Predicting Earnings Manipulation - Logistic Regression, Decision Trees and Random Forest

We will try to predict earnings manipulations by Indian firms using machine learning algorithms. The data comes from a case study by the Indian Institute of Management Bangalore. Entire dataset contains 1200 non-manipulators and 39 manipulators and sample dataset contains 220 cases including 39 manipulators.

```
#LOAD and fix STR of data
rm(list=ls())
library(readx1)
sample data <- read excel("IMB579-XLS-ENG.xlsx", sheet = "Sample for Model</pre>
Development")
#rename and convert to categorical variable (1 YES manipulator/0 NO not
manipulator)
sample data<- sample data[, -10] #remove manipulator yes/no column</pre>
names(sample data)[10] <- "c Manipulator"</pre>
C_Manipulator <- as.factor(sample_data$c_Manipulator)</pre>
sample data <- data.frame(sample data, C Manipulator)</pre>
sample data<- sample data[, -10] #remove c maniputator column that is not
categorical
sample data<- sample data[, -1] #remove company.ID column</pre>
str(sample_data)
## 'data.frame':
                    220 obs. of 9 variables:
## $ DSRI
                   : num 1.62 1 1 1.49 1 ...
## $ GMI
                  : num 1.13 1.61 1.02 1 1.37 ...
## $ AQI
                   : num 7.185 1.005 1.241 0.466 0.637 ...
## $ SGI
                   : num 0.366 13.081 1.475 0.673 0.861 ...
## $ DEPI
                  : num 1.38 0.4 1.17 2 1.45 ...
## $ SGAI
                   : num 1.6241 5.1982 0.6477 0.0929 1.7415 ...
## $ ACCR
                   : num -0.1668 0.0605 0.0367 0.2734 0.123 ...
## $ LEVI
                   : num 1.161 0.987 1.264 0.681 0.939 ...
## $ C_Manipulator: Factor w/ 2 levels "0", "1": 2 2 2 2 2 2 2 2 2 2 ...
```

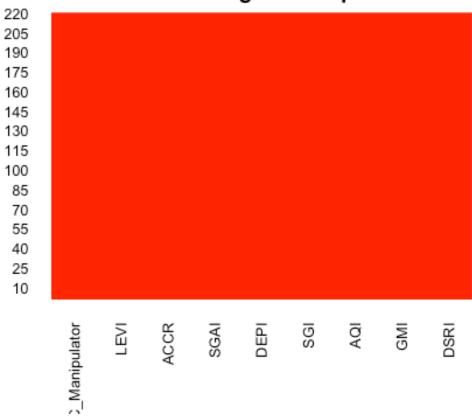
EDA

```
library(dplyr)
count(sample_data, vars= C_Manipulator)

## vars n
## 1 0 181
## 2 1 39
```

```
library(Amelia)
library(mlbench)
missmap(sample_data, col=c("blue", "red"), legend=FALSE)
```

Missingness Map

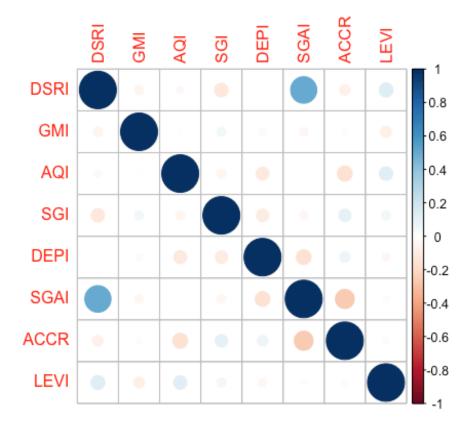


```
#no missing data in dataset

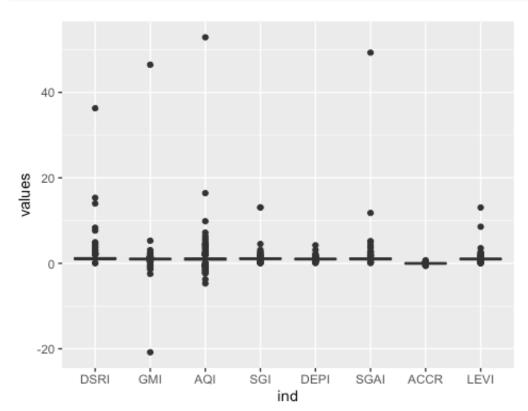
library(corrplot)

correlations <- cor(sample_data[,1:8])
 corrplot(correlations, method="circle")

library(ggplot2)</pre>
```



