

Week 5 - Logistic Regression Assignment

Due Yesterday by 1:29pm Points 15 Submitting a file upload

For the E-commerce data set, perform regression on "Churn/Stay" using the other numeric variables as the independent variables, in R. Use `glm(...family=binomial(link=logit))`.

- Generate the equation.
- Use the summary to view the diagnostics, and write a summary of the results
- Do you like the model or not? Why?
- How would you determine what is the "best" linear model?
- How will you add interactions? Which ones?
- Print results and graphs/charts, and submit Assignment 5.

Submission

✓ Turned In!
May 21 at 2:52pm (late)
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Week 5 - Logistic Regression Assignment

For the E-commerce data set, perform regression on "Churn/Stay" using the other numeric variables as the independent variables, in R. Use `glm(...family=binomial(link=logit))`.

a. Generate the equation.

Four equations generated -

```
m1<-glm(e$st~e$sls+e$sco+e$eco+e$csec+e$osec+e$qcec+e$spec+e$ad,
data =e, family = "binomial")
```

```
m2<-glm(e$st~e$sls+e$eco+e$qcec+e$ad, data =e, family = "binomial")
```

```
m3<-glm(e$st~e$sls*e$sco+e$eco*e$csec+e$osec*e$qcec+e$spec*e$ad, data
=e, family = "binomial")
```

```
m4<-glm(e$st~e$sls+e$sco*e$eco*e$csec+e$qcec*e$spec*e$ad, data =e,
family = "binomial")
```

b. Use the summary to view the diagnostics, and write a summary of the results

Ans-B- As seen below.

```
# Model - glm(m3) - AIC: 10803
# Model - glm(m4) -AIC: 10761
```

```
anova(object=m2, test="Chisq")
```

```
# Null- Resid deviance ==13842 at Df==9999
# e$ad- Resid deviance ==10863 at Df==9995
```

```
anova(object=m3, test="Chisq")
```

```
# Null- Resid deviance ==13842 at Df==9999
# e$spec:e$ad- Resid deviance ==10777 at Df==9987
```

```
anova(object=m4, test="Chisq")
```

```
# Null- Resid deviance ==13842 at Df==9999
# e$qcec:e$spec:e$ad- Resid deviance ==10729 at Df==9984
```

```
# c . Do you like the model or not? Why?
```

Ans-C-Model – glm(m4) is liked.

```
# d. How would you determine what is the "best" linear model?
```

Ans-D-Best model chosen basis various parameters of Confusion Matrix ,
anova(Chisq) and Hosmer-Lemeshow test.

```
# e. How will you add interactions? Which ones?
```

Interactions added in Model –glm(m3)

```
# f. Print results and graphs/charts, and submit Assignment 5.
```

```
#Reading Data
```

```
e<-read.csv("C:/STAT/Wk-5-EC-Logistic/ec.csv")
```

```
# First model -all independent variables.
```

```
attach(e)
```

```
m1<-glm(e$st~e$sls+e$sco+e$eco+e$csec+e$osec+e$qcec+e$spec+e$ad,  
data =e, family = "binomial")
```

```
summary(m1)
```

```
##
```

```
## Call:
```

```
## glm(formula = e$st ~ e$sls + e$sco + e$eco + e$csec + e$osec +  
## e$qcec + e$spec + e$ad, family = "binomial", data = e)
```

```
##
```

