# Exam3\_Jyoti\_Chaudhary

Jyoti Chaudhary

May 7, 2018

#### Problem 1

a.

```
data1 <- read.csv(paste(getwd(),"/Test_S18_Data.csv",sep = ""),header=T)
mdl1= glm ( data1$Owner ~ data1$Income, data = data1, family = binomial)
summary(mdl1) ##### PARAMETER ESTIMATES</pre>
```

```
##
## Call:
## glm(formula = data1$Owner ~ data1$Income, family = binomial,
##
      data = data1)
##
## Deviance Residuals:
##
      Min
               10 Median
                                 3Q
                                         Max
## -1.7236 -0.6104 -0.4412 0.6081
                                      1.5858
##
## Coefficients:
               Estimate Std. Error z value Pr(>|z|)
##
## (Intercept) -6.22899 2.67576 -2.328 0.0199 *
## data1$Income 0.13199
                          0.06072 2.174 0.0297 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
##
      Null deviance: 26.920 on 19 degrees of freedom
## Residual deviance: 17.634 on 18 degrees of freedom
## AIC: 21.634
##
## Number of Fisher Scoring iterations: 5
```

```
# The coefficients obtained are:
mdl1$coefficients
```

```
## (Intercept) data1$Income
## -6.2289868 0.1319937
```

The fitted model is:

```
\hat{y} = \hat{\pi} = rac{1}{1 + e^{-(-6.2289 + 0.1319x)}} = rac{1}{1 + e^{6.2289 - 0.1319x}}
```

where x = Income

If the model is good at 5% level?

```
anova(mdl1, test="Chisq")
```

```
## Analysis of Deviance Table
## Model: binomial, link: logit
##
## Response: data1$0wner
##
## Terms added sequentially (first to last)
##
##
                Df Deviance Resid. Df Resid. Dev Pr(>Chi)
##
## NULL
                                          26.921
## data1$Income 1
                     9.2869
                                   18
                                          17.634 0.002308 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

hoslem.test(data1\$Owner, fitted(mdl1), g=8)

```
##
## Hosmer and Lemeshow goodness of fit (GOF) test
##
## data: data1$Owner, fitted(mdl1)
## X-squared = 3.4905, df = 6, p-value = 0.7452
```

vcov(mdl1)

```
## (Intercept) data1$Income
## (Intercept) 7.1596962 -0.158412390
## data1$Income -0.1584124 0.003687358
```

```
# Test of model significance
0.1319/sqrt(0.003687)
```

```
## [1] 2.172242
```

```
(0.1319/sqrt(0.003687))^2
```

```
## [1] 4.718636
```

```
qchisq(.95,df = 1)
```

```
## [1] 3.841459
```

Null deviance= 26.921 Residual deviance= 17.634, it shows a significant improvement over null model. As D/df = 0.98 which is close to 1.00, this indicates that the model is adequate and good fit for the data. Also, the p-value (=0.002308) is very low which further indicates that model is adequate.

We can also do HL-test to test goodness of fit which shows Chi-square = 3.4905 with 6 degrees of freedom and p-value = 0.7452. Hence it is good fit to the data.

Also, as  $(0.1319/\text{sqrt}(0.003687))^2 > \text{qchisq}(.95,\text{df} = 1)$ , the model is adequate and significant.

## b)

Find and interpret the odds ratio for "Income".

```
OR=exp(coef(mdl1)[2])
OR
```

```
## data1$Income
## 1.141101
```

This odds ratio implies that for every unit increase in speed, the odds of hitting the target decrease by 1.75%.

Using odds ratio we can say that for every 1 unit increase of income, the odds of owning the device increase by 100x(1.141101 - 1) = 14.1%.

### c)

Predict the probability of ownership if income is \$40,000. Also nd a 98% condence interval for the true probability.

