classification_Xinming

Xinming Liu

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Set A - Debate 1

```
## SetA Preprocessing
users <- read.csv("user_setA/users.csv", stringsAsFactors = FALSE)</pre>
#head(a.users)
tweets <- read.csv("user_setA/tweets_debateVP.csv", stringsAsFactors = FALSE)</pre>
# tweets2 <- read.csv("user_setA/tweets_debate2.csv", stringsAsFactors = FALSE)
# tweets3 <- read.csv("user_setA/tweets_debate3.csv", stringsAsFactors = FALSE)
# tweets4 <- read.csv("user_setA/tweets_debateVP.csv", stringsAsFactors = FALSE)
# tweets <- rbind(tweets1, tweets2, tweets3, tweets4)</pre>
#head(tweets)
tweets<-tweets%>%
  mutate(userID=as.numeric((userID)))
data <- data.table(users, key="userID")[</pre>
 data.table(tweets, key="userID"),
 allow.cartesian=TRUE
data <- subset(data, party=='D' | party=='R')</pre>
#data <- subset(data, state_code=="PA")</pre>
#head(data)
use <- data.frame(data$text, factor(data$party), stringsAsFactors = FALSE)</pre>
colnames(use) <- c("text", "party")</pre>
#head(use)
## Create DocumentTermMatrix
corpus <- Corpus(VectorSource(use$text))</pre>
corpus = clean_corpus(corpus)
#td.mat = TermDocumentMatrix(corpus)
dt.mat = DocumentTermMatrix(corpus)
## dt.mat is not a matrix here
## Feature words extraction (this may create NA values) due to limited memory
dt.mat.use = removeSparseTerms(dt.mat, 0.95)
## Sparsity = 0.95 (7 terms remaining) ~ 0.97 (17 terms remaining) seems acceptable
## Attach class label
alldata <- as.matrix(dt.mat.use)</pre>
alldata <- cbind(alldata, use$party)</pre>
colnames(alldata) [ncol(alldata)] <- "Class"</pre>
## Class=1 for Democrats, Class=2 for Republican
alldata <- as.data.frame(alldata)</pre>
```

```
alldata$Class <- as.factor(alldata$Class)
levels(alldata$Class) <- c("Democrats", "Republican")</pre>
```

Use 10-fold CV with 70% data for training, 30% data for testing

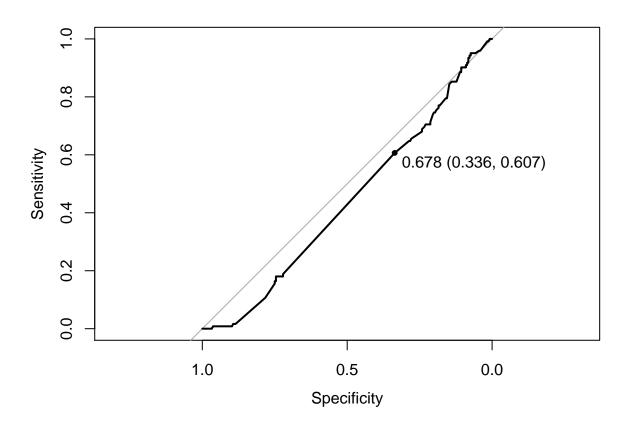
```
## Train-Test Split
set.seed(9000)
## 70% for training, 30% for testing
TrainingDataIndex <- createDataPartition(alldata$Class, p=0.7, list = FALSE)
trainingData <- alldata[TrainingDataIndex,]
testData <- alldata[-TrainingDataIndex,]
## 10-fold CV (cannot do repeatedcv due to CPU performance)
TrainingParameters <- trainControl(method = "cv", number = 10, classProbs = TRUE, summaryFunction = two</pre>
```

Classification using kNN (too many ties)

```
# ## Training
# fit <- train(Class ~ ., data = trainingData,
                   method = "knn",
#
                    trControl = TrainingParameters,
#
                   tuneGrid = expand.grid(k = seq(1, 10, length = 10)),
#
                   preProcess = c("scale", "center"),
#
                   na.action = na.omit
# )
# fit
# fit$bestTune
# ## Testing
# pred <- predict(fit, testData)</pre>
# ## Evaluation
# confusionMatrix(pred, testData$Class)
# ## Rank terms by importance
# importance <- varImp(fit, scale=FALSE)</pre>
# plot(importance)
```

