# Churn Data Modeling Model Accuracy

Lecture 8 (Practical) 07/08/2021

# Model Diagonostics; overall model significance

```
> #Model Diagonostics;overall model significance
> with(lr5,pchisq(null.deviance-deviance,df.null-df.residual, lower.tail = F))
[1] 5.069035e-103
```

• The p-value is small, hence the overall model is significant at 0.01.

```
- - 1-pchisq(2056.9 - 1560.6, df=(2498-2491))
[1] 0
```

• The p-value associated with chi-square test for deviance also shows that the model is significant at 0.01

# The analysis of deviance is given below.

```
> anova(1r5, test='Chisq')
Analysis of Deviance Table
Model: binomial, link: logit
Response: training$Churn
Terms added sequentially (first to last)
                    Df Deviance Resid. Df Resid. Dev Pr(>Chi)
                                     2498
                                              2056.9
NULL
training$CSC_Hi
                                              1889.4 < 2.2e-16 ***
                     1 167.504
                                     2497
training$Int.l..Plan 1 127.410
                                              1762.0 < 2.2e-16 ***
                                     2496
                                              1726.9 3.191e-09 ***
training$VMP.ind
                     1 35.063
                                     2495
                                              1609.4 < 2.2e-16 ***
training$Day.Mins
                     1 117.573
                                     2494
training$Eve.Mins
                     1 25.484
                                     2493
                                              1583.9 4.460e-07 ***
                                              1572.3 0.0006771 ***
training$Night.Mins
                    1 11.551
                                     2492
                                              1560.6 0.0006289 ***
training$Intl.Mins
                     1 11.689
                                     2491
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

All variables are significant at 0.01

## Prediction using the fitted model

- The first five observations in the training data gives the probabilities as given above. (View Training data for more details of row numbers)
- As the first observation in the training data(row 1345) is predicted with low probability of churning (considering cut-off probability 0.5), the predicted class is non-churner. But view the data, row 1345 customer is a churner. So there is a mis-classification.
- Let us find the mis-classification error now.

#### Mis-classification error

The sum table will give more details

```
sumtable5<-addmargins(tab5,FUN=sum)
sumtable5
actual
predicted False. True. sum
0 2047 294 2341
1 93 65 158
sum 2140 359 2499
```

#### Mis-classification error

```
actual
predicted False. True. sum
0 2047 294 2341
1 93 65 158
sum 2140 359 2499
```

- It says that out of total 2499 customers, the training data has 359 churners, but the model predicted only 158 as churners. Also the training data has 2140 non-churners, but the model predicted as 2341 non-churners.
- That is the model incorrectly classified some actual churners as nonchurners.
- To check the accuracy of the model, we can find out sensitivity, specificity etc.

# Sensitivity

```
#TAN : Total Actually Negative= TN + FP
 #TAP : Total Actually Positive= FN + TP
 TAP <- sum(tab5[,2])
 TAN \leftarrow sum(tab5[,1])
 TP \leftarrow tab5[2,2]
 TN < -tab5[1,1]
 FP < -tab5[2,1]
 FN < -tab5[1,2]
TPR <- TP / TAP
TPR
[1] 0.1810585
```

• The TPR is sensitivity which is 0.18 only. Also we have already seen that the model classified some actual churners as non-churners.

# Specificity

```
FPR <- FP / TAN
FPR

[1] 0.04345794

specificity <- TN/TAN
specificity

[1] 0.9565421</pre>
```

The false negative rate is 0.04 and accordingly specificity is 0.95.
 Which means the model predicted actual non-churners as non-churners.

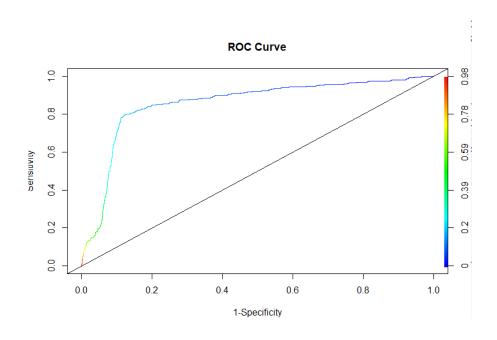
### Mis-classification error

```
#The proportion of observations correctly classified is
sum(diag(tab5))/sum(tab5)
[1] 0.8451381
```

- The correct classification rate is 0.845 which is more than 0.8 and looks fine to implement the model in practice. Accordingly the misclassification rate is
- #The proportion of observations misclassified is 1-sum(diag(tab5))/sum(tab5)
   [1] 0.1548619
- 15% is the mis-classification error.

#### **ROC Curve**

```
ROCRpred <- prediction(p, training[,21])
ROCRperf <- performance(ROCRpred, 'tpr','fpr')
ROCRperf
plot(ROCRperf, colorize=T, main= "ROC Curve", ylab = "Sensitivity", xlab = "1-Specificity")
abline(a=0,b=1)</pre>
```



• The curve is almost closer to left-hand border and top border, so the model accuracy is good.

#### Area under the curve

```
auc <- performance(ROCRpred, measure = "auc")
auc <- auc@y.values[[1]]
auc
[1] 0.8566045</pre>
```

• The area under the curve is 0.8566 which is more than 0.8 and can be considered as a good model and can be applied in practical situation.