```
nwdt=with(data1, data.frame(Income=40.0)) ##### NEW DATA POINT
nwdt
```

```
## Income
## 1 40
```

```
pct=0.98

nwdt2=subset(cbind(nwdt,predict(mdl1,newdata=nwdt, type="link", se=TRUE)),select = -c(residual.s cale))
```

```
## Warning: 'newdata' had 1 row but variables found have 20 rows
```

```
nwdt2
```

```
##
      Income
                      fit
                             se.fit
## 1
          40 -2.49356611 1.0708517
## 2
          40 -2.37477182 1.0258664
## 3
          40 -2.18998069 0.9578898
## 4
          40 -2.11078449 0.9296156
          40 -1.99199019 0.8883196
## 5
## 6
          40 -1.95239210 0.8748798
## 7
          40 -1.60920857 0.7667092
## 8
          40 -1.51681301 0.7406436
## 9
          40 -1.43761681 0.7195498
## 10
          40 -0.92284153 0.6182707
## 11
          40 -0.76444913 0.6026069
## 12
          40 -0.72485103 0.6000113
## 13
          40 -0.63245547 0.5960797
## 14
          40
              0.04071221 0.6558365
## 15
              0.77987672 0.8562786
          40
              1.22865518 1.0149217
## 16
          40
## 17
          40
              1.25505391 1.0247843
## 18
              1.71703173 1.2041506
          40
              4.27770879 2.3030711
## 19
          40
## 20
          40
              4.59449358 2.4441485
```

```
nwdt3=within(nwdt2,{PredictedProb <- plogis(fit)
LL <- plogis(fit - (qnorm((1+pct)/2) * se.fit))
UL <- plogis(fit + (qnorm((1+pct)/2) * se.fit))})
nwdt3</pre>
```

```
##
                     fit
                             se.fit
                                           UL
                                                        LL PredictedProb
      Income
          40 -2.49356611 1.0708517 0.4994019 0.006795070
## 1
                                                              0.07631045
## 2
          40 -2.37477182 1.0258664 0.5029375 0.008481974
                                                              0.08511681
## 3
          40 -2.18998069 0.9578898 0.5095999 0.011910351
                                                              0.10065384
          40 -2.11078449 0.9296156 0.5129533 0.013742912
                                                              0.10805304
## 4
          40 -1.99199019 0.8883196 0.5186289 0.016981046
## 5
                                                              0.12004647
## 6
          40 -1.95239210 0.8748798 0.5207088 0.018205346
                                                              0.12429276
## 7
          40 -1.60920857 0.7667092 0.5434957 0.032519958
                                                              0.16669852
## 8
          40 -1.51681301 0.7406436 0.5513636 0.037694865
                                                              0.17993130
## 9
          40 -1.43761681 0.7195498 0.5588032 0.042633741
                                                              0.19191467
          40 -0.92284153 0.6182707 0.6260882 0.086183239
## 10
                                                              0.28437927
## 11
          40 -0.76444913 0.6026069 0.6541710 0.102815771
                                                              0.31768109
## 12
          40 -0.72485103 0.6000113 0.6617234 0.107102453
                                                              0.32632565
## 13
          40 -0.63245547 0.5960797 0.6801004 0.117207520
                                                              0.34695398
## 14
          40
              0.04071221 0.6558365 0.8272721 0.184674642
                                                              0.51017665
              0.77987672 0.8562786 0.9411371 0.229325255
## 15
          40
                                                              0.68565354
## 16
          40
              1.22865518 1.0149217 0.9731355 0.243717419
                                                              0.77358311
## 17
              1.25505391 1.0247843 0.9743957 0.244354779
                                                              0.77817349
          40
## 18
          40
              1.71703173 1.2041506 0.9892102 0.252704183
                                                              0.84774611
              4.27770879 2.3030711 0.9999346 0.253499251
## 19
          40
                                                              0.98631545
## 20
              4.59449358 2.4441485 0.9999657 0.251346053
                                                              0.98999380
          40
```

```
data1$Income2 <- data1$Income^2
mdl2= glm ( data1$Owner ~ data1$Income + data1$Income2 , data = data1, family = binomial)
summary(mdl2) ##### PARAMETER ESTIMATES</pre>
```

