

```

> rm(list = ls())
> setwd("E:/Data of R")
>
> #Question 2
> library(VGAM)
> data2=read.csv("GSS.csv",header = T)
>
> names(data2)
[1] "gender"      "race"        "democrat"    "republican"  "independent"
>
> data2
  gender race democrat republican independent
1  male white     132         176         127
2  male black      42           6          12
3 female white    172        129         130
4 female black     56           4          15
>
> #(a)
>
> model21=vglm(cbind(democrat,republican,independent)~gender+race,data =
  data2,
+           family = multinomial)
> model21

```

Call:

```
vglm(formula = cbind(democrat, republican, independent) ~ gender +
  race, family = multinomial, data = data2)
```

Coefficients:

```

(Intercept):1 (Intercept):2 gendermale:1 gendermale:2 racewhite:1
  1.3882465    -1.1771027    -0.2201865     0.3525732    -1.1182884
racewhite:2
  1.1598459

```

Degrees of Freedom: 8 Total; 2 Residual

Residual deviance: 0.1982117

Log-likelihood: -20.17843

This is a multinomial logit model with 3 levels

```
> summary(model21)
```

Call:

```
vglm(formula = cbind(democrat, republican, independent) ~ gender +
  race, family = multinomial, data = data2)
```

Pearson residuals:

```
log(mu[,1]/mu[,3]) log(mu[,2]/mu[,3])
```

|   |          |          |
|---|----------|----------|
| 1 | -0.07696 | -0.05743 |
| 2 | 0.19896  | 0.22498  |
| 3 | 0.07201  | 0.05961  |
| 4 | -0.18480 | -0.23505 |

Coefficients:

|               | Estimate | Std. Error | z value | Pr(> z ) |     |
|---------------|----------|------------|---------|----------|-----|
| (Intercept):1 | 1.3882   | 0.2296     | 6.045   | 1.49e-09 | *** |
| (Intercept):2 | -1.1771  | 0.3807     | -3.092  | 0.00199  | **  |
| gendermale:1  | -0.2202  | 0.1583     | -1.391  | 0.16412  |     |
| gendermale:2  | 0.3526   | 0.1651     | 2.136   | 0.03271  | *   |
| racewhite:1   | -1.1183  | 0.2335     | -4.789  | 1.68e-06 | *** |
| racewhite:2   | 1.1598   | 0.3801     | 3.051   | 0.00228  | **  |

---

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Number of linear predictors: 2

Names of linear predictors: log(mu[,1]/mu[,3]), log(mu[,2]/mu[,3])

Residual deviance: 0.1982 on 2 degrees of freedom

Log-likelihood: -20.1784 on 2 degrees of freedom

Number of iterations: 3

No Hauck-Donner effect found in any of the estimates

Reference group is level 3 of the response

>

> #(b)

> deviance(model21)

[1] 0.1982117

> qchisq(0.05,4\*2-3\*2,lower.tail = FALSE)

[1] 5.991465

>

> #0.1982117<5.991465

> #We can not reject the null/reduced model in favor of the saturated model,

> #indicating the model fit with main effects for all the predictors provides

> #a reasonable fit.

>

> #(c)

> coefficients(model21)[3]

gendermale:1

-0.2201865

> 1/exp(coefficients(model21)[3])

gendermale:1

```

1.246309
> #The estimated coefficient for the Gender Male dummy in the Democrat vs.
  Independent
> #is -0.2202. This means that men are 1.246309 times less likely choose D
  emocrat
> #over Independent.
>
> coefficients(model21)[4]
gendermale:2
  0.3525732
> exp(coefficients(model21)[4])
gendermale:2
  1.422724
> #The estimated coefficient for the Gender Male dummy in the Republican v
  s. Independent
> #is 0.3525732. This means that men are 1.422724 times more likely choos
  e Republican
> #over Independent.
>
> #Gender effect is not significant overall. Gender has a statistically si
  gnificant
> #effect when comparing Republican over Independent.
>
> #(d)
>
> #Prob. being Independent for black females
> pi3=1/(1+exp(1.3882465)+exp(-1.1771027))
> pi3
[1] 0.1881118
>
> #Prob. being Democrat for black females
> pi1=exp(1.3882465)*pi3
> pi1
[1] 0.7539177
>
> #(e)
> #Intercept1 is >0, which means that pi1/pi3>1, pi1>pi3
> #Intercept2 is <0, which means that pi2/pi3<1, pi2<pi3
> #So,pi1>pi3>pi2
> #PiD_hat > PiI_hat > PiR_hat
>
> #(f)
> coefficients(model21)
(Intercept):1 (Intercept):2 gendermale:1 gendermale:2 racewhite:1
  1.3882465 -1.1771027 -0.2201865 0.3525732 -1.1182884
racewhite:2
  1.1598459
>

