

Logistic Regression Analysis - Medical Appointments

Prediction: No Show Appointments

Participants: Sarah Fite, Matthew Przybyla, and David Tran

Summary and Statistics

```
head(Data_Sub)
```

```
##   Gender Age Scholarship Hypertension Diabetes Alcoholism Handicap
## 1      F  62           0             1         0           0         0
## 2      M  56           0             0         0           0         0
## 3      F  62           0             0         0           0         0
## 4      F   8           0             0         0           0         0
## 5      F  56           0             1         1           0         0
## 6      F  76           0             1         0           0         0
##   SMS_received NOSHOW
## 1              0     No
## 2              0     No
## 3              0     No
## 4              0     No
## 5              0     No
## 6              0     No
```

```
str(Data_Sub)
```

```
## 'data.frame':   110527 obs. of  9 variables:
##  $ Gender      : Factor w/ 2 levels "F","M": 1 2 1 1 1 1 1 1 1 ...
##  $ Age         : int   62 56 62 8 56 76 23 39 21 19 ...
##  $ Scholarship : int   0 0 0 0 0 0 0 0 0 0 ...
##  $ Hypertension: int   1 0 0 0 1 1 0 0 0 0 ...
##  $ Diabetes    : int   0 0 0 0 1 0 0 0 0 0 ...
##  $ Alcoholism  : int   0 0 0 0 0 0 0 0 0 0 ...
##  $ Handicap    : int   0 0 0 0 0 0 0 0 0 0 ...
##  $ SMS_received: int   0 0 0 0 0 0 0 0 0 0 ...
##  $ NOSHOW      : Factor w/ 2 levels "No","Yes": 1 1 1 1 1 1 2 2 1 1 ...
```

```
sapply(Data_Sub, sd)
```

```
## Warning in var(if (is.vector(x) || is.factor(x)) x else as.double(x), na.rm = na.rm): Calling
var(x) on a factor x is deprecated and will become an error.
## Use something like 'all(duplicated(x)[-1L])' to test for a constant vector.

## Warning in var(if (is.vector(x) || is.factor(x)) x else as.double(x), na.rm = na.rm): Calling
var(x) on a factor x is deprecated and will become an error.
## Use something like 'all(duplicated(x)[-1L])' to test for a constant vector.
```

```
##      Gender      Age Scholarship Hypertension      Diabetes
## 0.4769790 23.1101759 0.2976748 0.3979213 0.2582651
## Alcoholism Handicap SMS_received      NOSHOW
## 0.1716856 0.1615427 0.4668727 0.4014440
```

```
xtabs(~NOSHOW + Age, data = Data_Sub)
```

```
##      Age
## NOSHOW  0   1   2   3   4   5   6   7   8   9  10  11  12
## No 2900 1859 1366 1236 1017 1169 1205 1126 1106 1008 970 948 820
## Yes 639 415 252 277 282 320 316 301 318 364 304 247 272
##      Age
## NOSHOW 13  14  15  16  17  18  19  20  21  22  23  24  25
## No 800 802 889 1049 1113 1137 1151 1082 1097 1025 1006 921 980
## Yes 303 316 322 353 396 350 394 355 355 351 343 321 352
##      Age
## NOSHOW 26  27  28  29  30  31  32  33  34  35  36  37  38
## No 971 1048 1116 1073 1152 1119 1174 1176 1204 1089 1236 1216 1309
## Yes 312 329 332 330 369 320 331 348 322 289 344 317 320
##      Age
## NOSHOW 39  40  41  42  43  44  45  46  47  48  49  50  51
## No 1196 1101 1038 1007 1035 1164 1198 1177 1127 1128 1354 1322 1284
## Yes 340 301 308 265 309 323 255 283 267 271 298 291 283
##      Age
## NOSHOW 52  53  54  55  56  57  58  59  60  61  62  63  64
## No 1449 1332 1262 1168 1372 1325 1216 1357 1175 1143 1100 1195 1149
## Yes 297 319 268 257 263 278 253 267 236 200 212 179 182
##      Age
## NOSHOW 65  66  67  68  69  70  71  72  73  74  75  76  77
## No 934 1008 825 843 714 630 574 514 629 513 463 480 448
## Yes 167 179 148 169 118 94 121 101 96 89 81 91 79
##      Age
## NOSHOW 78  79  80  81  82  83  84  85  86  87  88  89  90
## No 452 329 430 371 326 219 276 226 218 157 114 144 86
## Yes 89 61 81 63 66 61 35 49 42 27 12 29 23
##      Age
## NOSHOW 91  92  93  94  95  96  97  98  99 100 102 115
## No 53 66 43 27 18 16 9 5 1 4 2 2
## Yes 13 20 10 6 6 1 2 1 0 0 0 3
```