```
##
## Call:
## glm(formula = data1$Owner ~ data1$Income + data1$Income2, family = binomial,
##
      data = data1)
##
## Deviance Residuals:
      Min 1Q Median 3Q
##
                                        Max
## -1.8217 -0.5728 -0.3388 0.5787
                                     1.5251
##
## Coefficients:
##
                  Estimate Std. Error z value Pr(>|z|)
## (Intercept) -11.544155 7.989792 -1.445
                                               0.148
## data1$Income 0.360315 0.315612 1.142
                                               0.254
## data1$Income2 -0.002307 0.002950 -0.782
                                               0.434
##
## (Dispersion parameter for binomial family taken to be 1)
##
##
      Null deviance: 26.920 on 19 degrees of freedom
## Residual deviance: 17.121 on 17 degrees of freedom
## AIC: 23.121
##
## Number of Fisher Scoring iterations: 5
```

```
anova(mdl2, test="Chisq")
```

```
## Analysis of Deviance Table
##
## Model: binomial, link: logit
##
## Response: data1$0wner
##
## Terms added sequentially (first to last)
##
##
                Df Deviance Resid. Df Resid. Dev Pr(>Chi)
##
## NULL
                                  19
                                         26.921
## data1$Income 1 9.2869
                                  18
                                         17.634 0.002308 **
## data1$Income2 1 0.5122
                                  17
                                       17.121 0.474182
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

The deviance has NOT significantly reduced due to quadratic term with a high p-value of 0.474182. Hence its evident that the interaction term is not required in the model.

To test Beta3 = 0 vs Beta3 not = 0, Difference in Deviance = 17.634 - 17.121 = 0.513 which is lesser than chi-square(0.1,1) = 2.706. Hence the quadratic term is not significant at 10% level.

#### Problem 3

a.

```
data2 <- read.csv(paste(getwd(),"/Test_S18_Data_2.csv",sep = ""),header=T)

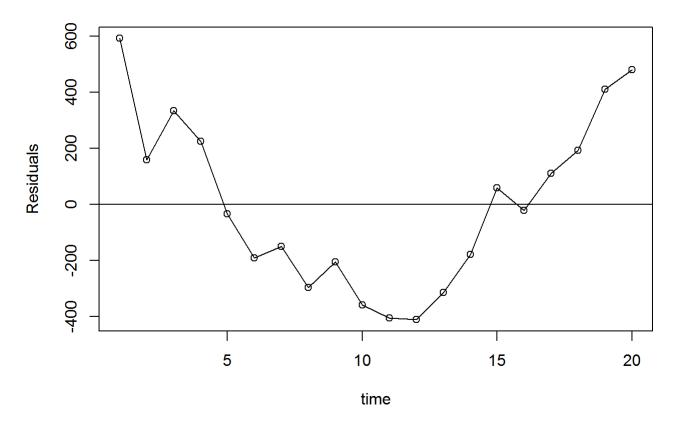
# fitting linear regression data

mod3 = lm(Yt ~ t , data = data2)

# plotting residuals against time

plot(data2$t, mod3$res, ylab="Residuals", xlab="time", main="Residual Plot over Time - Model-1", type="o")
abline(0, 0)</pre>
```

#### Residual Plot over Time - Model-1



```
# the plot indicates positive autocorrelation
 # Dubin watson test to determine correlation
 library(car)
 durbinWatsonTest(mod3, max.lag=1, simulate=TRUE, reps=10000, method="normal", alternative="posit
 ive")
     lag Autocorrelation D-W Statistic p-value
 ##
 ##
                 0.681758
                               0.308722
 ## Alternative hypothesis: rho > 0
n = 20, k = 1, dw = 0.308722, alpha = 0.05, dL = 1.201, dU = 1.411 As dw < dL, reject H0, which means there is
evidence to support the conclusion that the residuals are positively autocorrelated.
cochrane orcutt method
 ##### CALCULATING RHOHAT USING FORMULA ##########
 res3=resid(mod3)
 n= length(res3)
 rho = sum(res3[1:(n-1)]*res3[2:n])/sum(res3^2)
 rho
 ## [1] 0.681758
 library(DataCombine)
 ## Warning: package 'DataCombine' was built under R version 3.4.4
 data3=slide(data=data2, Var="Yt", TimeVar="t", NewVar="Yt_1", slideBy = -1,
            keepInvalid = FALSE, reminder = TRUE)
 ##
```