ggplot(stack(sample_data), aes(x = ind, y = values)) +
 geom_boxplot()



```
#function to calculate accuracy
my.accuracy <- function(actual, predictions)</pre>
  y<- as.vector(table(predictions, actual))</pre>
  names(y) <- c("TN", "FP", "FN", "TP")</pre>
  accuracy<- (y["TN"]+ y["TP"])/ sum(y)
  return(as.numeric(accuracy))
}
#function to calculate recall
my.recall <- function(actual, predictions)</pre>
  y<- as.vector(table(predictions, actual))</pre>
  names(y) <- c("TN", "FP", "FN", "TP")</pre>
  recall <- y["TP"]/ (y["TP"] + y["FN"])
  return(as.numeric(recall))
}
#function to calculate precision TP/TP+FP
my.precision <- function(actual, predictions)</pre>
  y<- as.vector(table(predictions, actual))</pre>
  names(y) <- c("TN", "FP", "FN", "TP")</pre>
  precision \leftarrow y["TP"]/ (y["TP"] + y["FP"])
  return(as.numeric(precision))
#function to calculate F-1 score 2*(Recall * Precision) / (Recall +
Precision)
my.f1 <- function(actual, predictions)</pre>
  y<- as.vector(table(predictions, actual))</pre>
  names(y) <- c("TN", "FP", "FN", "TP")</pre>
  f1 <- (2*((y["TP"]/ (y["TP"] + y["FN"]))* y["TP"]/ (y["TP"] + y["FP"]))) /
((y["TP"]/(y["TP"] + y["FN"]) + (y["TP"]/(y["TP"] + y["FP"]))))
  return(as.numeric(f1))
```

MODEL 1: LOGISTIC REGRESSION using sample data- NO BALANCING

```
#SPLIT SAMPLE DATA
set.seed(123)
indx<- sample(2, size=nrow(sample_data),replace= T, prob = c(0.7, 0.3))
train <- sample_data[indx==1,]
test <- sample_data[indx==2, ]
actual<- test$C_Manipulator

#TRAIN MODEL
logitModel <- glm(C_Manipulator ~ . , data = train, family = "binomial")
summary(logitModel)</pre>
```

```
##
## Call:
## glm(formula = C_Manipulator ~ ., family = "binomial", data = train)
## Deviance Residuals:
                                   3Q
##
       Min
                 10
                      Median
                                           Max
## -1.6588 -0.4022
                    -0.2812 -0.2009
                                        2,4824
##
## Coefficients:
##
               Estimate Std. Error z value Pr(>|z|)
                            2.0376 -4.096 4.2e-05 ***
## (Intercept) -8.3466
## DSRI
                 0.9096
                            0.3706
                                    2.455 0.014108 *
## GMI
                 1.3304
                            0.4578
                                     2.906 0.003659 **
## AQI
                 0.4764
                            0.1641 2.903 0.003693 **
                            0.6788
                                     3.315 0.000917 ***
## SGI
                 2.2498
## DEPI
                 0.2925
                            0.6000
                                     0.487 0.625909
## SGAI
                 0.1892
                            0.4570
                                     0.414 0.678823
                            2.1026 3.387 0.000705 ***
## ACCR
                 7.1227
                            0.7333 -0.177 0.859182
## LEVI
                -0.1301
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
##
       Null deviance: 138.688 on 159 degrees of freedom
## Residual deviance: 87.272 on 151 degrees of freedom
## AIC: 105.27
##
## Number of Fisher Scoring iterations: 8
#TEST MODEL
predictions <- predict(logitModel, newdata = test, type = "response")</pre>
actual <- test$C Manipulator
#confusion matrix
predict_logitmodel <- rep(0, length(predictions)) #all zeros for predictions</pre>
predict logitmodel[predictions >= 0.5 ] <- 1 #replace with 1 of prob >= 0.5
table(predict logitmodel,actual , dnn= c("Predictions", "Actual"))
#confusion matrix
##
              Actual
## Predictions
               0
##
            0 46 9
##
             1 0
#Evaluation of model
my.accuracy(actual, predict_logitmodel)
## [1] 0.85
my.recall(actual, predict_logitmodel)
## [1] 0.3571429
```

Balancing TRAINING data with 4 different techniques (over, under, both, rose)

```
library(ROSE)
set.seed(123)
#UNDERSAMPLED - uses all instances from minority training class(uses 25 out
of 25 observations in minority class) and ONLY the same num of instances for
majority class
sample data balanced under <- ovun.sample(C Manipulator ~ ., data = train,</pre>
method = "under")$data
table(sample data balanced under$C Manipulator)
## 0 1
## 25 25
#OVERSAMPLED - duplicates examples from the minority class
sample data balanced over <- ovun.sample(C Manipulator ~ ., data = train,</pre>
method = "over", N=500)$data
table(sample data balanced over$C Manipulator)
##
   0 1
## 135 365
#BOTH-the minority class is oversampled with replacement and majority class
is undersampled without replacement
sample data balanced both <- ovun.sample(C Manipulator ~ ., data = train,</pre>
method = "both", N=500)$data
table(sample data balanced both$C Manipulator)
## 0 1
## 244 256
#ROSE-generate data synthetically as well.
sample data balanced rose <- ROSE(C Manipulator ~ ., data = train)$data
table(sample data balanced rose$C Manipulator)
## 0 1
## 86 74
```

