R Notebook

Code ▼

This is an R Markdown (http://rmarkdown.rstudio.com) Notebook. When you execute code within the notebook, the results appear beneath the code.

Try executing this chunk by clicking the *Run* button within the chunk or by placing your cursor inside it and pressing *Ctrl+Shift+Enter*.

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```
#question 1
#part a
library(openxlsx)
xlsxFile <- system.file("extdata","hw_9_data.xlsx", package = "openxlsx")
df <- read.xlsx(xlsxFile = "hw_9_data.xlsx", sheet = 1, skipEmptyRows = FALSE)

mdl1= glm( y ~ x, data = df, family = binomial)
#part b
summary(mdl1)</pre>
```

```
Call:
glm(formula = y \sim x, family = binomial, data = df)
Deviance Residuals:
   Min
             1Q Median
                               3Q
                                       Max
-2.0620 -0.4868 0.3915 0.5476
                                    2.1682
Coefficients:
             Estimate Std. Error z value Pr(>|z|)
(Intercept) 6.070884
                       2.108996
                                  2.879 0.00399 **
            -0.017705
                       0.006076 -2.914 0.00357 **
Х
Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for binomial family taken to be 1)
    Null deviance: 34.617 on 24 degrees of freedom
Residual deviance: 20.364 on 23 degrees of freedom
AIC: 24.364
Number of Fisher Scoring iterations: 4
```

```
#install.packages("ResourceSelection")
library(ResourceSelection)
hoslem.test(df$y,fitted(mdl1), g=8) ###### NOT TESTED
```

```
Hosmer and Lemeshow goodness of fit (GOF) test
data: df$y, fitted(mdl1)
X-squared = 4.8315, df = 6, p-value = 0.5656
                                                                                              Hide
#part c
OR=exp(coef(mdl1)) ####### ODDS RATIO OF THE VARIABLES
OR
(Intercept)
433.0632952
            0.9824511
                                                                                              Hide
#part d
df$xsq=df$x^2
mdl1b = glm ( y \sim x + xsq, data = df, family = binomial)
summary(mdl1b) ##### PARAMETER ESTIMATES
Call:
glm(formula = y \sim x + xsq, family = binomial, data = df)
Deviance Residuals:
   Min
              1Q
                   Median
                                3Q
                                        Max
-2.0642 -0.4879
                   0.3890
                            0.5469
                                     2.1651
Coefficients:
              Estimate Std. Error z value Pr(>|z|)
(Intercept) 6.193e+00 9.030e+00
                                    0.686
                                             0.493
            -1.847e-02 5.513e-02 -0.335
                                             0.738
Х
             1.100e-06 7.921e-05
xsq
                                    0.014
                                             0.989
(Dispersion parameter for binomial family taken to be 1)
    Null deviance: 34.617 on 24 degrees of freedom
Residual deviance: 20.363 on 22 degrees of freedom
AIC: 26.363
Number of Fisher Scoring iterations: 4
```

```
#question 1
#part a
library(openxlsx)
xlsxFile <- system.file("extdata","hw_9_data.xlsx", package = "openxlsx")
df <- read.xlsx(xlsxFile = "hw_9_data.xlsx", sheet = 2, skipEmptyRows = FALSE)

mdl2= glm( y ~ x1+x2, data = df, family = binomial)
#part b
summary(mdl2)</pre>
```

```
Call:
glm(formula = y \sim x1 + x2, family = binomial, data = df)
Deviance Residuals:
   Min
             10 Median
                               3Q
                                      Max
-1.5635 -0.8045 -0.1397 0.9535
                                   1.7915
Coefficients:
             Estimate Std. Error z value Pr(>|z|)
(Intercept) -7.047e+00 4.674e+00 -1.508
                                           0.132
x1
            7.382e-05 6.371e-05
                                           0.247
                                   1.159
x2
            9.879e-01 5.274e-01
                                 1.873
                                           0.061 .
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for binomial family taken to be 1)
    Null deviance: 27.726 on 19 degrees of freedom
Residual deviance: 21.082 on 17 degrees of freedom
AIC: 27.082
Number of Fisher Scoring iterations: 5
```

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```
hoslem.test(df$y,fitted(mdl2), g=8)
```

```
Hosmer and Lemeshow goodness of fit (GOF) test

data: df$y, fitted(mdl2)

X-squared = 4.7136, df = 6, p-value = 0.581
```

```
#part c
OR=exp(coef(md12))
OR
```

```
(Intercept) x1 x2
0.0008699617 1.0000738195 2.6855513881
```

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```
#part d
nwdt=with(df, data.frame(x1=45000, x2=5))
##################
pct=0.95
########################
nwdt2=subset(cbind(nwdt,predict(mdl2,newdata=nwdt, type="link", se=TRUE)),select = -c(residual.s cale))
#################################
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>
```

	x1 <dbl></dbl>	x2 <dbl></dbl>	fit <dbl></dbl>	se.fit <dbl></dbl>	UL <dbl></dbl>	LL <dbl></dbl>	PredictedProb <dbl></dbl>
1	45000	5	1.214124	0.8630815	0.9481291	0.3828464	0.7710279
1 row							

```
#part e
df$x12=df$x1*df$x2
mdl2b= glm ( y ~ x1 + x2 + x12, data = df, family = binomial)
summary(mdl2b)
```

```
Call:
glm(formula = y \sim x1 + x2 + x12, family = binomial, data = df)
Deviance Residuals:
     Min
                     Median
                10
                                    3Q
                                             Max
-1.63981 -0.62754 -0.05642
                              0.66213
                                        1.85666
Coefficients:
              Estimate Std. Error z value Pr(>|z|)
(Intercept) 3.144e-01 6.394e+00
                                   0.049
                                             0.961
x1
            -1.411e-04 1.412e-04 -0.999
                                             0.318
x2
            -2.462e+00 2.081e+00 -1.183
                                             0.237
x12
             1.014e-04 6.297e-05
                                  1.610
                                             0.107
(Dispersion parameter for binomial family taken to be 1)
    Null deviance: 27.726 on 19 degrees of freedom
Residual deviance: 16.551 on 16 degrees of freedom
AIC: 24.551
Number of Fisher Scoring iterations: 6
                                                                                              Hide
#part f
pct2=0.95
CI_lower1 = coefficients(mdl2)[2] - (qnorm((1+pct2)/2))*summary(mdl2)$coefficients[2,2]
CI upper1 = coefficients(mdl2)[2] + (qnorm((1+pct2)/2))*summary(mdl2)$coefficients[2,2]
CI lower1
           x1
-5.105847e-05
                                                                                              Hide
CI upper1
          x1
0.0001986921
                                                                                              Hide
CI_lower2 = coefficients(mdl2)[3] - (qnorm((1+pct2)/2))*summary(mdl2)$coefficients[3,3]
CI\_upper2 = coefficients(mdl2)[3] + (qnorm((1+pct2)/2))*summary(mdl2)$coefficients[3,3]
CI lower2
```

```
x2
-2.683594

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CI_upper2

x2
4.659366
```

Add a new chunk by clicking the *Insert Chunk* button on the toolbar or by pressing *Ctrl+Alt+I*.

When you save the notebook, an HTML file containing the code and output will be saved alongside it (click the *Preview* button or press *Ctrl+Shift+K* to preview the HTML file).

The preview shows you a rendered HTML copy of the contents of the editor. Consequently, unlike *Knit*, *Preview* does not run any R code chunks. Instead, the output of the chunk when it was last run in the editor is displayed.