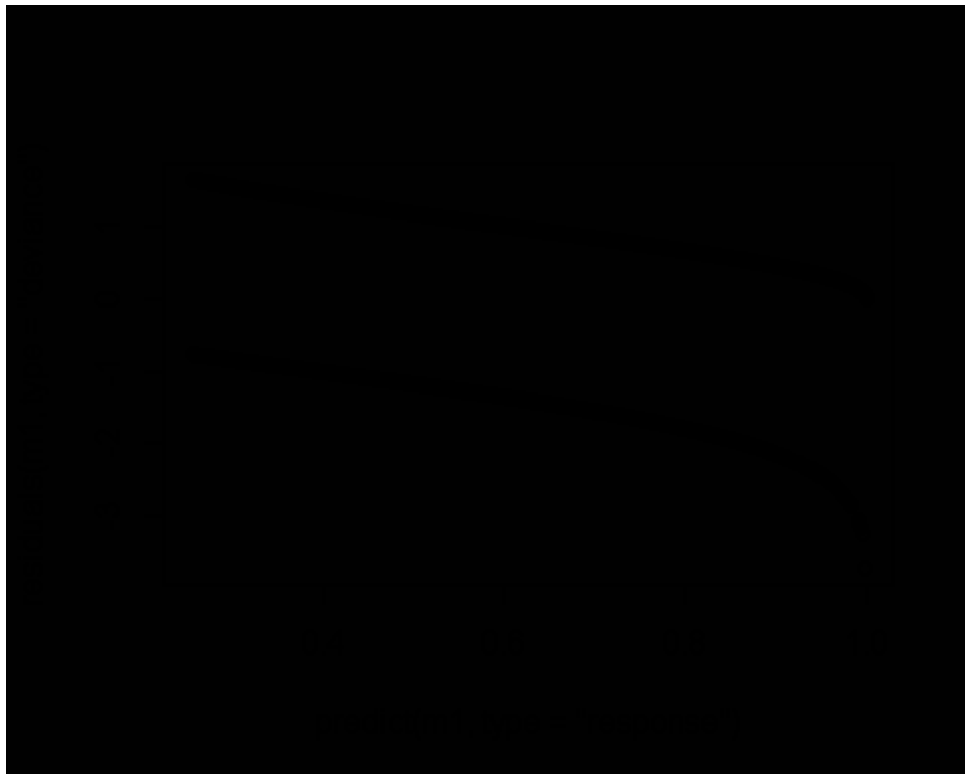
```
## Deviance Residuals:
```

```
##   Min      1Q  Median      3Q      Max  
## -3.7229 -0.8522  0.1636  0.9333  1.6499
```

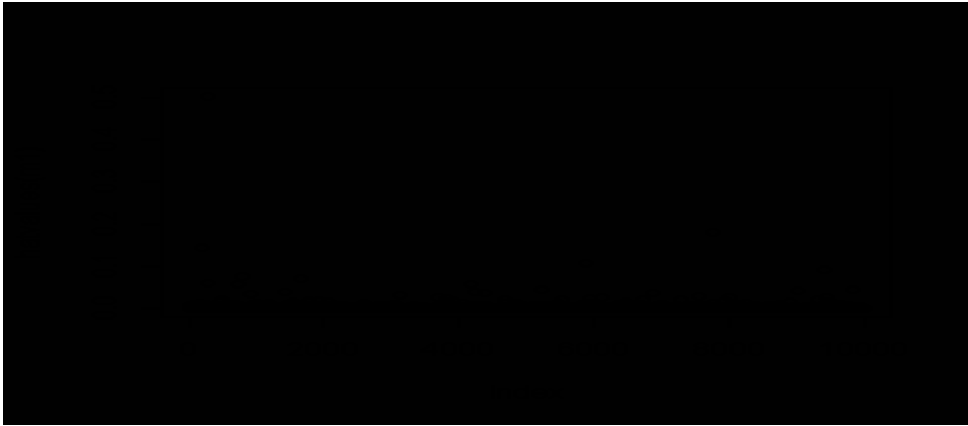
```
##
```

```
## Coefficients:
##           Estimate Std. Error z value Pr(>|z|)
## (Intercept) -1.109e+00  3.419e-02 -32.442 < 2e-16 ***
## e$sls       1.479e-05  2.753e-06  5.373 7.75e-08 ***
## e$sco      -4.030e-04  1.413e-03  -0.285  0.775
## e$eco       1.664e-03  2.344e-04  7.099 1.26e-12 ***
## e$csec     -1.628e-02  9.958e-03  -1.635  0.102
## e$osec      8.666e-03  1.003e-02   0.864  0.388
## e$qcec     -2.414e-03  2.641e-04  -9.139 < 2e-16 ***
## e$spec     -4.492e-03  2.877e-03  -1.561  0.118
## e$ad       7.214e-02  3.971e-03  18.169 < 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: 13842  on 9999  degrees of freedom
## Residual deviance: 10789  on 9991  degrees of freedom
## AIC: 10807
##
## Number of Fisher Scoring iterations: 6

# From Summary- FOUR undermentioned variables, not Significant to Churn
Status:-
# e$sco - p-Value==0.775
# e$csec- p-Value==0.102
# e$osec- p-Value==0.388
# e$spec- p-Value==0.118
## Remove FOUR variables - check if model is more Significant.
#
plot(predict(m1, type="response"),residuals(m1, type= "deviance"))
```



```
plot(hatvalues(m1))
```



```
plot(rstudent(m1))
```



```
plot(cooks.distance(m1))
```



Diagnostic plots helpful when response variable takes on many values.