MODEL 2,3,4,5: LOGISTIC REGRESSION using different sample techniques

```
set.seed(123)
#***UNDERSAMPLIG****
sample_under_fit <- glm(C_Manipulator ~ . , data =</pre>
sample data balanced under, family = "binomial")
summary(sample_under_fit)
##
## Call:
## glm(formula = C_Manipulator ~ ., family = "binomial", data =
sample data balanced under)
##
## Deviance Residuals:
##
        Min
                   10
                         Median
                                       30
                                                Max
                                            1.74083
## -2.89210 -0.46380 -0.01759
                                  0.40329
##
## Coefficients:
               Estimate Std. Error z value Pr(>|z|)
##
## (Intercept) -14.7099
                            5.6874 -2.586
                                             0.0097 **
## DSRI
                            2.0445 2.320
                 4.7427
                                             0.0204 *
                                     1.504
## GMI
                 2.3973
                            1.5939
                                             0.1326
## AQI
                 1.8086
                            0.7859
                                   2.301
                                             0.0214 *
## SGI
                 6.8424
                            2.6860
                                    2.547
                                             0.0109 *
                -1.7355
## DEPI
                            1.2769 -1.359
                                             0.1741
## SGAI
                -0.6365
                            0.9269 -0.687
                                             0.4923
## ACCR
                10.2933
                            4.4202
                                     2.329
                                             0.0199 *
                -2.0795
## LEVI
                            1.7785 -1.169
                                             0.2423
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
##
       Null deviance: 69.315 on 49 degrees of freedom
## Residual deviance: 30.185 on 41 degrees of freedom
## AIC: 48.185
##
## Number of Fisher Scoring iterations: 9
sample_predictions_under <- predict(sample_under_fit, newdata = test, type =</pre>
"response")
#confusion matrix
predict_under_sample<- rep(0, length(sample predictions_under)) #all zeros</pre>
```

```
for predictions
predict under sample[sample predictions under >= 0.5 ] <- 1</pre>
                                                                   #replace
with 1 of prob >= 0.5
table(predict_under_sample,actual , dnn= c("Predictions", "Actual"))
              Actual
## Predictions 0 1
##
             0 37 3
##
             1 9 11
#***OVERSAMPLIG****
#train
sample_over_fit <- glm(C_Manipulator ~ . , data = sample_data_balanced_over,</pre>
family = "binomial")
summary(sample_over_fit)
##
## Call:
## glm(formula = C_Manipulator ~ ., family = "binomial", data =
sample_data_balanced_over)
##
## Deviance Residuals:
                      Median
                                   30
                                           Max
##
       Min
                 10
## -3.7250 -0.1922
                      0.2110
                               0.4881
                                        1.6669
## Coefficients:
                Estimate Std. Error z value Pr(>|z|)
##
                            1.39169 -7.291 3.08e-13 ***
## (Intercept) -10.14674
                                    5.758 8.53e-09 ***
## DSRI
                 2.02334
                            0.35142
## GMI
                 1.96464
                            0.42087
                                     4.668 3.04e-06 ***
                            0.12354
                                    6.934 4.08e-12 ***
## AQI
                 0.85664
## SGI
                 4.21524
                            0.59038
                                      7.140 9.34e-13 ***
## DEPI
                -0.05482
                            0.52391
                                    -0.105
                                               0.917
## SGAI
                            0.30412
                                      0.041
                 0.01251
                                               0.967
                                    7.029 2.09e-12 ***
## ACCR
                 9.68868
                            1.37848
## LEVI
                -0.23833
                            0.43402 -0.549
                                               0.583
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
       Null deviance: 583.26 on 499 degrees of freedom
## Residual deviance: 310.63 on 491 degrees of freedom
## AIC: 328.63
##
## Number of Fisher Scoring iterations: 9
#test
sample_predictions_over <- predict(sample_over_fit, newdata = test, type =</pre>
"response")
```

```
#confusion matrix
predict over sample<- rep(0, length(sample predictions over))</pre>
                                                                #all zeros
for predictions
predict over sample[sample predictions over >= 0.5 ] <- 1</pre>
                                                                #replace
with 1 of prob >= 0.5
table(predict_over_sample,actual , dnn= c("Predictions", "Actual"))
##
              Actual
## Predictions 0 1
             0 35 2
##
             1 11 12
##
#***BOTH***
#train
sample_both_fit <- glm(C_Manipulator ~ . , data = sample_data_balanced_both,</pre>
family = "binomial")
summary(sample_both_fit)
##
## Call:
## glm(formula = C Manipulator ~ ., family = "binomial", data =
sample_data_balanced_both)
##
## Deviance Residuals:
       Min
                 1Q
                      Median
                                   3Q
                                           Max
## -3.6787 -0.5121
                      0.0000
                               0.5631
                                        1.9443
##
## Coefficients:
                Estimate Std. Error z value Pr(>|z|)
## (Intercept) -14.67630
                            1.57498 -9.318 < 2e-16 ***
## DSRI
                            0.29525
                                    7.191 6.45e-13 ***
                 2.12304
## GMI
                 1.81684
                            0.29933
                                      6.070 1.28e-09 ***
                                    6.730 1.70e-11 ***
## AQI
                 0.65673
                            0.09758
                            0.55758
                                    8.325 < 2e-16 ***
## SGI
                 4.64189
                            0.75546
                                    4.663 3.11e-06 ***
## DEPI
                 3.52297
                            0.26861 -0.165
## SGAI
                -0.04422
                                               0.869
## ACCR
                8.40806
                            1.06452
                                      7.898 2.82e-15 ***
## LEVI
               -0.38377
                            0.41262 -0.930
                                               0.352
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
       Null deviance: 692.86 on 499 degrees of freedom
##
## Residual deviance: 366.42 on 491 degrees of freedom
## AIC: 384.42
## Number of Fisher Scoring iterations: 9
```

```
#test
sample predictions both <- predict(sample both fit, newdata = test, type =</pre>
"response")
#confusion matrix
predict both sample<- rep(0, length(sample predictions both))</pre>
                                                               #all zeros
for predictions
predict both sample[sample predictions both >= 0.5 ] <- 1</pre>
                                                               #replace
with 1 of prob >= 0.5
table(predict_both_sample,actual , dnn= c("Predictions", "Actual"))
##
             Actual
## Predictions 0 1
##
            0 43 3
##
            1 3 11
#***ROSE***
#train
sample rose fit <- glm(C_Manipulator ~ . , data = sample data_balanced_rose,
family = "binomial")
summary(sample_rose_fit)
##
## Call:
## glm(formula = C_Manipulator ~ ., family = "binomial", data =
sample data balanced rose)
## Deviance Residuals:
##
      Min
                1Q
                     Median
                                  3Q
                                          Max
## -1.9413 -0.9968 -0.7160
                              0.9469
                                       2.2675
##
## Coefficients:
               Estimate Std. Error z value Pr(>|z|)
##
## (Intercept) -1.48760
                          0.71124 -2.092 0.03648 *
                                    1.394 0.16346
## DSRI
               0.04479
                          0.03214
## GMI
               0.68807
                          0.28385 2.424 0.01535 *
                                    2.103 0.03543 *
## AQI
               0.05178
                          0.02462
                                    0.471 0.63735
## SGI
               0.05354
                          0.11357
## DEPI
                                    0.248 0.80418
               0.12462
                          0.50263
## SGAI
               0.07291
                          0.03518
                                    2.072 0.03822 *
## ACCR
               3.07205
                          0.95629
                                    3.212 0.00132 **
## LEVI
               0.10692
                          0.15213
                                    0.703 0.48217
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
##
       Null deviance: 220.91 on 159 degrees of freedom
## Residual deviance: 194.05 on 151 degrees of freedom
## AIC: 212.05
```

```
##
## Number of Fisher Scoring iterations: 4
#test
sample predictions rose <- predict(sample rose fit, newdata = test, type =</pre>
"response")
#confusion matrix
predict rose sample<- rep(0, length(sample predictions rose)) #all zeros</pre>
for predictions
predict_rose_sample[sample_predictions_rose >= 0.5 ] <- 1</pre>
                                                                 #replace
with 1 of prob >= 0.5
table(predict_rose_sample,actual , dnn= c("Predictions", "Actual"))
##
              Actual
## Predictions 0 1
##
             0 42 6
##
             1 4 8
```