```

```

> c1=c(coefficients(model21)[1],coefficients(model21)[3],coefficients(model21)[5])
> c1
(Intercept):1  gendermale:1  racewhite:1
      1.3882465    -0.2201865    -1.1182884
> #log(pi1/pi3)=1.3882465-0.2201865*male-1.1182884*white
>
> c2=c(coefficients(model21)[2],coefficients(model21)[4],coefficients(model21)[6])
> c2
(Intercept):2  gendermale:2  racewhite:2
      -1.1771027    0.3525732    1.1598459
> #log(pi2/pi3)=-1.1771027+0.3525732*male+1.159845*white
>
> #log(pi1/pi3)-log(pi2/pi3)=log(pi1/pi2)=c1-c2
> c1-c2
(Intercept):1  gendermale:1  racewhite:1
      2.5653492    -0.5727597    -2.2781343
> #log=(PiD/PiR)=log(pi1/pi2)=2.5653492-0.5727597*male-2.2781343*white
>
> #(g)
> model22=vglm(cbind(democrat,independent,repUBLICan)~gender+race,data =
  data2,
+           family = multinomial)
> summary(model22)

```

Call:

```
vglm(formula = cbind(democrat, independent, republican) ~ gender +
      race, family = multinomial, data = data2)
```

Pearson residuals:

|   | $\log(\mu_{[,1]}/\mu_{[,3]})$ | $\log(\mu_{[,2]}/\mu_{[,3]})$ |
|---|-------------------------------|-------------------------------|
| 1 | -0.03865                      | 0.08791                       |
| 2 | 0.03288                       | -0.29853                      |
| 3 | 0.03077                       | -0.08827                      |
| 4 | -0.01045                      | 0.29881                       |

Coefficients:

|               | Estimate | Std. Error | z value | Pr(> z ) |     |
|---------------|----------|------------|---------|----------|-----|
| (Intercept):1 | 2.5653   | 0.3437     | 7.465   | 8.35e-14 | *** |
| (Intercept):2 | 1.1771   | 0.3807     | 3.092   | 0.001986 | **  |
| gendermale:1  | -0.5728  | 0.1575     | -3.636  | 0.000277 | *** |
| gendermale:2  | -0.3526  | 0.1651     | -2.136  | 0.032707 | *   |
| racewhite:1   | -2.2781  | 0.3428     | -6.646  | 3.02e-11 | *** |
| racewhite:2   | -1.1598  | 0.3801     | -3.051  | 0.002279 | **  |

---

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Number of linear predictors: 2

Names of linear predictors:  $\log(\mu[,1]/\mu[,3])$ ,  $\log(\mu[,2]/\mu[,3])$

Residual deviance: 0.1982 on 2 degrees of freedom

Log-likelihood: -20.1784 on 2 degrees of freedom

Number of iterations: 3

No Hauck-Donner effect found in any of the estimates

Reference group is level 3 of the response

>

> coefficients(model22)

```
(Intercept):1 (Intercept):2 gendermale:1 gendermale:2 racewhite:1
  2.5653492    1.1771027   -0.5727597   -0.3525732   -2.2781343
```

```
racewhite:2
```

```
  -1.1598459
```

> #log=(PiD/PiR)=log(pi1/pi2)=2.5653492-0.5727597\*male-2.2781343\*white

> #It is the same as the result in part (f)