Classification using SVM

```
## Support Vector Machines with Linear Kernel
##
## 857 samples
## 10 predictor
##
    2 classes: 'Democrats', 'Republican'
##
## Pre-processing: scaled (10), centered (10)
## Resampling: Cross-Validated (10 fold)
## Summary of sample sizes: 771, 772, 770, 771, 771, ...
## Resampling results across tuning parameters:
##
##
            ROC
                       Sens
                                 Spec
    0.1000 0.5896150 0.9860254 0.02770936
##
##
    0.2125  0.6138639  0.9824561  0.04187192
##
    0.3250 0.5815255 0.9825166 0.03448276
    ##
##
    0.5500 0.6024238 0.9947368 0.01379310
##
    0.6625 0.5954164 0.9754991 0.05246305
##
    0.7750 0.6191039 0.9754991 0.05665025
    0.8875 0.5901019 0.9859649 0.03879310
##
##
    1.0000 0.6121968 0.9842105 0.03842365
## ROC was used to select the optimal model using the largest value.
## The final value used for the model was C = 0.775.
fit.lSVM$bestTune
##
        C
## 7 0.775
## Testing
pred.lSVM <- predict(fit.lSVM, testData, type="prob")</pre>
## Evaluation
#confusionMatrix(pred, testData$Class)
result.roc <- roc(testData$Class, pred.1SVM$Democrats)</pre>
plot(result.roc, print.thres="best", print.thres.best.method="closest.topleft")
```