Here response variable is Binomial - thus ignoring Diagnostic Plots.

*# Create **Confusion Matrix** for this glm().*
Compare Classification capability with other glm()'s.

```
ModelFit_m1<-m1$fit
ProbabModelFit_m1<-m1$fit>0.5
P1<-as.data.frame(ProbabModelFit_m1)
# "if" loop to convert TRUE and FALSE into 1 and 0.
v1<-vector()
for (i in 1:nrow(P1))
{
  if(P1[i,1]=="TRUE")
  {
    v1[i]<-1
  }
  if(P1[i,1]=="FALSE")
  {
    v1[i]<-0
  }
}
V1<-as.data.frame(cbind(e$st,v1))
colnames(V1)<-c("yhat","m1")
# write.table(V1,"C:/STAT/V1.csv", sep=",")
```

Confusion Matrix – Model – glm(m1)

TN(f[0,0])	FP (f[0,1])		
3904	869	4773	Total ACTUAL NO- 0
1897	3330	5227	Total ACTUAL YES - 1
FN (f[1,0])	TP(f[1,1])		
5801	4199		
Total [PRED] NO- 0 CHURNED Count	Total [[PRED] YES-1 STAYED Count		

Percentage of Misclassification OR - Misclassification Rate:	28%	//=(FP+FN)/n	Less is better.
Sensitivity OR - True Positive Rate	64%	//=TP/(TP+FN)	More is better.
Specificity OR - True Negative rate (1- FalsePositive Rate)	82%	//=TN/(TN+FP)	More is better.

Confusion Matrix – Model – glm(m2)

Percentage of Misclassification OR - Misclassification Rate:	28%	$//=(FP+FN)/n$	Less is better.
Sensitivity OR - True Positive Rate	65%	$//=TP/(TP+FN)$	More is better.
Specificity OR - True Negative rate (1 - FalsePositive Rate)	81%	$//=TN/(TN+FP)$	More is better.

As seen from the Conf Matrix – Model – glm(m1) , which has all the Independent variables is preferred as Significant.

Also as seen below - **Prefer a glm()- with Smaller Mean Deviance / Residual Deviance-m1==Residual deviance:10789 .Whereas- m2==Residual deviance: 10863**

Second glm()-as seen below -

```
m2<-glm(e$st~e$sls+e$eco+e$qcec+e$ad, data =e, family = "binomial")
summary(m2)
```

```
##
## Call:
## glm(formula = e$st ~ e$sls + e$eco + e$qcec + e$ad, family = "binomial",
## data = e)
##
## Deviance Residuals:
## Min 1Q Median 3Q Max
## -3.6332 -0.8610 0.1405 0.9674 1.6083
##
## Coefficients:
## Estimate Std. Error z value Pr(>|z|)
## (Intercept) -1.035e+00 3.193e-02 -32.419 < 2e-16 ***
## e$sls 6.453e-06 2.583e-06 2.498 0.0125 *
## e$eco 5.969e-04 1.842e-04 3.240 0.0012 **
## e$qcec -9.973e-04 1.966e-04 -5.071 3.95e-07 ***
## e$ad 6.181e-02 3.122e-03 19.798 < 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: 13842 on 9999 degrees of freedom
## Residual deviance: 10863 on 9995 degrees of freedom
## AIC: 10873
##
## Number of Fisher Scoring iterations: 6
```

Independent Variables - all FOUR have Significant p-Values.