>

> #(h)

> # Democrat vs. Independence

> d1=cbind(rep(1,259),rep(1,259),c(rep(1,132),rep(0,127)))

> d2=cbind(rep(1,54),rep(0,54),c(rep(1,42),rep(0,12)))

> d3=cbind(rep(0,302),rep(1,302),c(rep(1,172),rep(0,130)))

> d4=cbind(rep(0,71),rep(0,71),c(rep(1,56),rep(0,15)))

> data21=rbind(d1,d2,d3,d4)

> colnames(data21)=c("gender","race","democrat")

> data21=data.frame(data21)

> logit1=glm(democrat~gender+race,data=data21,family=binomial(link = logit))

>

> # Republican vs. Independence

> d1=cbind(rep(1,303),rep(1,303),c(rep(1,176),rep(0,127)))

> d2=cbind(rep(1,18),rep(0,18),c(rep(1,6),rep(0,12)))

> d3=cbind(rep(0,259),rep(1,259),c(rep(1,129),rep(0,130)))

> d4=cbind(rep(0,19),rep(0,19),c(rep(1,4),rep(0,15)))

> data22=rbind(d1,d2,d3,d4)

> colnames(data22)=c("gender","race","republican")

> data22=data.frame(data22)

> logit2=glm(republican~gender+race,data=data22,family=binomial(link = logit))

>

> summary(logit1)

Call:

glm(formula = democrat ~ gender + race, family = binomial(link = logit),

```
data = data21)
```

Deviance Residuals:

| Min     | 1Q      | Median | 3Q     | Max    |
|---------|---------|--------|--------|--------|
| -1.7943 | -1.1992 | 0.7357 | 1.0655 | 1.1558 |

Coefficients:

|             | Estimate | Std. Error | z value | Pr(> z )     |
|-------------|----------|------------|---------|--------------|
| (Intercept) | 1.3866   | 0.2296     | 6.040   | 1.54e-09 *** |
| gender      | -0.2181  | 0.1585     | -1.376  | 0.169        |
| race        | -1.1175  | 0.2335     | -4.785  | 1.71e-06 *** |

---

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 930.60 on 685 degrees of freedom

Residual deviance: 902.32 on 683 degrees of freedom

AIC: 908.32

Number of Fisher Scoring iterations: 4

```
> summary(logit2)
```

Call:

```
glm(formula = republican ~ gender + race, family = binomial(link = logit),
```

```
data = data22)
```

Deviance Residuals:

| Min    | 1Q     | Median | 3Q    | Max   |
|--------|--------|--------|-------|-------|
| -1.322 | -1.171 | 1.040  | 1.040 | 1.697 |

Coefficients:

|             | Estimate | Std. Error | z value | Pr(> z )   |
|-------------|----------|------------|---------|------------|
| (Intercept) | -1.1698  | 0.3814     | -3.067  | 0.00216 ** |
| gender      | 0.3486   | 0.1660     | 2.099   | 0.03580 *  |
| race        | 1.1543   | 0.3809     | 3.031   | 0.00244 ** |

---

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 828.79 on 598 degrees of freedom

Residual deviance: 813.75 on 596 degrees of freedom

AIC: 819.75

Number of Fisher Scoring iterations: 4

```
> summary(model21)
```

Call:

```
vglm(formula = cbind(democrat, republican, independent) ~ gender +  
      race, family = multinomial, data = data2)
```

Pearson residuals:

|   | $\log(\mu[,1]/\mu[,3])$ | $\log(\mu[,2]/\mu[,3])$ |
|---|-------------------------|-------------------------|
| 1 | -0.07696                | -0.05743                |
| 2 | 0.19896                 | 0.22498                 |
| 3 | 0.07201                 | 0.05961                 |
| 4 | -0.18480                | -0.23505                |