Split dataset into “Train” (80%) and “Test” (20%)

```
Split <- sample(2, nrow(Data_Sub), replace=TRUE, prob = c(0.8, 0.2))
Train <- Data_Sub[Split==1,]
Test <- Data_Sub[Split==2,]
```

Fitting the Model

```
model <- glm(NOSHOW ~., family=binomial(link='logit'), data=Train)
model2 <- glm(NOSHOW ~ Age + Scholarship + Hypertension + Diabetes + Alcoholism + SMS_received,
              family=binomial(link='logit'), data=Train)
predict <- predict(model, type = 'response')

summary(model)
```

```
##
## Call:
## glm(formula = NOSHOW ~ ., family = binomial(link = "logit"),
##      data = Train)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -0.9538  -0.6843  -0.6083  -0.5372   2.1124
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  -1.3831649   0.0196638  -70.341 < 2e-16 ***
## GenderM      -0.0142421   0.0182145   -0.782  0.43427
## Age          -0.0065303   0.0004391  -14.872 < 2e-16 ***
## Scholarship   0.1954285   0.0273091   7.156  8.3e-13 ***
## Hypertension -0.0617751   0.0274333   -2.252  0.02433 *
## Diabetes      0.0987910   0.0380787   2.594  0.00948 **
## Alcoholism    0.1280998   0.0495697   2.584  0.00976 **
## Handicap      0.0166565   0.0545769    0.305  0.76022
## SMS_received  0.6360021   0.0173020  36.759 < 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: 88946  on 88363  degrees of freedom
## Residual deviance: 87232  on 88355  degrees of freedom
## AIC: 87250
##
## Number of Fisher Scoring iterations: 4
```

```
summary(model2)
```

```
##
## Call:
## glm(formula = NOSHOW ~ Age + Scholarship + Hypertension + Diabetes +
##       Alcoholism + SMS_received, family = binomial(link = "logit"),
##       data = Train)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -0.9525  -0.6838  -0.6072  -0.5372   2.1200
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  -1.3898475   0.0176308  -78.830   < 2e-16 ***
## Age           -0.0064831   0.0004353  -14.893   < 2e-16 ***
## Scholarship    0.1983339   0.0270540    7.331 2.28e-13 ***
## Hypertension  -0.0615190   0.0274156   -2.244  0.02484 *
## Diabetes       0.0988917   0.0380688    2.598  0.00938 **
## Alcoholism     0.1234178   0.0492134    2.508  0.01215 *
## SMS_received  0.6364802   0.0172780   36.838   < 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: 88946  on 88363  degrees of freedom
## Residual deviance: 87233  on 88357  degrees of freedom
## AIC: 87247
##
## Number of Fisher Scoring iterations: 4
```

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

```
## Analysis of Deviance Table
##
## Model: binomial, link: logit
##
## Response: NOSHOW
##
## Terms added sequentially (first to last)
##
##
```

	Df	Deviance	Resid. Df	Resid. Dev	Pr(>Chi)
## NULL			88363	88946	
## Gender	1	1.31	88362	88945	0.25265
## Age	1	324.73	88361	88620	< 2.2e-16 ***
## Scholarship	1	50.63	88360	88570	1.116e-12 ***
## Hypertension	1	3.61	88359	88566	0.05729 .
## Diabetes	1	3.92	88358	88562	0.04767 *
## Alcoholism	1	3.19	88357	88559	0.07391 .
## Handicap	1	0.36	88356	88558	0.54606
## SMS_received	1	1326.54	88355	87232	< 2.2e-16 ***

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

```
confint(model)
```

```
## Waiting for profiling to be done...
```

```
##
```