EVALUATION OF LOGISTIC REGRESSION MODELS using balanced data

```
print("Accuracy/Recall/Precision/F-1 score using UNDER SAMPLING")
my.accuracy(actual, predict_under_sample)
## [1] 0.8
my.recall(actual, predict_under_sample)
## [1] 0.7857143
my.precision(actual, predict_under_sample)
## [1] 0.55
my.f1(actual, predict_under_sample)
## [1] 0.6470588
print("Accuracy/Recall/Precision/F-1 score using OVER SAMPLING")
my.accuracy(actual, predict_over_sample)
## [1] 0.7833333
my.recall(actual, predict over sample)
## [1] 0.8571429
my.precision(actual, predict_over_sample)
## [1] 0.5217391
```

```
my.f1(actual, predict over sample)
## [1] 0.6486486
print("Accuracy/Recall/Precision/F-1 score using BOTH UNDER AND OVER
SAMPLING")
my.accuracy(actual, predict_both_sample)
## [1] 0.9
my.recall(actual, predict_both_sample)
## [1] 0.7857143
my.precision(actual, predict_both_sample)
## [1] 0.7857143
my.f1(actual, predict both sample)
## [1] 0.7857143
print("Accuracy/Recall/Precision/F-1 score using ROSE")
my.accuracy(actual, predict_rose_sample)
## [1] 0.8333333
my.recall(actual, predict_rose_sample)
## [1] 0.5714286
my.precision(actual, predict rose sample)
## [1] 0.6666667
my.f1(actual, predict_rose_sample)
## [1] 0.6153846
```

	UNDERSAMPLING	OVERSAMPLING	ВОТН	ROSE
Accuracy	0.8	0.7833333	0.9	0.8333333
Recall	0.7857143	0.8571429	0.7857143	0.5714286
Precision	0.55	0.5217391	0.7857143	0.6666667
F-1 Score	0.6470588	0.6486486	0.7857143	0.6153846

#BEST F-1 SCORE WAS OBTAINED BY BALANCING DATA WITH BOTH UNDER AND OVER SAMPLING.