Coefficients:

|               | Estimate | Std. Error | z value | Pr(> z )     |
|---------------|----------|------------|---------|--------------|
| (Intercept):1 | 1.3882   | 0.2296     | 6.045   | 1.49e-09 *** |
| (Intercept):2 | -1.1771  | 0.3807     | -3.092  | 0.00199 **   |
| gendermale:1  | -0.2202  | 0.1583     | -1.391  | 0.16412      |
| gendermale:2  | 0.3526   | 0.1651     | 2.136   | 0.03271 *    |
| racewhite:1   | -1.1183  | 0.2335     | -4.789  | 1.68e-06 *** |
| racewhite:2   | 1.1598   | 0.3801     | 3.051   | 0.00228 **   |

---

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Number of linear predictors: 2

Names of linear predictors:  $\log(\mu[,1]/\mu[,3])$ ,  $\log(\mu[,2]/\mu[,3])$

Residual deviance: 0.1982 on 2 degrees of freedom

Log-likelihood: -20.1784 on 2 degrees of freedom

Number of iterations: 3

No Hauck-Donner effect found in any of the estimates

Reference group is level 3 of the response

>

> #(i)

> #The coefficients and their statistical significance in two separate logistic

> #models are the same as the corresponding parts of the baseline category logit

> #model. That is because the submodel of the baseline category logit is exactly

> #the logistic model. Taking first logistic model as a example, the sumation

> #of  $y=0$  is exactly the number of people who choose independent

```

>
> #Question 3
> library(gee)
> data3=read.csv("attitudes.csv", header = T)
> names(data3)
[1] "gender" "response" "question" "case"
> dim(data3)
[1] 5550 4
>
> #(a)
> model31=gee(response~gender+as.factor(question),id=case,family=binomial,
+           constr="unstructured", scale.fix=T,data = data3)
Beginning Cgee S-function, @(#) geeformula.q 4.13 98/01/27
running glm to get initial regression estimate
              (Intercept)              gender as.factor(question)2
              0.023939537              0.003582051              -0.097329124
as.factor(question)3
              -0.149347113
> summary(model31)

```

GEE: GENERALIZED LINEAR MODELS FOR DEPENDENT DATA  
gee S-function, version 4.13 modified 98/01/27 (1998)

Model:  
Link: Logit  
Variance to Mean Relation: Binomial  
Correlation Structure: Unstructured

Call:  
gee(formula = response ~ gender + as.factor(question), id = case,  
data = data3, family = binomial, constr = "unstructured",  
scale.fix = T)

Summary of Residuals:

| Min        | 1Q         | Median     | 3Q        | Max       |
|------------|------------|------------|-----------|-----------|
| -0.5070710 | -0.4827461 | -0.4684470 | 0.5172539 | 0.5315530 |

Coefficients:

|                      | Estimate     | Naive S.E. | Naive z     | Robust S.E. | Robust z   |
|----------------------|--------------|------------|-------------|-------------|------------|
| (Intercept)          | 0.022967184  | 0.06773175 | 0.33909035  | 0.06778176  | 0.3388402  |
| gender               | 0.005318694  | 0.08785139 | 0.06054194  | 0.08782143  | 0.0605626  |
| as.factor(question)2 | -0.097328985 | 0.02753097 | -3.53525447 | 0.02753163  | -3.5351691 |
| as.factor(question)3 | -0.149346974 | 0.02973943 | -5.02184998 | 0.02973863  | -5.0219851 |



Estimated Scale Parameter: 1

Number of Iterations: 2

working correlation

|      | [,1]      | [,2]      | [,3]      |
|------|-----------|-----------|-----------|
| [1,] | 1.0000000 | 0.8248498 | 0.7958825 |
| [2,] | 0.8248498 | 1.0000000 | 0.8312594 |
| [3,] | 0.7958825 | 0.8312594 | 1.0000000 |

>

> #(b)

> model31\$working.correlation

|      | [,1]      | [,2]      | [,3]      |
|------|-----------|-----------|-----------|
| [1,] | 1.0000000 | 0.8248498 | 0.7958825 |
| [2,] | 0.8248498 | 1.0000000 | 0.8312594 |
| [3,] | 0.7958825 | 0.8312594 | 1.0000000 |

