Random Forest

```
## Loading required package: lattice
## Loading required package: ggplot2
```

Stratified sampling

Random Forest

- can handles the overfitting problem you faced with decision trees. This approach having the majority's vote count as the actual classification decision.
- can deal with "small n large p"-problem, high-order interactions, correlated predictor variables.

```
formular1 =
as.factor(target)~new_outstanding_principal_balance+initial_loan_amount+fico+
sales_channel__c+type+current_collection_method+term+average_bank_balance__c+
lender1
my_forest <- train(formular1, method="rf", data=training,</pre>
trControl=trainControl(method="cv"), number=3)
## Loading required package: randomForest
## randomForest 4.6-10
## Type rfNews() to see new features/changes/bug fixes.
print(my_forest)
## Random Forest
##
## 358 samples
## 18 predictor
     2 classes: '0', '1'
##
##
## No pre-processing
## Resampling: Cross-Validated (10 fold)
##
## Summary of sample sizes: 322, 322, 322, 322, 323, ...
##
```

```
## Resampling results across tuning parameters:
##
##
    mtry Accuracy
                     Kappa
                                Accuracy SD Kappa SD
     2
          0.7151587 0.2768319 0.04241798
                                            0.1280355
##
     7
          0.6677778 0.2236602 0.08039734
##
                                            0.1616732
          0.6594444 0.2058128 0.07842160
                                            0.1513176
##
##
## Accuracy was used to select the optimal model using the largest value.
## The final value used for the model was mtry = 2.
my_forest$finalModel
##
## Call:
## randomForest(x = x, y = y, mtry = param$mtry, number = 3)
                 Type of random forest: classification
                       Number of trees: 500
## No. of variables tried at each split: 2
##
          OOB estimate of error rate: 29.33%
##
## Confusion matrix:
        1 class.error
##
     0
## 0 36 93 0.72093023
## 1 12 217 0.05240175
```

Apply Random Forest to the testing data

```
forest pred <- predict(my forest, testing)</pre>
forest pred df <- data.frame(forest pred, target=testing$target)</pre>
confusionForest=table(forest pred df$forest pred, forest pred df$target)
print(confusionForest)
##
##
      0 1
##
     0 13 6
## 1 21 79
Observed Accuracy =
(confusionForest[1,1]+confusionForest[2,2])/sum(confusionForest)
print(Observed_Accuracy)
## [1] 0.7731092
Expected Accuracy =
(sum(confusionForest[,1])*sum(confusionForest[1,])/sum(confusionForest)+sum(c
onfusionForest[,2])*sum(confusionForest[2,])/sum(confusionForest))/sum(confus
ionForest)
```

```
kappa = (Observed_Accuracy-Expected_Accuracy)/(1-Expected_Accuracy)
print(kappa)
## [1] 0.359322
```

Random forest variable importance measures and plots

```
forestImp <- round(varImp(my forest$finalModel, scale = FALSE),2)</pre>
forestImp1=forestImp[order(forestImp$Overall),,drop=FALSE]
forestImp[order(-forestImp$Overall),,drop=FALSE]
##
                                                      Overall
## new_outstanding_principal_balance
                                                        28.00
## average_bank_balance__c
                                                        17.50
## fico
                                                        16.58
## initial_loan_amount
                                                        11.35
## term
                                                          5.91
## typeLoan - Renewal
                                                          3.82
## current_collection_methodTransfer Account Vendors
                                                          2.38
## current collection methodSplit Funding
                                                          2.34
## sales channel cFAP: Managed Application Program
                                                          2.10
## lender1
                                                          1.91
## sales channel cReferral
                                                          1.24
## sales_channel__cPromontory
                                                          0.39
counts <- forestImp1[,1]</pre>
par(mai=c(1,2,1,1))
bp=barplot(counts, main="Feature Importance plot", horiz=TRUE,
names.arg=rownames(forestImp1),col="light blue",las=1)
text(x= counts, bp, labels=as.character(counts), pos=4)
```

Feature Importance plot



