1
                     0.052018 0.065843 0.790 0.4295
          dummv2
          Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' '1
          (Dispersion parameter for binomial family taken to be 1)
             Null deviance: 7689.5 on 5549 degrees of freedom
          Residual deviance: 7684.2 on 5546 degrees of freedom
         AIC: 7692.2
          Number of Fisher Scoring iterations: 3
```

(b) I fit the GEE logistic regression in R by using 'gee' library. The code and output are attached. In the R output, 'naive SE' means the model-based standard error. 'robust SE' means the empirical (sandwich) standard error. The results are summarized in the following tables.

Exchangeable correlation structure:

	row.names	Estimate	Model-based SE	Model-based Pv	Empirical SE	Empirical Pv
1	alpha	0.4687000	0.016920	0.000e+00	0.016850	0.000e+00
2	dummy1	0.0373000	0.007023	1.091e-07	0.007420	4.991e-07
3	dummy2	0.0129700	0.007023	6.471e-02	0.006745	5.442e-02
4	gender	0.0008939	0.021940	9.675e-01	0.021920	9.675e-01

	row.names	Q1	Q2	Q3
1	Q1	1.0000	0.8173	0.8173
2	Q2	0.8173	1.0000	0.8173
3	Q3	0.8173	0.8173	1.0000

Unstructured correlation:

	row.names	Estimate	Model-based SE	Model-based Pv	Empirical SE	Empirical Pv
1	alpha	0.468400	0.016920	0.000e+00	0.016850	0.000e+00
2	dummy1	0.037300	0.007427	5.119e-07	0.007420	4.991e-07
3	dummy2	0.012970	0.006763	5.508e-02	0.006745	5.442e-02
4	gender	0.001375	0.021930	9.500e-01	0.021920	9.500e-01

	row.names	Q1	Q2	Q3
1	Q1	1.0000	0.8257	0.7957
2	Q2	0.8257	1.0000	0.8306
3	Q3	0.7957	0.8306	1.0000

In either assumption of correlation structure, we have observed a very high estimated correlation between the three questions (around 0.8). This high correlation is indeed expected from Table 11.13. Due to the high correlation, the estimates and standard errors differ significantly from part (a) where we treat the three answers of the same individual as independent. Notice that by taking into consideration the correlation between 3 questions, the evidence for β_1 become much stronger, which implies that the attitudes toward different questions do differ.

R code:

Exchangeable:

```
> library(gee)
> exc <- gee(response~gender+dummy1+dummy2, id=case, data=dat,corstr='exchangeable')</pre>
Beginning Cgee S-function, @(#) geeformula.q 4.13 98/01/27
running glm to get initial regression estimate
(Intercept)
               gender
                            dummy1
0.4686880977 0.0008939434 0.0372972973 0.0129729730
> summary(exc)
GEE: GENERALIZED LINEAR MODELS FOR DEPENDENT DATA
gee S-function, version 4.13 modified 98/01/27 (1998)
Model:
Link:
                            Identity
Variance to Mean Relation: Gaussian
Correlation Structure:
                           Exchangeable
gee(formula = response ~ gender + dummy1 + dummy2, id = case,
    data = dat, corstr = "exchangeable")
Summary of Residuals:
      Min 1Q
                         Median
-0.5068793 -0.4825550 -0.4686881 0.5174450 0.5313119
Coefficients:
                Estimate Naive S.E.
                                         Naive z Robust S.E.
(Intercept) 0.4686880977 0.016917979 27.70355085 0.016848315 27.81809838
          0.0008939434 0.021938148 0.04074835 0.021921409 0.04077947
           0.0372972973 0.007022767 5.31091176 0.007419920 5.02664385 0.0129729730 0.007022767 1.84727366 0.006744609 1.92345826
dummy2
Estimated Scale Parameter: 0.2497433
Number of Iterations: 1
Working Correlation
         [,1] [,2]
                              [,3]
[1,] 1.0000000 0.8173312 0.8173312
[2,] 0.8173312 1.0000000 0.8173312
[3,] 0.8173312 0.8173312 1.0000000
```

Unstructured:

```
> unstr <- gee(response~gender+dummy1+dummy2, id=case, data=dat,corstr='unstructured')
Beginning Cgee S-function, @(#) geeformula.q 4.13 98/01/27
running glm to get initial regression estimate
(Intercept) gender dummy1 dummy2
0.4686880977 0.0008939434 0.0372972973 0.0129729730
> summary(unstr)
GEE: GENERALIZED LINEAR MODELS FOR DEPENDENT DATA
gee S-function, version 4.13 modified 98/01/27 (1998)
Model:
Link:
                          Identity
 Variance to Mean Relation: Gaussian
Correlation Structure: Unstructured
gee(formula = response ~ gender + dummy1 + dummy2, id = case,
   data = dat, corstr = "unstructured")
Summary of Residuals:
                                   3Q
                      Median
     Min
             10
                                               Max
-0.5070910 -0.4827666 -0.4684182 0.5172334 0.5315818
Coefficients:
              Estimate Naive S.E.
                                   Naive z Robust S.E. Robust z
(Intercept) 0.468418173 0.016915490 27.69167072 0.016853850 27.7929485
         0.001375486 0.021932036 0.06271586 0.021916157 0.0627613
gender
dummy1
           0.037297297 0.007427091 5.02179084 0.007419920 5.0266439
dummy2
           0.012972973 0.006762812 1.91828085 0.006744609 1.9234583
Estimated Scale Parameter: 0.2497433
Number of Iterations: 2
Working Correlation
        [,1] [,2]
                          [,3]
[1,] 1.0000000 0.8256973 0.7956920
[2,] 0.8256973 1.0000000 0.8306043
[3,] 0.7956920 0.8306043 1.0000000
```