> #The working correlation indicates that in the same case, the correlation between 1 and 2 is

> #0.8248498, the correlation between 1 and 3 is 0.7958825 and the correlation

> #between 2 and 3 is 0.8312594

>

> #These estimation is large (closing to 1), which indicates that accounting

> #for clustering is necessary

>

> #(c)

> model32=gee(response~gender+as.factor(question), id=case, family=binomial,

+ corstr="exchangeable", scale.fix=T, data = data3)

Beginning Cgee S-function, @(#) geeformula.q 4.13 98/01/27

running glm to get initial regression estimate

| (Intercept)          | gender      | as.factor(question)2 |
|----------------------|-------------|----------------------|
| 0.023939537          | 0.003582051 | -0.097329124         |
| as.factor(question)3 |             |                      |
| -0.149347113         |             |                      |

>

> model31\$working.correlation

|      | [,1]      | [,2]      | [,3]      |
|------|-----------|-----------|-----------|
| [1,] | 1.0000000 | 0.8248498 | 0.7958825 |
| [2,] | 0.8248498 | 1.0000000 | 0.8312594 |
| [3,] | 0.7958825 | 0.8312594 | 1.0000000 |

> model32\$working.correlation

|      | [,1]      | [,2]      | [,3]      |
|------|-----------|-----------|-----------|
| [1,] | 1.0000000 | 0.8173308 | 0.8173308 |
| [2,] | 0.8173308 | 1.0000000 | 0.8173308 |
| [3,] | 0.8173308 | 0.8173308 | 1.0000000 |

> #The working correlation matrix indicates that the correlation between observations

```

> #within a subject is estimated to be 0.8173308
> #Because we specified an exchangeable correlation structure, this correlation
> #is the same for all pairs in a group
>
> #0.8173308 is between 0.8248498 and 0.7958825 and it is also closed to either
> #of them, which indicates that is reasonable to use exchangeable correlation
>
> #(d)
> summary(model32)

```

GEE: GENERALIZED LINEAR MODELS FOR DEPENDENT DATA  
 gee S-function, version 4.13 modified 98/01/27 (1998)

Model:

```

Link:                      Logit
Variance to Mean Relation: Binomial
Correlation Structure:     Exchangeable

```

Call:

```

gee(formula = response ~ gender + as.factor(question), id = case,
    data = data3, family = binomial, corstr = "exchangeable",
    scale.fix = T)

```

Summary of Residuals:

|  | Min        | 1Q         | Median     | 3Q        | Max       |
|--|------------|------------|------------|-----------|-----------|
|  | -0.5068644 | -0.4825396 | -0.4687095 | 0.5174604 | 0.5312905 |

Coefficients:

|                      | Estimate     | Naïve S.E. | Naïve z     | Robust S.E. |
|----------------------|--------------|------------|-------------|-------------|
| (Intercept)          | 0.024021377  | 0.06774107 | 0.35460579  | 0.06779334  |
| gender               | 0.003437873  | 0.08787462 | 0.03912248  | 0.08784072  |
| as.factor(question)2 | -0.097329120 | 0.02811568 | -3.46173752 | 0.02753161  |
| as.factor(question)3 | -0.149347107 | 0.02813360 | -5.30849707 | 0.02973865  |

  

|                      | Robust z    |
|----------------------|-------------|
| (Intercept)          | 0.35433237  |
| gender               | 0.03913758  |
| as.factor(question)2 | -3.53517666 |
| as.factor(question)3 | -5.02198729 |