## Lagging t by 1 time units.

```
##
             Υt
                  Yt_1 t_1
       t
       1 4710.0
## 1
                    NA
                         NA
## 2
       2 4187.7 4710.0
                          1
                          2
## 3
       3 4275.6 4187.7
## 4
       4 4076.9 4275.6
                          3
## 5
       5 3731.4 4076.9
                          4
## 6
       6 3484.3 3731.4
                          5
## 7
       7 3437.2 3484.3
                          6
       8 3203.2 3437.2
                          7
## 8
## 9
       9 3206.2 3203.2
                          8
## 10 10 2963.1 3206.2
                          9
## 11 11 2828.5 2963.1
## 12 12 2734.8 2828.5
## 13 13 2743.9 2734.8
## 14 14 2789.6 2743.9
## 15 15 2939.4 2789.6
## 16 16 2770.6 2939.4
## 17 17 2815.0 2770.6
## 18 18 2808.4 2815.0
## 19 19 2938.1 2808.4
## 20 20 2918.6 2938.1
```

```
pst2$yprime= pst2$Yt - rho*pst2$Yt_1
pst2$xprime= pst2$t - rho*pst2$t_1
pst2
```

```
##
             Υt
                  Yt_1 t_1
                                       xprime
       t
                              yprime
## 1
       1 4710.0
                    NA
                       NA
                                  NA
                                            NA
## 2
       2 4187.7 4710.0
                         1 976.6200 1.318242
## 3
       3 4275.6 4187.7
                         2 1420.6022 1.636484
       4 4076.9 4275.6
                         3 1161.9757 1.954726
## 4
## 5
       5 3731.4 4076.9
                         4 951.9410 2.272968
## 6
       6 3484.3 3731.4
                         5 940.3883 2.591210
## 7
       7 3437.2 3484.3
                         6 1061.7507 2.909452
## 8
       8 3203.2 3437.2
                         7 859.8615 3.227694
## 9
       9 3206.2 3203.2
                         8 1022.3929 3.545936
## 10 10 2963.1 3206.2
                           777.2476 3.864178
## 11 11 2828.5 2963.1
                        10
                            808.3830 4.182420
## 12 12 2734.8 2828.5
                            806.4476 4.500662
                        11
## 13 13 2743.9 2734.8
                        12
                            879.4283 4.818904
## 14 14 2789.6 2743.9
                            918.9243 5.137147
                        13
## 15 15 2939.4 2789.6
                        14 1037.5680 5.455389
## 16 16 2770.6 2939.4
                        15
                            766.6406 5.773631
## 17 17 2815.0 2770.6
                        16
                            926.1214 6.091873
## 18 18 2808.4 2815.0
                        17
                            889.2513 6.410115
## 19 19 2938.1 2808.4
                        18 1023.4509 6.728357
## 20 20 2918.6 2938.1 19 915.5269 7.046599
```

```
mod5 = lm(yprime ~ xprime , data = pst2)
mod5
```

```
summary(mod5)
```

```
##
## Call:
## lm(formula = yprime ~ xprime, data = pst2)
##
## Residuals:
      Min
               1Q Median
##
                              3Q
                                     Max
## -189.72 -107.33 -0.07
                           63.65 369.68
##
## Coefficients:
              Estimate Std. Error t value Pr(>|t|)
##
## (Intercept) 1112.59 84.24 13.208 2.29e-10 ***
               -37.69 18.59 -2.027 0.0586.
## xprime
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 141.3 on 17 degrees of freedom
    (1 observation deleted due to missingness)
## Multiple R-squared: 0.1947, Adjusted R-squared: 0.1473
## F-statistic: 4.109 on 1 and 17 DF, p-value: 0.05863
```

```
dw2=durbinWatsonTest(lm(yprime ~ xprime,data = pst2) , max.lag=1, alternative="positive")
dw2
```

```
## lag Autocorrelation D-W Statistic p-value
## 1 0.09297401 1.778269 0.213
## Alternative hypothesis: rho > 0
```

```
n = 19, k = 1, dw = 1.778269, alpha = 0.05, dL = 1.180, dU = 1.401
```

As dw > dU , which means that there is no auto-correlation present among errors. So we conclude that there is no problem with autocorrelated errors in the transformed model. The CochraneOrcutt method has been effective in removing the autocorrelation.