	2.5 %	97.5 %
## (Intercept)	-1.421772186	-1.344690711
## GenderM	-0.049976895	0.021423687
## Age	-0.007391725	-0.005670456
## Scholarship	0.141725615	0.248780368
## Hypertension	-0.115644517	-0.008104809
## Diabetes	0.023865006	0.173142209
## Alcoholism	0.030132402	0.224478586
## Handicap	-0.091898433	0.122139506
## SMS_received	0.602081037	0.669904593

Misclassification Rate

```
p <- predict(model2, Data_Sub)
table <- table(p, Data_Sub$Handicap)
Classification_Rate = sum(diag(table))/sum(table)
Classification_Rate
```

```
## [1] 9.047563e-06
```

```
Misclassification_Rate = 1- sum(diag(table))/sum(table)
Misclassification_Rate
```

```
## [1] 0.999991
```

Accessing the predicability of the model

```
fitted.results <- predict(model, newdata=subset(Test, select=c(1,2,3,4,5,6,7,8)), type='response')
fitted.results <- ifelse(fitted.results > 0.5, 1, 0)
misClasificError <- mean(fitted.results != Test$NOSHOW)
print(paste('Accuracy', 1-misClasificError))
```

```
## [1] "Accuracy 0"
```

Model Performance Evaluation

```
pred <- predict(model, Train, type= "response")
head(pred)
```

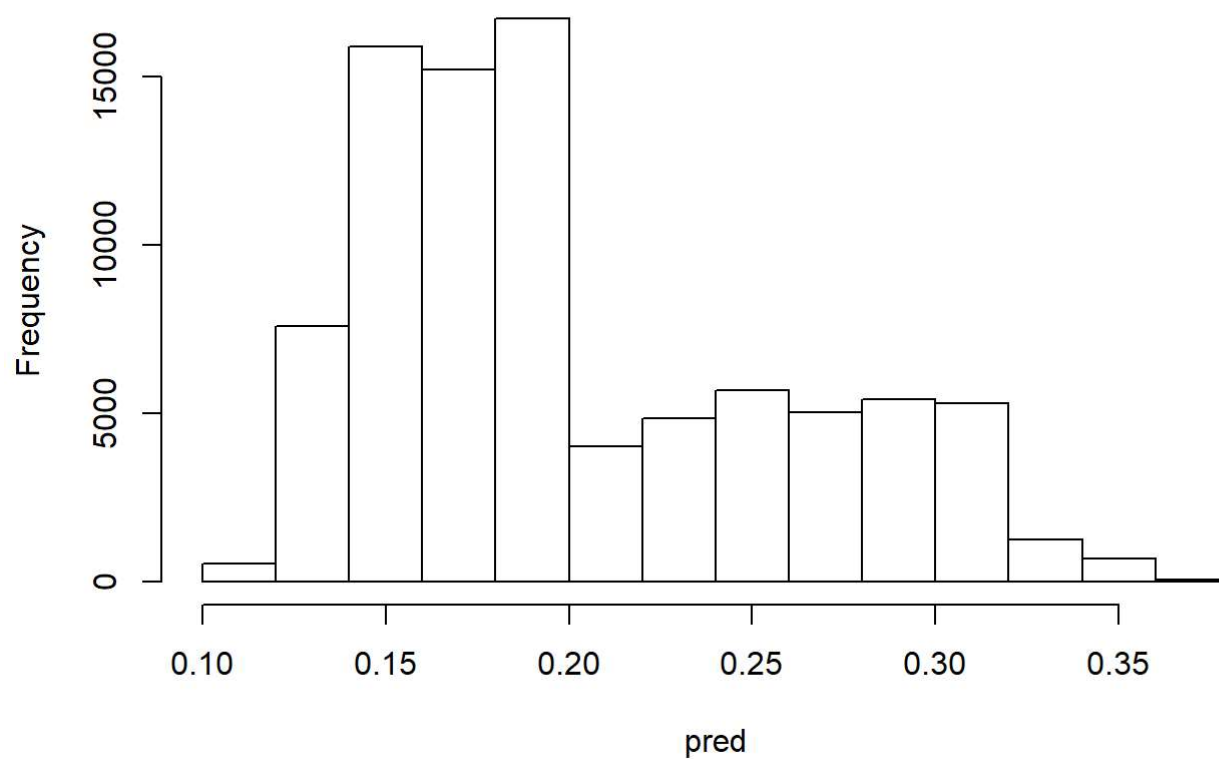
```
##           1           3           4           6           7           8
## 0.1358943 0.1433129 0.1922576 0.1255115 0.1775024 0.1627584
```

```
head(Train)
```