Estimated Scale Parameter: 1

Number of Iterations: 2

Working Correlation

|      | [,1]      | [,2]      | [,3]      |
|------|-----------|-----------|-----------|
| [1,] | 1.0000000 | 0.8173308 | 0.8173308 |

```

[2,] 0.8173308 1.0000000 0.8173308
[3,] 0.8173308 0.8173308 1.0000000
> exp(coefficients(model32)[2])
gender
1.003444
> #The estimated coefficient for the gender is 0.003437873
> #This means that females are 1.003444 times more likely to support legal
  ized abortion
>
> qnorm(0.975)
[1] 1.959964
>
> #Naive z 0.03912248 < 1.959964
> #Can not reject H0
>
> #(e)
> lb=0.003437873-qnorm(0.975)*0.08787462
> ub=0.003437873+qnorm(0.975)*0.08787462
>
> #the 95% CI for      is
> c(lb,ub)
[1] -0.1687932  0.1756690
>
> #the 95% CI for      is
> c(exp(lb),exp(ub))
[1] 0.8446836 1.1920434
>
> #(f)
> #response=0.024021377+0.003437873*female-0.097329120*Q2-0.149347107*Q3
>
> #estimated odds of support for legalized abortion in scenario 2 for a ma
  le
> odds=0.024021377-0.097329120
> odds
[1] -0.07330774
>
> #(g)
> model33=gee(response~gender+question,id=case,family=binomial,
+             corstr="exchangeable", scale.fix=T,data = data3)
Beginning Cgee S-function, @(#) geeformula.q 4.13 98/01/27
running glm to get initial regression estimate
(Intercept)      gender      question
0.091085560 0.003581948 -0.074688363
> summary(model33)

```

GEE: GENERALIZED LINEAR MODELS FOR DEPENDENT DATA  
 gee S-function, version 4.13 modified 98/01/27 (1998)

Model:

Link: Logit  
Variance to Mean Relation: Binomial  
Correlation Structure: Exchangeable

call:

```
gee(formula = response ~ gender + question, id = case, data = data3,
    family = binomial, corstr = "exchangeable", scale.fix = T)
```

### Summary of Residuals:

| Min        | 1Q         | Median     | 3Q        | Max       |
|------------|------------|------------|-----------|-----------|
| -0.5049607 | -0.4862922 | -0.4668213 | 0.5137078 | 0.5331787 |

Coefficients:

|             | Estimate     | Naive S.E. | Naive z     | Robust S.E. | Robust z    |
|-------------|--------------|------------|-------------|-------------|-------------|
| (Intercept) | 0.091155117  | 0.07149404 | 1.27500299  | 0.07221536  | 1.26226763  |
| gender      | 0.003376898  | 0.08787193 | 0.03842977  | 0.08783897  | 0.03844419  |
| question    | -0.074688407 | 0.01406974 | -5.30844210 | 0.01487266  | -5.02186055 |

Estimated Scale Parameter: 1

Number of Iterations: 2

## Working Correlation

|      | [,1]      | [,2]      | [,3]      |
|------|-----------|-----------|-----------|
| [1,] | 1.0000000 | 0.8172912 | 0.8172912 |
| [2,] | 0.8172912 | 1.0000000 | 0.8172912 |
| [3,] | 0.8172912 | 0.8172912 | 1.0000000 |

>

```
> #Comparing to part(c)
```

```
> summary(model32)
```

## GEE: GENERALIZED LINEAR MODELS FOR DEPENDENT DATA

gee S-function, version 4.13 modified 98/01/27 (1998)

### Model :

Link: Logit  
Variance to Mean Relation: Binomial  
Correlation Structure: Exchangeable

call:

```
gee(formula = response ~ gender + as.factor(question), id = case,
    data = data3, family = binomial, corstr = "exchangeable",
    scale.fix = T)
```

### Summary of Residuals:

| Min        | 1Q         | Median     | 3Q        | Max       |
|------------|------------|------------|-----------|-----------|
| -0.5068644 | -0.4825396 | -0.4687095 | 0.5174604 | 0.5312905 |

Coefficients:

|                      | Estimate     | Naive S.E. | Naive z     | Robust S.E. |
|----------------------|--------------|------------|-------------|-------------|
| (Intercept)          | 0.024021377  | 0.06774107 | 0.35460579  | 0.06779334  |
| gender               | 0.003437873  | 0.08787462 | 0.03912248  | 0.08784072  |
| as.factor(question)2 | -0.097329120 | 0.02811568 | -3.46173752 | 0.02753161  |
| as.factor(question)3 | -0.149347107 | 0.02813360 | -5.30849707 | 0.02973865  |