#F-1 SCORE INCREASED WITH ALL BALANCING TECHNIQUES COMPARED TO THE ORIGINAL SAMPLE DATA. (0.5263158)

LOGISTIC REGRESSION MODELS using entire dataset

```
set.seed(123)
data <- read_excel("IMB579-XLS-ENG.xlsx", sheet = "Complete Data")
data<- data[, -10] #remove manipulator yes/no column
names(data)[10] <- "c_Manipulator"
C_Manipulator <- as.factor(data$c_Manipulator)
data <- data.frame(data, C_Manipulator)
data<- data[, -10] #remove c_maniputalor column that is not categorical
data<- data[, -1] #remove company.ID column</pre>
```

Create balanced datasets(4)

```
#UNDERSAMPLED
down data <- ovun.sample(C Manipulator ~ ., data = data, method =</pre>
"under")$data
table(down data$C Manipulator)
## 0 1
## 38 39
#OVERSAMPLED
up data <- ovun.sample(C Manipulator ~ ., data = data, method = "over",
N=2400)$data
table(up_data$C_Manipulator)
##
      0
## 1200 1200
#BOTH
both_data <- ovun.sample(C_Manipulator ~ ., data = data, method = "both",
p=0.5, N=1250)$data
table(both_data$C_Manipulator)
## 0 1
## 622 628
#ROSE-generate data synthetically as well.
rose data <- ROSE(C Manipulator ~ ., data = data)$data
table(rose data$C Manipulator)
## 0 1
## 634 605
```

#LOGISTIC REGRESSION MODEL USING NO BALANCING AND 4 BALANCING TECHNIQUES(under, over, both, rose)