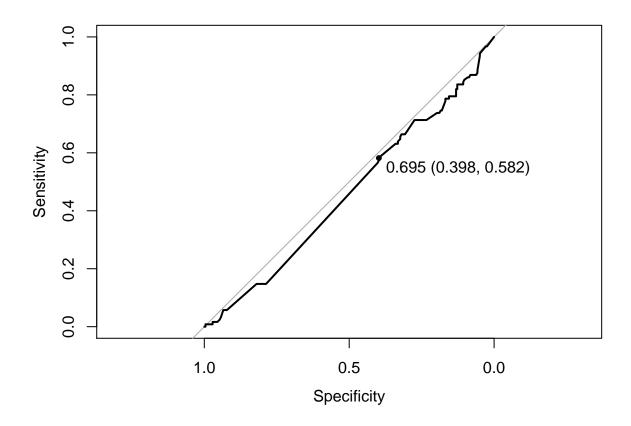


result.coords <- coords(result.roc, "best", best.method="closest.topleft", ret=c("threshold", "accuracy
print(result.coords)</pre>

```
## threshold accuracy
## threshold 0.6782905 0.4262295
```

```
# set.seed(9002)
# ## Using polynomial kernel
# ## Training
\textit{\# \#\# Due to CPU performance, cannot apply a grid to tune parameters}
# fit.pSVM <- train(Class ~ ., data = trainingData,</pre>
                    method = "sumPoly",
#
                    metric = "ROC",
                    trControl = TrainingParameters,
                    preProcess = c("scale", "center"),
#
                    na.action = na.omit
# )
# fit.pSVM
\# \ fit.pSVM\$bestTune
# ## Testing
# pred.pSVM <- predict(fit.pSVM, testData)</pre>
# ## Evaluation
```

```
# #confusionMatrix(pred, testData$Class)
# result.roc <- roc(testData$Class, pred.pSVM$Democrats)</pre>
# plot(result.roc, print.thres="best", print.thres.best.method="closest.topleft")
\# result.coords <- coords(result.roc, "best", best.method="closest.topleft", ret=c("threshold", "accura
# print(result.coords)
set.seed(9003)
## Using radial basis kernel
## Training
## Due to CPU performance, cannot apply a grid to tune parameters
fit.rSVM <- train(Class ~ ., data = trainingData,</pre>
                 method = "svmRadial",
                 metric = "ROC",
                 trControl= TrainingParameters,
                 preProcess = c("scale", "center"),
                 na.action = na.omit
fit.rSVM
## Support Vector Machines with Radial Basis Function Kernel
##
## 857 samples
## 10 predictor
    2 classes: 'Democrats', 'Republican'
##
##
## Pre-processing: scaled (10), centered (10)
## Resampling: Cross-Validated (10 fold)
## Summary of sample sizes: 771, 771, 772, 771, 771, 772, ...
## Resampling results across tuning parameters:
##
##
           ROC
                      Sens
                                 Spec
##
    0.25  0.6108551  0.9789474  0.04199507
##
   0.50 0.6079819 0.9685118 0.06958128
     1.00 0.6253011 0.9579855 0.12931034
##
## Tuning parameter 'sigma' was held constant at a value of 0.1100243
## ROC was used to select the optimal model using the largest value.
## The final values used for the model were sigma = 0.1100243 and C = 1.
fit.rSVM$bestTune
         sigma C
## 3 0.1100243 1
## Testing
pred.rSVM <- predict(fit.rSVM, testData, type="prob")</pre>
## Evaluation
#confusionMatrix(pred, testData$Class)
result.roc <- roc(testData$Class, pred.rSVM$Democrats)</pre>
plot(result.roc, print.thres="best", print.thres.best.method="closest.topleft")
```



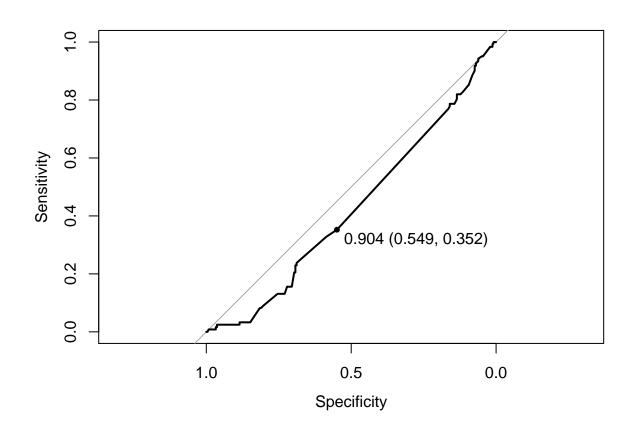
```
result.coords <- coords(result.roc, "best", best.method="closest.topleft", ret=c("threshold", "accuracy print(result.coords)
```

```
## threshold accuracy
## threshold 0.6946895 0.4590164
```