  

|                      | Robust z    |
|----------------------|-------------|
| (Intercept)          | 0.35433237  |
| gender               | 0.03913758  |
| as.factor(question)2 | -3.53517666 |
| as.factor(question)3 | -5.02198729 |

Estimated Scale Parameter: 1

Number of Iterations: 2

Working Correlation

|      | [,1]      | [,2]      | [,3]      |
|------|-----------|-----------|-----------|
| [1,] | 1.0000000 | 0.8173308 | 0.8173308 |
| [2,] | 0.8173308 | 1.0000000 | 0.8173308 |
| [3,] | 0.8173308 | 0.8173308 | 1.0000000 |

>

> #(h)

> model34=glm(response~gender+as.factor(question),family=binomial,data=data3)

> summary(model34)

Call:

glm(formula = response ~ gender + as.factor(question), family = binomial, data = data3)

Deviance Residuals:

| Min    | 1Q     | Median | 3Q    | Max   |
|--------|--------|--------|-------|-------|
| -1.189 | -1.148 | -1.125 | 1.207 | 1.231 |

Coefficients:

|                      | Estimate  | Std. Error | z value | Pr(> z ) |
|----------------------|-----------|------------|---------|----------|
| (Intercept)          | 0.023940  | 0.055528   | 0.431   | 0.6664   |
| gender               | 0.003582  | 0.054138   | 0.066   | 0.9472   |
| as.factor(question)2 | -0.097329 | 0.065783   | -1.480  | 0.1390   |
| as.factor(question)3 | -0.149347 | 0.065825   | -2.269  | 0.0233 * |

---

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

```
>  
> #Comparing to part(c)  
> summary(model32)
```

GEE: GENERALIZED LINEAR MODELS FOR DEPENDENT DATA  
gee S-function, version 4.13 modified 98/01/27 (1998)

Model:  
Link: Logit  
Variance to Mean Relation: Binomial  
Correlation Structure: Exchangeable

Call:  
gee(formula = response ~ gender + as.factor(question), id = case,  
data = data3, family = binomial, corstr = "exchangeable",  
scale.fix = T)

Summary of Residuals:

|  | Min        | 1Q         | Median     | 3Q        | Max       |
|--|------------|------------|------------|-----------|-----------|
|  | -0.5068644 | -0.4825396 | -0.4687095 | 0.5174604 | 0.5312905 |

Coefficients:

|                      | Estimate     | Naive S.E. | Naive z     | Robust S.E. |
|----------------------|--------------|------------|-------------|-------------|
| (Intercept)          | 0.024021377  | 0.06774107 | 0.35460579  | 0.06779334  |
| gender               | 0.003437873  | 0.08787462 | 0.03912248  | 0.08784072  |
| as.factor(question)2 | -0.097329120 | 0.02811568 | -3.46173752 | 0.02753161  |
| as.factor(question)3 | -0.149347107 | 0.02813360 | -5.30849707 | 0.02973865  |

  

|                      | Robust z    |
|----------------------|-------------|
| (Intercept)          | 0.35433237  |
| gender               | 0.03913758  |
| as.factor(question)2 | -3.53517666 |
| as.factor(question)3 | -5.02198729 |

Estimated Scale Parameter: 1  
Number of Iterations: 2

Working Correlation

|      | [,1]      | [,2]      | [,3]      |
|------|-----------|-----------|-----------|
| [1,] | 1.0000000 | 0.8173308 | 0.8173308 |
| [2,] | 0.8173308 | 1.0000000 | 0.8173308 |
| [3,] | 0.8173308 | 0.8173308 | 1.0000000 |

```
>  
> #The coefficients estimates of these two models are the same  
> #The standard error of exchangeable model is smaller than the standard error  
of independence model
```

```
>
> #(i)
> #It is because that as for GEE, there is a large correlation within subject, so
> #standard error is smaller than GLM. However, using the same data, the total
> #standard errors are always the same. Therefore, GEE has larger between-subject
> #standard error.
>
>
> #
```