Comparison between models- AIC - AIC of 2nd Model -m2 Higher at- AIC: 10873

Prefer a glm() - with Smaller Mean Deviance / Residual Deviance -

m1==Residual deviance: 10789 # Whereas - m2==Residual deviance: 10863

Confusion Matrix created in Excel as seen below -

```
ModelFit_m2<-m2$fit
```

```
ProbabModelFit_m2<-m2$fit>0.5
```

```
P2<-as.data.frame(ProbabModelFit_m2)
```

"if" loop to convert TRUE and FALSE into 1 and 0.

```
v2<-vector()
for (i in 1:nrow(P2))
{
  if(P2[i,1]=="TRUE")
  {
    v2[i]<-1
  }
  if(P2[i,1]=="FALSE")
  {
    v2[i]<-0
  }
}

V2<-as.data.frame(cbind(e$st,v2))
colnames(V2)<-c("yhat","m2")
write.table(V2,"C:/STAT/Wk-5-EC-Logistic/V2.csv", sep=",")
## CIs using profiled log-likelihood
confint(m2)
```

Waiting for profiling to be done...

```
##           2.5 %      97.5 %
## (Intercept) -1.097900e+00 -9.727387e-01
## e$sls       1.463126e-06  1.157731e-05
## e$eco       2.391078e-04  9.606299e-04
## e$qcec      -1.376291e-03 -6.050908e-04
## e$ad        5.571897e-02  6.795780e-02
```

Analysis of the Residual Deviance from the ANOVA summary ...Analysis of Deviance Table

```
anova(object=m1, test="Chisq")
```

Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred

Analysis of Deviance Table

##

Model: binomial, link: logit

##

Response: e\$st

##

Terms added sequentially (first to last)

##

##

		Df	Deviance	Resid. Df	Resid. Dev	Pr(>Chi)
##	NULL			9999	13842	
##	e\$sls	1	2190.16	9998	11652	< 2.2e-16 ***
##	e\$eco	1	405.70	9997	11246	< 2.2e-16 ***
##	e\$eco	1	4.77	9996	11242	0.02902 *
##	e\$csec	1	24.05	9995	11218	9.386e-07 ***
##	e\$osec	1	4.01	9994	11214	0.04510 *
##	e\$qcec	1	101.70	9993	11112	< 2.2e-16 ***
##	e\$spec	1	0.35	9992	11112	0.55221
##	e\$ad	1	322.37	9991	10789	< 2.2e-16 ***


```
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

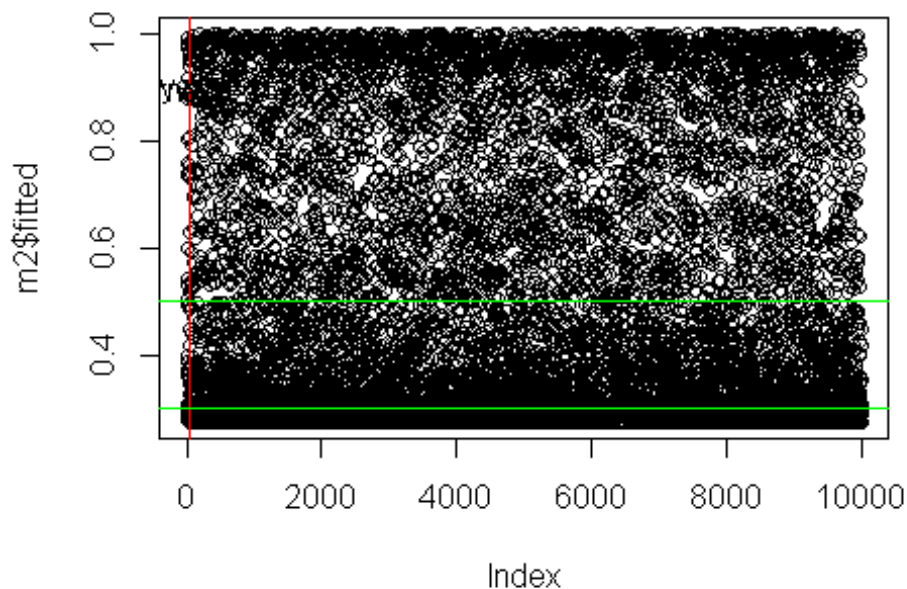
# Null- Resid deviance ==13842 at Df==9999
# e$ad- Resid deviance ==10789 at Df==9991
# The deviance was reduced by points on 8 degrees of freedom, for a p-value
of...
x<-13842-10789
x

## [1] 3053

1 - pchisq(3053, df=8)

## [1] 0

plot(m2$fitted)
abline(v=30.5,col="red")
abline(h=.3,col="green")
abline(h=.5,col="green")
text(15,.9,"Churned = 0")
text(40,.9,"Stayed = 1")
```



```
# Hosmer - The Hosmer-Lemeshow test...
library("ResourceSelection", lib.loc="~/R/win-library/3.1")

## ResourceSelection 0.2-4 2014-05-19

hl <- hoslem.test(m1$y, fitted(m1), g=10)
hl

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

