## **Homework 5**

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## Problem 1, Part 1

a)

```
Titanic<-read.csv("http://users.stat.umn.edu/~parky/TitanicPartial.csv")
attach(Titanic)
table(Pclass,Survived)

## Survived
## Pclass 0 1
## 1 64 122
## 2 90 83
## 3 270 85</pre>
```

(i) First Class:

```
Prob1=122/(122+64)
Odds1=Prob1/(1-Prob1)
Odds1
## [1] 1.90625
```

Second Class:

```
Prob2=83/(83+90)
Odds2=Prob2/(1-Prob2)
Odds2
## [1] 0.9222222
```

Third Class:

```
Prob3=85/(85+270)
Odds3=Prob3/(1-Prob3)
Odds3
## [1] 0.3148148
```

(ii) Odds ratio:

```
Odds1/Odds3
## [1] 6.055147
```

(iii) Passengers in First Class were over 6 times more likely to survive than passengers in Third Class.

```
b)
m0<-glm(Survived~as.factor(Pclass), family=binomial, data=Titanic)</pre>
summary(m0)
##
## Call:
## glm(formula = Survived ~ as.factor(Pclass), family = binomial,
##
       data = Titanic)
##
## Deviance Residuals:
       Min
                       Median
##
                  1Q
                                              Max
## -1.4607 -0.7399 -0.7399
                                 0.9184
                                           1.6908
##
## Coefficients:
                       Estimate Std. Error z value Pr(>|z|)
##
                                               4.180 2.92e-05 ***
## (Intercept)
                         0.6451
                                     0.1543
## as.factor(Pclass)2 -0.7261
                                     0.2168
                                             -3.350 0.000808 ***
## as.factor(Pclass)3 -1.8009
                                     0.1982 -9.086 < 2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
       Null deviance: 964.52 on 713 degrees of freedom
## Residual deviance: 869.81 on 711 degrees of freedom
## AIC: 875.81
##
## Number of Fisher Scoring iterations: 4
(i) Logit Form: log(theta/(1-theta)) = 0.6451-0.7261*x_1-1.8009*x_2
Probability Form: theta=1/(1+e^{(-(0.6451-0.7261*x_1-1.8009*x_2)))}
(ii) For a person in First Class (the default), the odds of survival were
exp(0.6451)
## [1] 1.906178
In other words, beta0=log(odds of survival for 1st class)
(iii) The odds ratio for a person in third class is
exp(1.8009)
## [1] 6.055095
In other words, beta2=-log(odds ratio)
(iv) The estimated probability of surviving for a randomly selected person in First Class is
1/(1+exp(-0.6451))
## [1] 0.6559054
```

## Problem 1, Part 2

(a) It looks like females in 1st class had the highest odds of survival.

(b)

```
m1=glm(Survived~as.factor(Pclass)+Sex, family=binomial, data=Titanic)
m2=glm(Survived~as.factor(Pclass)+ as.factor(Pclass)*Sex + Age,
family=binomial, data=Titanic)
summary(m1)
##
## Call:
## glm(formula = Survived ~ as.factor(Pclass) + Sex, family = binomial,
      data = Titanic)
##
## Deviance Residuals:
      Min
                     Median
                10
                                   30
                                           Max
## -2.2029
           -0.7371
                    -0.4556
                               0.6617
                                        2.1526
## Coefficients:
##
                      Estimate Std. Error z value Pr(>|z|)
                                   0.2385
                                            9.784 < 2e-16 ***
## (Intercept)
                        2.3338
## as.factor(Pclass)2 -0.9261
                                   0.2580
                                          -3.590 0.000331 ***
## as.factor(Pclass)3 -1.9748
                                   0.2379 -8.300 < 2e-16 ***
                                   0.2032 -12.658 < 2e-16 ***
## Sexmale
                       -2.5721
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
       Null deviance: 964.52 on 713
                                     degrees of freedom
##
## Residual deviance: 672.43 on 710 degrees of freedom
## AIC: 680.43
## Number of Fisher Scoring iterations: 4
summary(m2)
##
## Call:
## glm(formula = Survived ~ as.factor(Pclass) + as.factor(Pclass) *
       Sex + Age, family = binomial, data = Titanic)
##
## Deviance Residuals:
##
      Min
                1Q
                     Median
                                   3Q
                                           Max
## -3.1093 -0.6339 -0.4620
                               0.3896
                                        2.5299
##
## Coefficients:
                               Estimate Std. Error z value Pr(>|z|)
                               4.909854 0.682782 7.191 6.43e-13 ***
## (Intercept)
```

```
## as.factor(Pclass)2
                                         0.734046 -1.581 0.11391
                             -1.160437
## as.factor(Pclass)3
                                         0.648309 -6.431 1.27e-10 ***
                             -4.169187
                                         0.628523 -5.790 7.03e-09 ***
## Sexmale
                             -3.639225
## Age
                             -0.041934
                                         0.008184 -5.124 2.99e-07 ***
## as.factor(Pclass)2:Sexmale -0.668839
                                         0.815889 -0.820 0.41235
## as.factor(Pclass)3:Sexmale 2.192651
                                                   3.200 0.00138 **
                                         0.685268
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
      Null deviance: 964.52 on 713 degrees of freedom
##
## Residual deviance: 613.43 on 707 degrees of freedom
## AIC: 627.43
##
## Number of Fisher Scoring iterations: 6
c)
pred1=predict(m1, newdata = data.frame(Sex = 'male', Age = 60, Pclass = 3))
1/(1+exp(-pred1))
##
## 0.09858161
pred2=predict(m2, newdata = data.frame(Sex = 'male', Age = 60, Pclass = 3))
1/(1+exp(-pred2))
##
           1
## 0.03834764
```

The probability of survival is higher using model 1.

d) H<sub>0</sub>: beta4, beta5, and beta6 all equal 0 H<sub>1</sub>: At least one of these betas does not equal 0 Test statistic:

```
672.43 - 613.43

## [1] 59

pchisq(59, df=3, lower=F)

## [1] 9.613049e-13
```

Since this is much smaller than 0.05, we reject the null hypothesis and conclude that model 1 is not adequate compared to model 2.