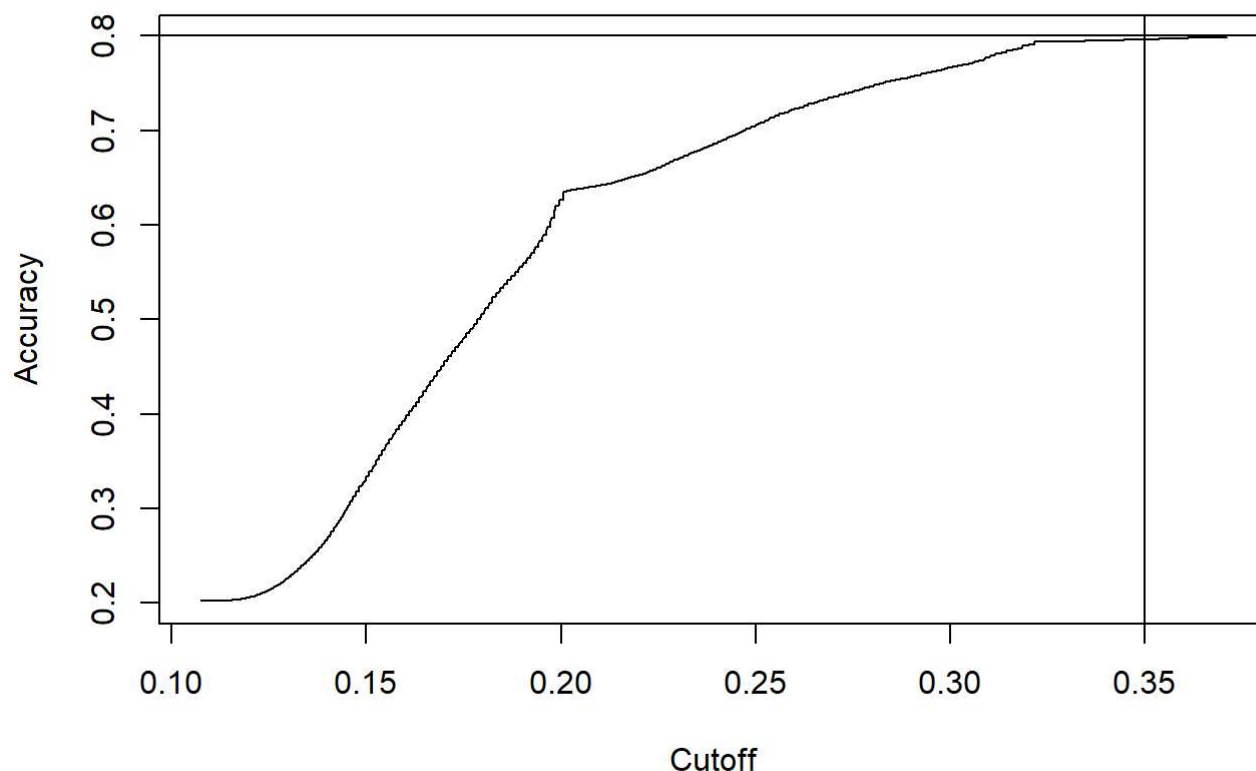
```
##   Gender Age Scholarship Hypertension Diabetes Alcoholism Handicap
## 1      F  62           0             1         0           0         0
## 3      F  62           0             0         0           0         0
## 4      F   8           0             0         0           0         0
## 6      F  76           0             1         0           0         0
## 7      F  23           0             0         0           0         0
## 8      F  39           0             0         0           0         0
##   SMS_received NOSHOW
## 1             0     No
## 3             0     No
## 4             0     No
## 6             0     No
## 7             0     Yes
## 8             0     Yes
```

```
hist(pred)
```

Histogram of pred



```
predf <- prediction(pred, Train$NOSHOW)
eval <- performance(predf, "acc")
plot(eval)
abline(h=0.80, v=0.35)
```