```
##
## data: m1$y, fitted(m1)
## X-squared = 66.6066, df = 8, p-value = 2.32e-11

cbind(hl$observed,hl$expected)

##          y0 y1  yhat0  yhat1
## [0.255,0.264] 793 207 738.41580 261.5842
## (0.264,0.279] 754 246 727.79106 272.2089
## (0.279,0.305] 729 271 708.95382 291.0462
## (0.305,0.344] 682 318 677.52100 322.4790
## (0.344,0.417] 554 446 623.44816 376.5518
## (0.417,0.525] 470 530 532.75843 467.2416
## (0.525,0.679] 380 620 397.52050 602.4795
## (0.679,0.844] 276 724 239.10715 760.8929
## (0.844,0.943] 106 894 99.11548 900.8845
## (0.943,1]    29 971 28.36861 971.6314

pihat1 <- m1$fitted
pihat1cat <- cut(pihat1, breaks=c(0,quantile(pihat1, probs=seq(0.1,0.9,0.1)),1),
labels=FALSE)

m3<-glm(e$st~e$sls*e$sco+e$eco*e$csec+e$osec*e$qcec+e$spec*e$ad, data
=e, family = "binomial")
summary(m3)

##
## Call:
## glm(formula = e$st ~ e$sls * e$sco + e$eco * e$csec + e$osec *
## e$qcec + e$spec * e$ad, family = "binomial", data = e)
##
## Deviance Residuals:
##   Min       1Q   Median       3Q      Max
## -3.3791 -0.8512  0.1592  0.9319  2.3418
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  -1.099e+00  3.449e-02 -31.870 < 2e-16 ***
## e$sls         1.752e-05  3.374e-06   5.192 2.08e-07 ***
## e$sco         5.631e-04  2.055e-03   0.274 0.78406
## e$eco         1.553e-03  2.487e-04   6.243 4.28e-10 ***
## e$csec        -1.662e-02  1.006e-02  -1.651 0.09866 .
## e$osec         9.403e-03  1.015e-02   0.927 0.35402
## e$qcec        -2.268e-03  3.075e-04  -7.376 1.63e-13 ***
## e$spec        -2.338e-02  7.698e-03  -3.037 0.00239 **
## e$ad          6.722e-02  4.299e-03  15.638 < 2e-16 ***
## e$sls:e$sco   -1.633e-08  9.037e-09  -1.807 0.07079 .
## e$eco:e$csec   7.816e-08  2.485e-07   0.314 0.75315
## e$osec:e$qcec -1.949e-08  3.640e-07  -0.054 0.95729
## e$spec:e$ad    6.444e-04  2.618e-04   2.461 0.01386 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
```

```
##
## Null deviance: 13842 on 9999 degrees of freedom
## Residual deviance: 10777 on 9987 degrees of freedom
## AIC: 10803
##
## Number of Fisher Scoring iterations: 7

# Confusion Matrix created in Excel as seen below -
ModelFit_m3<-m3$fit
ProbabModelFit_m3<-m3$fit>0.5
P3<-as.data.frame(ProbabModelFit_m3)
# "if" loop to convert TRUE and FALSE into 1 and 0.

v3<-vector()
for (i in 1:nrow(P3))
{
  if(P3[i,1]=="TRUE")
  {
    v3[i]<-1
  }
  if(P3[i,1]=="FALSE")
  {
    v3[i]<-0
  }
}