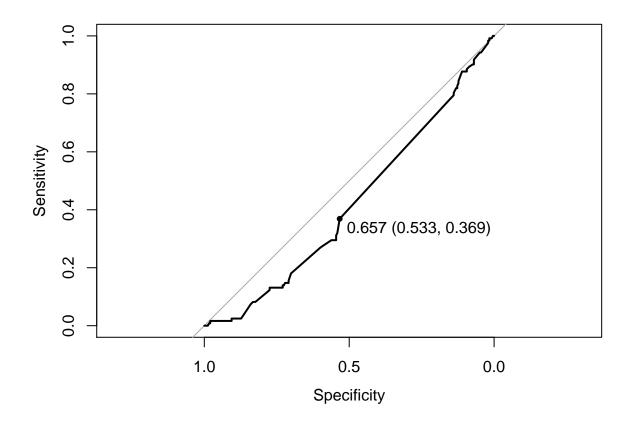
Classification using Naive Bayes

```
## Naive Bayes
##
## 857 samples
## 10 predictor
## 2 classes: 'Democrats', 'Republican'
##
```

```
## Pre-processing: scaled (10), centered (10)
## Resampling: Cross-Validated (10 fold)
## Summary of sample sizes: 771, 772, 772, 771, 771, 771, ...
## Resampling results across tuning parameters:
##
##
     usekernel ROC
                           Sens
                                      Spec
##
     FALSE
                0.6244815 0.8287961 0.2850985
      TRUE
                0.6294962 0.9982456 0.0000000
##
##
## Tuning parameter 'fL' was held constant at a value of 0
## Tuning
## parameter 'adjust' was held constant at a value of 1
## ROC was used to select the optimal model using the largest value.
## The final values used for the model were fL = 0, usekernel = TRUE and adjust
## = 1.
## Testing
pred.NB <- predict(fit.NB, testData, type="prob")</pre>
## Evaluation
#confusionMatrix(pred, testData$Class)
result.roc <- roc(testData$Class, pred.NB$Democrats)</pre>
plot(result.roc, print.thres="best", print.thres.best.method="closest.topleft")
```



```
result.coords <- coords(result.roc, "best", best.method="closest.topleft", ret=c("threshold", "accuracy
print(result.coords)
             threshold accuracy
## threshold 0.9042312 0.4836066
Classification using Logistic Regression
set.seed(9005)
## Training
fit.LR <- train(Class ~ ., data = trainingData,</pre>
                    method = "glm",
                    metric = "ROC",
                    preProcess=c("scale","center"),
                    trControl= TrainingParameters,
                    na.action = na.omit
)
fit.LR
## Generalized Linear Model
##
## 857 samples
## 10 predictor
    2 classes: 'Democrats', 'Republican'
##
## Pre-processing: scaled (10), centered (10)
## Resampling: Cross-Validated (10 fold)
## Summary of sample sizes: 771, 771, 770, 771, 772, 772, ...
## Resampling results:
##
##
     ROC
                Sens
                            Spec
##
     0.6378245 0.9002117 0.18633
## Testing
pred.LR <- predict(fit.LR, testData, type="prob")</pre>
## Evaluation
#confusionMatrix(pred, testData$Class)
result.roc <- roc(testData$Class, pred.LR$Democrats)</pre>
plot(result.roc, print.thres="best", print.thres.best.method="closest.topleft")
```



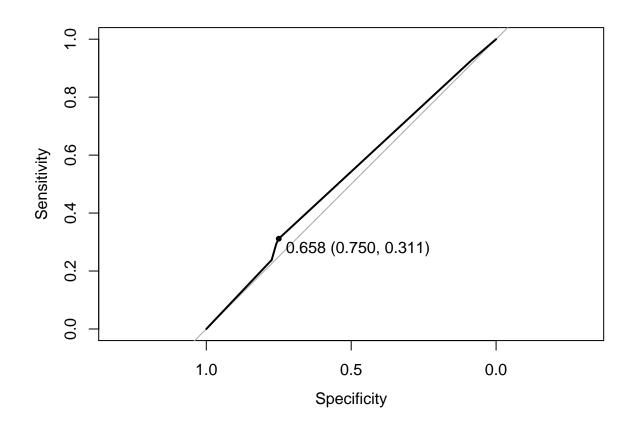
```
result.coords <- coords(result.roc, "best", best.method="closest.topleft", ret=c("threshold", "accuracy
print(result.coords)</pre>
```

```
## threshold accuracy
## threshold 0.6567117 0.4781421
```

Classification using Decision Tree

```
set.seed(9006)
## Training
fit.DT <- train(Class ~ ., data = trainingData,</pre>
                    method = "rpart",
                    metric = "ROC",
                    preProcess=c("scale","center"),
                     trControl= TrainingParameters,
                     na.action = na.omit
)
fit.DT
## CART
##
## 857 samples
    10 predictor
##
##
     2 classes: 'Democrats', 'Republican'
##
```

```
## Pre-processing: scaled (10), centered (10)
## Resampling: Cross-Validated (10 fold)
## Summary of sample sizes: 772, 771, 770, 772, 771, 772, ...
## Resampling results across tuning parameters:
##
##
                  ROC
                             Sens
                                         Spec
##
     0.001169591
                  0.5818926
                             0.9160315
                                        0.17204433
##
     0.010526316
                  0.5536807
                             0.9456443
                                        0.08115764
##
     0.012280702  0.5438750  0.9579250  0.05615764
##
## ROC was used to select the optimal model using the largest value.
## The final value used for the model was cp = 0.001169591.
## Testing
pred.DT <- predict(fit.DT, testData, type="prob")</pre>
## Evaluation
#confusionMatrix(pred, testData$Class)
result.roc <- roc(testData$Class, pred.DT$Democrats)</pre>
plot(result.roc, print.thres="best", print.thres.best.method="closest.topleft")
```