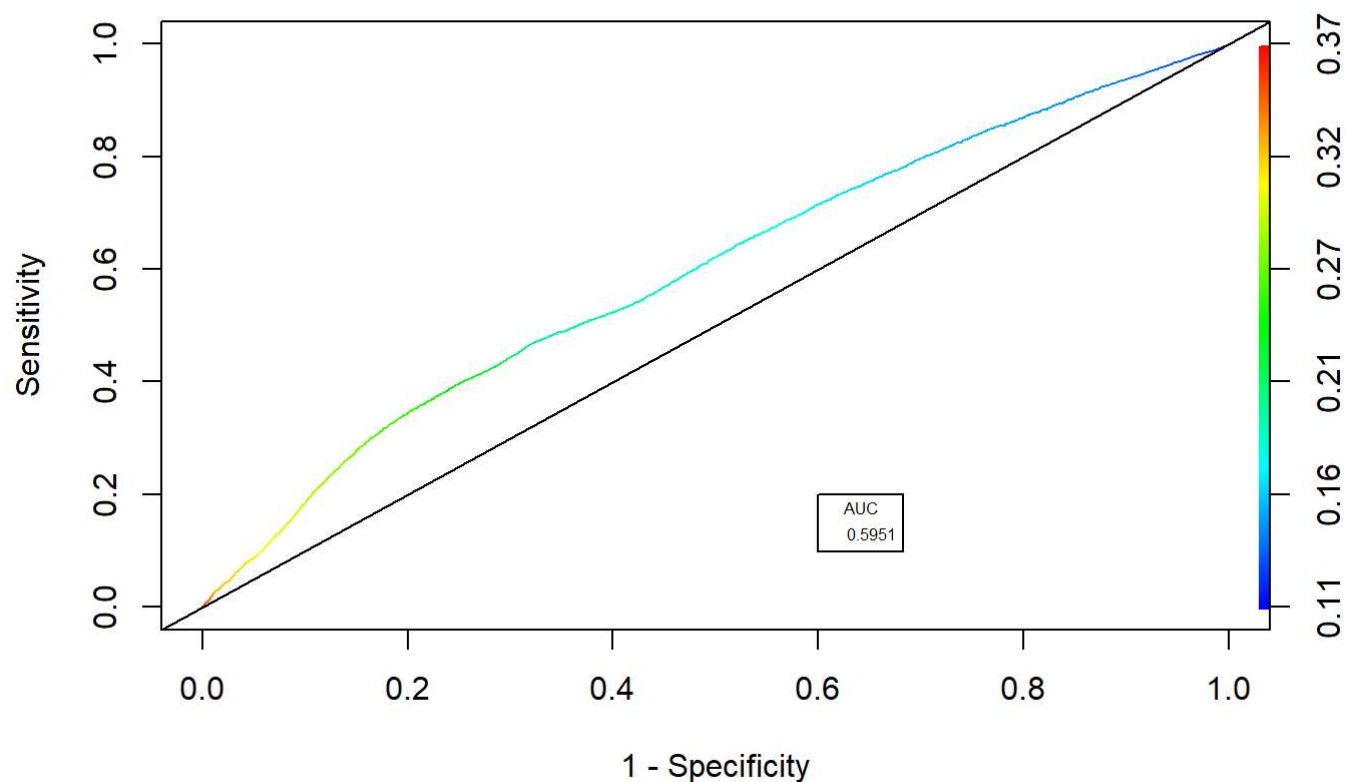
Receiver Operating Characteristic (ROC) Curve & Area Under Curve (AUC)

```

pred2 <- prediction(pred, Train$NOSHOW)
roc <- performance(pred2, "tpr", "fpr")
plot(roc,
      colorize=T,
      main = "ROC Curve",
      ylab = "Sensitivity",
      xlab = "1 - Specificity")
abline(a=0, b=1)
auc <- performance(pred2, "auc")
auc2 <- unlist(slot(auc, "y.values"))
auc <- round(auc2, 4)
legend(.6, .2, auc, title = "AUC", cex =.5)

```


ROC Curve



Identify Best Values

```
max <- which.max(slot(eval, "y.values")[[1]])
max
```

```
## [1] 3
```

```
acc <- slot(eval, "y.values")[[1]][max]
acc
```

```
## [1] 0.7979381
```

```
cut <- slot(eval, "x.values")[[1]][max]
cut
```

```
##      32330
## 0.3694332
```