V3<-as.data.frame(cbind(e$st,v3))
colnames(V3)<-c("yhat","m2")
# write.table(V3,"C:/STAT/Wk-5-EC-Logistic/V3.csv", sep=",")

m4<-glm(e$st~e$sls+e$sco*e$eco*e$csec+e$qcec*e$spec*e$ad, data =e,
family = "binomial")

## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred

summary(m4)

##
## Call:
## glm(formula = e$st ~ e$sls + e$sco * e$eco * e$csec + e$qcec *
## e$spec * e$ad, family = "binomial", data = e)
##
## Deviance Residuals:
## Min 1Q Median 3Q Max
## -4.3529 -0.8376 0.1294 0.9022 2.7156
##
## Coefficients:
## Estimate Std. Error z value Pr(>|z|)
## (Intercept) -1.160e+00 3.632e-02 -31.933 < 2e-16 ***
## e$sls 1.468e-05 2.798e-06 5.245 1.56e-07 ***
## e$sco 2.071e-03 2.612e-03 0.793 0.427819
## e$eco 2.078e-03 2.673e-04 7.772 7.71e-15 ***
## e$csec -4.906e-03 1.708e-03 -2.873 0.004067 **
## e$qcec -1.874e-03 4.361e-04 -4.298 1.73e-05 ***
```

```

## e$spec      -3.242e-02  9.544e-03  -3.397 0.000682 ***
## e$ad        5.823e-02  5.445e-03  10.693 < 2e-16 ***
## e$sco:e$eco -5.575e-06  1.414e-06  -3.943 8.05e-05 ***
## e$sco:e$csec 2.554e-06  6.066e-06   0.421 0.673768
## e$eco:e$csec -2.172e-06  4.496e-07  -4.830 1.36e-06 ***
## e$qcec:e$spec 8.738e-06  1.579e-05   0.553 0.579968
## e$qcec:e$ad  1.018e-05  1.125e-05   0.905 0.365387
## e$spec:e$ad  8.784e-04  3.314e-04   2.650 0.008040 **
## e$sco:e$eco:e$csec 7.860e-09 1.565e-09  5.021 5.14e-07 ***
## e$qcec:e$spec:e$ad 1.555e-07 4.964e-07  0.313 0.754070
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##    Null deviance: 13842  on 9999  degrees of freedom
## Residual deviance: 10729  on 9984  degrees of freedom

## AIC: 10761
##
## Number of Fisher Scoring iterations: 10

# Model - glm(m3) - AIC: 10803
# Model - glm(m4) -AIC: 10761
ModelFit_m4<-m4$fit
ProbabModelFit_m4<-m4$fit>0.5
P4<-as.data.frame(ProbabModelFit_m4)
# "if" loop to convert TRUE and FALSE into 1 and 0.

v4<-vector()
for (i in 1:nrow(P4))
{
  if(P4[i,1]=="TRUE")
  {
    v4[i]<-1
  }
  if(P4[i,1]=="FALSE")
  {
    v4[i]<-0
  }
}

V4<-as.data.frame(cbind(e$st,v4))
colnames(V4)<-c("yhat","m4")
# write.table(V4,"C:/STAT/V4.csv", sep=",")