MODELS using entire dataset (Unbalanced data, under, over, both, and rose sampled)

```
#***NOT BALANCED****
indx<- sample(2, size=nrow(data),replace= T, prob = c(0.7, 0.3))
train1 <- data[indx==1,]</pre>
```

```
test1 <- data[indx==2, ]
actual2<- test1$C Manipulator</pre>
#train
logitModel.fit <- glm(C_Manipulator ~ . , data = train1, family = "binomial")</pre>
summary(logitModel.fit)
## Call:
## glm(formula = C Manipulator ~ ., family = "binomial", data = train1)
## Deviance Residuals:
##
        Min
                  10
                                      3Q
                                               Max
                        Median
## -3.10782 -0.17717 -0.12640 -0.09576
                                           3.05648
##
## Coefficients:
##
               Estimate Std. Error z value Pr(>|z|)
## (Intercept) -6.19075
                          1.51533 -4.085 4.40e-05 ***
                          0.16662 3.874 0.000107 ***
## DSRI
               0.64549
## GMI
                                    2.550 0.010762 *
               0.78188
                          0.30658
## AQI
               0.27769
                          0.09708 2.861 0.004229 **
## SGI
               1.18102
                          0.30660 3.852 0.000117 ***
## DEPI
              -1.31165
                        1.15049 -1.140 0.254254
               ## SGAI
## ACCR
               9.12279
                          1.85525
                                   4.917 8.78e-07 ***
## LEVI
              -0.28205 0.30942 -0.912 0.362008
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '* 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##
       Null deviance: 235.72 on 902 degrees of freedom
## Residual deviance: 144.56 on 894 degrees of freedom
## AIC: 162.56
##
## Number of Fisher Scoring iterations: 8
logitModel.fit.predictions <- predict(logitModel.fit, newdata = test1, type</pre>
= "response")
#confusion matrix
predict logitModel.fit<- rep(0, length(logitModel.fit.predictions))</pre>
zeros for predictions
predict logitModel.fit[logitModel.fit.predictions >= 0.5 ] <- 1</pre>
#replace with 1 of prob >= 0.5
table(predict_logitModel.fit,actual2 , dnn= c("Predictions", "Actual"))
             Actual
## Predictions
                0
                    1
            0 321 11
##
                2
##
```

```
#***UNDERSAMPLING***
#split
indx<- sample(2, size=nrow(down data),replace= T, prob = c(0.7, 0.3))
train down <- down data[indx==1,]
test down<- down data[indx==2, ]</pre>
actual down<- test down$C Manipulator
#train
logitModel.fit.UNDER <- glm(C_Manipulator ~ . , data = train_down, family =</pre>
"binomial")
summary(logitModel.fit.UNDER)
## Call:
## glm(formula = C Manipulator ~ ., family = "binomial", data = train down)
##
## Deviance Residuals:
##
       Min
                 1Q
                      Median
                                   3Q
                                           Max
                      0.0000
## -2.1817 -0.7679
                               0.9191
                                        1.5822
##
## Coefficients:
##
               Estimate Std. Error z value Pr(>|z|)
## (Intercept) -5.1216
                            2.9362 -1.744
                                             0.0811 .
                            0.7949
                                     2.275
                                             0.0229 *
## DSRI
                 1.8083
## GMI
                 0.6987
                            0.5384
                                     1.298
                                             0.1944
## AQI
                 0.7630
                            0.3431
                                   2.224
                                             0.0261 *
## SGI
                 2.1790
                            1.2278
                                    1.775
                                             0.0760 .
## DEPI
                -0.4671
                            1.2498 -0.374
                                             0.7086
## SGAI
                -0.8592
                            0.7173 -1.198
                                             0.2310
## ACCR
                7.1009
                            3.8698 1.835
                                             0.0665 .
                            0.5173 -1.212
## LEVI
                -0.6268
                                             0.2257
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '* 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##
       Null deviance: 74.786 on 53 degrees of freedom
## Residual deviance: 51.917 on 45 degrees of freedom
## AIC: 69.917
##
## Number of Fisher Scoring iterations: 8
logitModel.fit.UNDER.predictions <- predict(logitModel.fit.UNDER, newdata =</pre>
test down, type = "response")
#confusion matrix
predict logitModel.UNDER.fit<- rep(0,
length(logitModel.fit.UNDER.predictions)) #all zeros for predictions
predict logitModel.UNDER.fit[logitModel.fit.UNDER.predictions >= 0.5 ] <- 1</pre>
#replace with 1 of prob >= 0.5
table(predict_logitModel.UNDER.fit,actual_down , dnn= c("Predictions",
"Actual"))
```

```
##
             Actual
## Predictions 0 1
##
             0 12 4
             1
               0
                 7
##
#***OVERSAMPLING***
#split
indx<-sample(2, size=nrow(up_data), replace= T, prob = c(0.7, 0.3))
train_up <- up_data[indx==1,]</pre>
test_up<- up_data[indx==2, ]</pre>
actual_up<- test_up$C_Manipulator
#train
logitModel.fit.OVER <- glm(C Manipulator ~ . , data = train up, family =</pre>
"binomial")
summary(logitModel.fit.OVER)
##
## Call:
## glm(formula = C Manipulator ~ ., family = "binomial", data = train up)
## Deviance Residuals:
##
      Min
                 10
                     Median
                                   30
                                          Max
## -4.3761 -0.5375
                     0.0000
                               0.6736
                                       1.7691
##
## Coefficients:
               Estimate Std. Error z value Pr(>|z|)
##
## (Intercept) -8.97978
                          0.59189 -15.171 < 2e-16 ***
## DSRI
               1.77400
                          0.15314 11.584 < 2e-16 ***
## GMI
                          0.19315 10.578 < 2e-16 ***
               2.04306
## AQI
               0.64578
                          0.04269 15.126 < 2e-16 ***
                        0.21936 11.796 < 2e-16 ***
## SGI
               2.58751
## DEPI
               0.49009 0.21517 2.278
                                            0.0227 *
## SGAI
               0.17636
                        0.14058
                                    1.255
                                            0.2096
              10.52655 0.73138 14.393 < 2e-16 ***
## ACCR
               -0.47889
                          0.11400 -4.201 2.66e-05 ***
## LEVI
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##
       Null deviance: 2398.2 on 1730 degrees of freedom
## Residual deviance: 1270.4 on 1722 degrees of freedom
## AIC: 1288.4
##
## Number of Fisher Scoring iterations: 9
logitModel.fit.OVER.predictions <- predict(logitModel.fit.OVER, newdata =</pre>
test up, type = "response")
#confusion matrix
```

```
predict logitModel.OVER.fit<- rep(0, length(logitModel.fit.OVER.predictions))</pre>
#all zeros for predictions
predict logitModel.OVER.fit[logitModel.fit.OVER.predictions >= 0.5 ] <- 1</pre>
#replace with 1 of prob >= 0.5