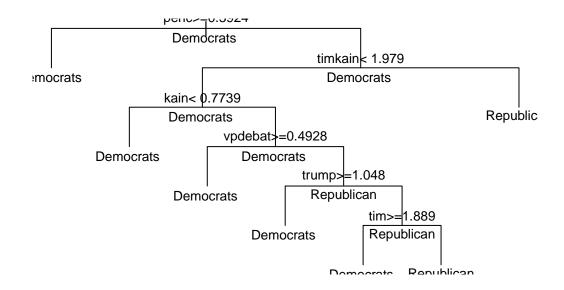


```
result.coords <- coords(result.roc, "best", best.method="closest.topleft", ret=c("threshold", "accuracy print(result.coords)
```

```
## threshold accuracy
## threshold 0.6582397 0.6038251
```

```
plot(fit.DT$finalModel, uniform=TRUE, main="Classification Tree")
text(fit.DT$finalModel, use.n.=TRUE, all=TRUE, cex=.8)
```

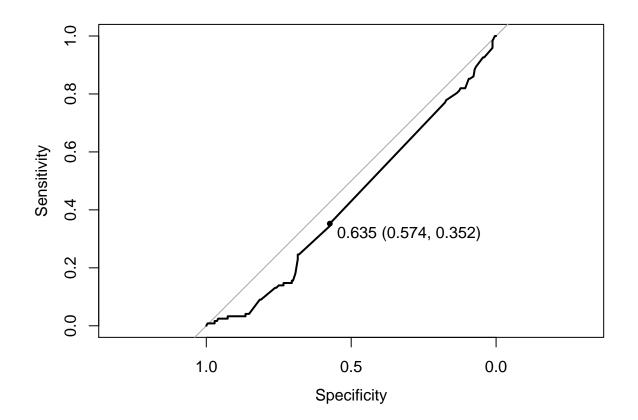
Classification Tree



Classification using AdaBoost

```
## Boosted Classification Trees
##
## 857 samples
## 10 predictor
## 2 classes: 'Democrats', 'Republican'
##
## Pre-processing: scaled (10), centered (10)
## Resampling: Cross-Validated (10 fold)
## Summary of sample sizes: 772, 770, 770, 771, 772, 771, ...
```

```
## Resampling results across tuning parameters:
##
     maxdepth iter
                                           Spec
##
                     ROC
                                Sens
                                1.0000000
                                           0.00000000
##
                50
                     0.6210489
##
     1
               100
                     0.6258082 1.0000000
                                           0.00000000
               150
                     0.6340619 1.0000000 0.007142857
##
     1
##
     2
                50
                     0.6149828 0.9842408 0.031773399
               100
##
     2
                     0.6179529 0.9596794
                                           0.080665025
                                           0.122906404
##
     2
               150
                     0.6272023 0.9509074
                50
##
     3
                     0.6229648 0.9544465
                                          0.109236453
##
     3
               100
                     0.6286640 0.9474894
                                           0.150862069
               150
                     0.6289168 0.9387477
##
     3
                                           0.164901478
##
## Tuning parameter 'nu' was held constant at a value of 0.1
## ROC was used to select the optimal model using the largest value.
## The final values used for the model were iter = 150, maxdepth = 1 and nu = 0.1.
## Testing
pred.ADA <- predict(fit.ADA, testData, type="prob")</pre>
## Evaluation
#confusionMatrix(pred, testData$Class)
result.roc <- roc(testData$Class, pred.ADA$Democrats)</pre>
plot(result.roc, print.thres="best", print.thres.best.method="closest.topleft")
```



```
result.coords <- coords(result.roc, "best", best.method="closest.topleft", ret=c("threshold", "accuracy
print(result.coords)</pre>
```

```
## threshold accuracy
## threshold 0.6350066 0.5
```