# Analysis of the Residual Deviance from the ANOVA summary ...Analysis of
Deviance Table
anova(object=m2, test="Chisq")

```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Analysis of Deviance Table
##
## Model: binomial, link: logit
##
## Response: e$st
##
## Terms added sequentially (first to last)
##
##
##      Df Deviance Resid. Df Resid. Dev Pr(>Chi)
## NULL                9999    13842
## e$sls  1 2190.16    9998    11652 < 2.2e-16 ***
## e$eco  1  129.00    9997    11523 < 2.2e-16 ***
## e$qcec 1  236.79    9996    11286 < 2.2e-16 ***
## e$ad   1  423.51    9995    10863 < 2.2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
# Null- Resid deviance ==13842 at Df==9999
# e$ad- Resid deviance ==10863 at Df==9995
```

```
anova(object=m3, test="Chisq")
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Analysis of Deviance Table
##
## Model: binomial, link: logit
##
## Response: e$st
##
## Terms added sequentially (first to last)
##
##
##      Df Deviance Resid. Df Resid. Dev Pr(>Chi)
## NULL                9999    13842
## e$sls    1 2190.16    9998    11652 < 2.2e-16 ***
## e$sco    1  405.70    9997    11246 < 2.2e-16 ***
## e$eco    1    4.77    9996    11242 0.029025 *
## e$csec   1  24.05    9995    11218 9.386e-07 ***
## e$osec   1   4.01    9994    11214 0.045101 *
## e$qcec   1 101.70    9993    11112 < 2.2e-16 ***
## e$spec   1   0.35    9992    11112 0.552213
## e$ad     1  322.37    9991    10789 < 2.2e-16 ***
## e$sls:e$sco 1   4.80    9990    10784 0.028393 *
## e$eco:e$csec 1   0.11    9989    10784 0.744254
## e$osec:e$qcec 1   0.03    9988    10784 0.862922
## e$spec:e$ad 1   7.54    9987    10777 0.006033 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
# Null- Resid deviance ==13842 at Df==9999
# e$spec:e$ad- Resid deviance ==10777 at Df==9987
```

```
anova(object=m4, test="Chisq")
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
## Analysis of Deviance Table
```

```
##
```

```
## Model: binomial, link: logit
```

```
##
```

```
## Response: e$st
```

```
##
```

```
## Terms added sequentially (first to last)
```

```
##
```

```
##
```

```
##          Df Deviance Resid. Df Resid. Dev Pr(>Chi)
```

```
## NULL          9999    13842
```

```
## e$sls          1 2190.16    9998    11652 < 2.2e-16 ***
```

```
## e$sco          1  405.70    9997    11246 < 2.2e-16 ***
```

```
## e$eco          1   4.77    9996    11242  0.02902 *
```

```
## e$csec         1  24.05    9995    11218 9.386e-07 ***
```

```
## e$qcec         1 101.15    9994    11116 < 2.2e-16 ***
```

```
## e$spec         1   0.30    9993    11116  0.58452
```

```
## e$ad           1 326.30    9992    10790 < 2.2e-16 ***
```

```
## e$sco:e$eco     1   0.90    9991    10789  0.34220
```

```
## e$sco:e$csec    1   0.80    9990    10788  0.37057
```

```
## e$eco:e$csec    1   1.37    9989    10787  0.24186
```

```
## e$qcec:e$spec   1   3.83    9988    10783  0.05043 .
```

```
## e$qcec:e$ad     1   1.29    9987    10782  0.25608
```

```
## e$spec:e$ad     1   4.78    9986    10777  0.02884 *
```

```
## e$sco:e$eco:e$csec 1  47.76    9985    10729 4.828e-12 ***
```

```
## e$qcec:e$spec:e$ad 1   0.10    9984    10729  0.75348
```

```
## ---
```

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

```
# Null- Resid deviance ==13842 at Df==9999
```

```
# e$qcec:e$spec:e$ad- Resid deviance ==10729 at Df==9984
```

```
#####
```

```
# Tried to Combine Binomial variables of EComm data and glm()- create
Confusion Matrix within R. Could not be done.
```

```

# install.packages('caret', dependencies = TRUE)
## The R inbuilt caret Package - Confusion Matrix is not used as the code is not
clear...
# library("caret", lib.loc=~R/win-library/3.1")
# lvs <- c("Stayed", "Churned")
# truth <- factor(
#   c(
#     rep(lvs, times = c(4199,5801))),
#     levels = rev(lvs))
# pred <- factor(
#   c(
#     rep(lvs, times = c(5227,4773))),
#     levels = rev(lvs))
#
# xtab <- table(pred, truth)
# xtab
# confusionMatrix(xtab)
# confusionMatrix(pred, truth)
# confusionMatrix(xtab, prevalence = 0.25)
#
#
#
# lvs <- c("normal", "abnormal")
# truth <- factor(rep(lvs, times = c(86, 258)),
#   levels = rev(lvs))
# pred <- factor(
#   c(
#     rep(lvs, times = c(54, 32)),
#     rep(lvs, times = c(27, 231))),
#     levels = rev(lvs))
#
# xtab <- table(pred, truth)
#
# confusionMatrix(xtab)
# confusionMatrix(pred, truth)
# confusionMatrix(xtab, prevalence = 0.25)

```