table(predict_logitModel.OVER.fit,actual_up , dnn= c("Predictions",
"Actual"))
##
             Actual
                0
## Predictions
                    1
            0 311 56
##
             1 49 253
##
#***BOTH***
#split
indx<- sample(2, size=nrow(both data),replace= T, prob = c(0.7, 0.3))
train_both <- both_data[indx==1,]</pre>
test_both<- both_data[indx==2, ]</pre>
actual both<- test both$C Manipulator
#the minority class is oversampled with replacement and majority class is
undersampled without replacement
#train
logitModel.fit.BOTH <- glm(C_Manipulator ~ . , data = train_both, family =</pre>
"binomial")
summary(logitModel.fit.BOTH)
## Call:
## glm(formula = C Manipulator ~ ., family = "binomial", data = train_both)
##
## Deviance Residuals:
##
      Min
                1Q
                     Median
                                   3Q
                                          Max
## -4.8537 -0.5095
                     0.0000
                              0.4688
                                        1.7349
##
## Coefficients:
               Estimate Std. Error z value Pr(>|z|)
## (Intercept) -8.92276
                          0.79701 -11.195 < 2e-16 ***
                                    9.364 < 2e-16 ***
## DSRI
                          0.21681
               2.03032
                                     6.195 5.82e-10 ***
## GMI
               1.55101
                          0.25035
                          0.06205 11.075 < 2e-16 ***
## AQI
               0.68715
## SGI
               2.60875
                          0.29080
                                    8.971 < 2e-16 ***
## DEPI
               0.67790
                          0.31784
                                    2.133
                                            0.0329 *
## SGAI
               0.10742
                          0.19080
                                    0.563
                                            0.5734
## ACCR
              10.21293
                          0.95445 10.700 < 2e-16 ***
## LEVI
              ## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
       Null deviance: 1233.73 on 889
                                      degrees of freedom
## Residual deviance: 632.34 on 881 degrees of freedom
```

```
## AIC: 650.34
##
## Number of Fisher Scoring iterations: 8
logitModel.fit.BOTH.predictions <- predict(logitModel.fit.BOTH, newdata =</pre>
test_both, type = "response")
#confusion matrix
predict logitModel.BOTH.fit<- rep(0, length(logitModel.fit.BOTH.predictions))</pre>
#all zeros for predictions
predict logitModel.BOTH.fit[logitModel.fit.BOTH.predictions >= 0.5 ] <- 1</pre>
#replace with 1 of prob >= 0.5
table(predict_logitModel.BOTH.fit,actual_both , dnn= c("Predictions",
"Actual"))
##
              Actual
## Predictions
                 0
                     1
##
             0 161 31
##
               20 148
#***ROSE***
#split
indx<- sample(2, size=nrow(rose_data),replace= T, prob = c(0.7, 0.3))
train_rose <- rose_data[indx==1,]</pre>
test_rose<- rose_data[indx==2, ]</pre>
actual rose<- test rose$C Manipulator
#train
logitModel.fit.ROSE <- glm(C Manipulator ~ . , data = train rose, family =
"binomial")
summary(logitModel.fit.ROSE)
## Call:
## glm(formula = C_Manipulator ~ ., family = "binomial", data = train_rose)
## Deviance Residuals:
       Min
                 1Q
                      Median
                                   3Q
                                            Max
## -2.0275 -1.0041 -0.5014
                               0.9723
                                         2.8542
##
## Coefficients:
##
                Estimate Std. Error z value Pr(>|z|)
## (Intercept) -0.736406
                           0.226330 -3.254 0.00114 **
## DSRI
                0.090829
                           0.019241
                                      4.721 2.35e-06 ***
## GMI
                0.034552
                           0.016716 2.067 0.03874 *
## AQI
                0.067753
                           0.012514
                                      5.414 6.16e-08 ***
                                     4.855 1.20e-06 ***
## SGI
                0.197038
                           0.040583
## DEPI
                           0.182147 -0.022 0.98205
               -0.004098
## SGAI
                0.035921
                           0.013894
                                     2.585 0.00973 **
                3.547205
                           0.437171
                                      8.114 4.90e-16 ***
## ACCR
## LEVI
                0.005422
                           0.040957
                                     0.132 0.89469
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
##
## (Dispersion parameter for binomial family taken to be 1)
##
       Null deviance: 1230.7 on 887 degrees of freedom
##
## Residual deviance: 1050.8 on 879 degrees of freedom
## AIC: 1068.8
##
## Number of Fisher Scoring iterations: 4
#test
logitModel.fit.ROSE.predictions <- predict(logitModel.fit.ROSE, newdata =</pre>
test rose, type = "response")
#confusion matrix
predict logitModel.ROSE.fit<- rep(0, length(logitModel.fit.ROSE.predictions))</pre>
#all zeros for predictions
predict_logitModel.ROSE.fit[logitModel.fit.ROSE.predictions >= 0.5 ] <- 1</pre>
#replace with 1 of prob >= 0.5
table(predict logitModel.ROSE.fit,actual rose , dnn= c("Predictions",
"Actual"))
##
              Actual
## Predictions
                 0
                     1
##
             0 142 74
##
             1 39 96
#EVALUATION
print("Accuracy/Recall/Precision/F-1 score NOT using balancing technique")
my.accuracy(actual2, predict_logitModel.fit)
## [1] 0.9613095
my.recall(actual2, predict_logitModel.fit)
## [1] 0.1538462
my.precision(actual2, predict logitModel.fit)
## [1] 0.5
my.f1(actual2, predict_logitModel.fit)
## [1] 0.2352941
print("Accuracy/Recall/Precision/F-1 score using UNDER SAMPLING")
my.accuracy(actual down, predict logitModel.UNDER.fit)
## [1] 0.826087
my.recall(actual_down, predict_logitModel.UNDER.fit)
## [1] 0.6363636
```

```
my.precision(actual_down, predict_logitModel.UNDER.fit)
## [1] 1
my.f1(actual_down, predict_logitModel.UNDER.fit)
## [1] 0.7777778
print("Accuracy/Recall/Precision/F-1 score using OVER SAMPLING")
my.accuracy(actual_up, predict_logitModel.OVER.fit)
## [1] 0.8430493
my.recall(actual_up, predict_logitModel.OVER.fit)
## [1] 0.8187702
my.precision(actual_up, predict_logitModel.OVER.fit)
## [1] 0.8377483
my.f1(actual_up, predict_logitModel.OVER.fit)
## [1] 0.8281506
print("Accuracy/Recall/Precision/F-1 score using BOTH UNDER AND OVER
SAMPLING")
my.accuracy(actual_both, predict_logitModel.BOTH.fit)
## [1] 0.8583333
my.recall(actual_both, predict_logitModel.BOTH.fit)
## [1] 0.8268156
my.precision(actual_both, predict_logitModel.BOTH.fit)
## [1] 0.8809524
my.f1(actual_both, predict_logitModel.BOTH.fit)
## [1] 0.8530259
print("Accuracy/Recall/Precision/F-1 score using ROSE")
my.accuracy(actual_rose, predict_logitModel.ROSE.fit)
## [1] 0.6780627
my.recall(actual_rose, predict_logitModel.ROSE.fit)
## [1] 0.5647059
my.precision(actual_rose, predict_logitModel.ROSE.fit)
```

```
## [1] 0.7111111

my.f1(actual_rose, predict_logitModel.ROSE.fit)

## [1] 0.6295082

#Model with highest F-1 score is obtained while using BOTH under and over sampling. It's important to notice that all models with the different balancing techniques have a much greater F-1 score compared to the model using the unbalanced data.
```

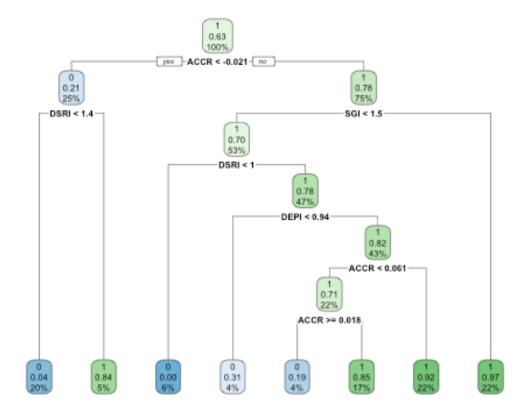
	UNBALANCED	UNDERSAMPLE	OVERSAMPLE	ВОТН	ROSE
Accuracy	0.9613095	0.826087	0.8430493	0.8583333	0.6780627
Recall	0.1538462	0.6363636	0.8187702	0.8268156	0.5647059
Precision	0.5	1	0.8377483	0.8809524	0.7111111
F-1 Score	0.2352941	0.77777781	0.8281506	0.8530259	0.6295082

CLASSIFICATION & REGRESSION TREE(CART model)-using sample data

```
Balanced_Data <- ovun.sample(C_Manipulator ~., data = sample_data, method =
"over", N = 500)$data

set.seed(123)
indx <- sample(2, nrow(Balanced_Data), replace = T, prob = c(0.7, 0.3))
train <- Balanced_Data [indx == 1, ]
test <- Balanced_Data [indx == 2, ]

library (rpart)
decision_tree <- rpart(C_Manipulator ~., data = train)
library(rpart.plot)
rpart.plot(decision_tree)</pre>
```



```
decision_tree_pred <- predict(decision_tree, test, type = "prob")</pre>
decision_tree_class <- predict(decision_tree, test, type = "class")</pre>
actual <- test$C_Manipulator</pre>
table(decision_tree_class, actual, dnn= c("predictions", "actual"))
##
              actual
## predictions
                0
             0 36
                  7
##
             1 14 86
##
my.accuracy(actual, decision_tree_class)
## [1] 0.8531469
my.recall(actual, decision_tree_class)
## [1] 0.9247312
my.precision(actual, decision_tree_class)
## [1] 0.86
my.f1(actual, decision_tree_class)
## [1] 0.8911917
```

	OVERSAMPLED		
Accuracy	0.8531469		
Recall	0.9247312		
Precision	0.86		
F-1 Score	0.8911917		

RANDOM FOREST-using sample data

```
set.seed(123)
library(randomForest)
Balanced_Data <- ovun.sample(C_Manipulator ~., data = sample_data, method =
"over", N = 500)$data
rf <- randomForest(C_Manipulator ~ ., data = Balanced_Data, mtry =</pre>
sqrt(ncol(Balanced_Data)-1), ntree = 300, proximity = T, importance = T)
rf
##
## Call:
## randomForest(formula = C_Manipulator ~ ., data = Balanced_Data,
                                                                         mtry
= sqrt(ncol(Balanced_Data) - 1), ntree = 300, proximity = T,
                                                               importance
= T)
##
                  Type of random forest: classification
##
                        Number of trees: 300
## No. of variables tried at each split: 3
##
##
           OOB estimate of error rate: 1.8%
## Confusion matrix:
       0 1 class.error
## 0 172 9 0.04972376
## 1
      0 319 0.00000000
#recall
319/(319+9) # = 0.972561
## [1] 0.972561
#precision
319/(319+0) # = 1
## [1] 1
#f-score
2 * 1 * 0.972561 / (1 + 0.972561) # = 0.9860897
 ## [1] 0.9860897
```

	OVERSAMPLED		
Accuracy	0.8531469		
Recall	0.972561		
Precision	1		
F-1 Score	0.9860897		

Conclusion: The final recommendation for predicting earnings manipulation would be to use balanced data and then deploy a random forest model. This model yields the highest F-1 Score of all models tried, which best measures accuracy of results for this case.

	RF-	Decision Tree-	Logistic Reg-				
	Oversampled	Oversampled	Unbalanced	Undersampled	Oversampled	Both	Rose
F-1	0.9860897	0.8911917	0.2352941	0.7777781	0.8281506	0.8530259	0.6295082
Score							