Machine Learning Code

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library(tidyverse) # map_df();

library(caret) # createDataPartition();

library(AppliedPredictiveModeling) # transparentTheme();

```
library(pROC) # roc();
library(rpart.plot) # rpart.plot();
library(ranger) # for variable importance of rf
library(pdp) # PDP
library(lime) # prediction explaination
library(factoextra) # clustering visualization
library(corrplot)
# import the raw data
DATA = read.csv(file = './OnlineNewsPopularity/OnlineNewsPopularity.csv', head = TRUE, sep = ',')
# check for missing data
sum( map_df(DATA, function(x) sum(is.na(x))) )
# NOTES: The original data contains no missing data.
1 Preprocessing
# raw data cleaning part1
Data_prep_1 =
 DATA[,-1] %>%
  mutate(., data_channel_is_other = ifelse(data_channel_is_lifestyle+data_channel_is_entertainment+data
  gather(., key = channel_type, value = data_channel_is,
            data_channel_is_lifestyle:data_channel_is_world, data_channel_is_other) %>%
 filter(., data_channel_is == '1') %>%
  select(., -data_channel_is) %>%
  mutate(., channel_type = str_replace(channel_type, 'data_channel_is_', ''),
            channel type = forcats::fct relevel(channel type, 'other'))
# In this part, we deal with the channel types. First, we created a new variable 'other' that refers to
# raw data cleaning part2
  gather(Data_prep_1, key = publish_day, value = day_is, weekday_is_monday:weekday_is_sunday) %>%
  filter(., day_is == '1') %>%
  select(., -day_is) %>%
  mutate(., publish_day = str_replace(publish_day, 'weekday_is_', ''),
            publish_day = forcats::fct_relevel(publish_day, 'monday'))
# In this part, similarly, we gather 7 columns into one single variable 'publish_day', and make 'monday
# delete uninformative variables
Data_prep_3 =
 select(Data_prep_2, -n_unique_tokens, -n_non_stop_words, -n_non_stop_unique_tokens, -timedelta)
# NOTES: the first three variables has nearly 0 variance, in addition to one or two outliers.
```

NOTES: timedelta cannot be used to predict 'popularity' before the article get published.

```
# Using 1500 as threshold to separate 'shares' into 'popular' and 'unpopular'.
Data_prep_4 =
 mutate(Data_prep_3, popularity = case_when(shares >= 1500 ~ 'popular',
                                             shares < 1500 ~ 'unpopular')) %>%
 mutate(., popularity = forcats::fct_relevel(popularity, 'unpopular')) %>%
 select(., -shares)
#Data_prep_4 %>% count(., popularity)
# NOTES: make 'unpopular' to be the reference category.
# NOTES: the raw dataset ended up with 19562 'popular' and 20082 'unpopular' articles.
# Randomly take a subset of the whole data as the original dataset.
set.seed(4)
subset = createDataPartition(Data_prep_4$popularity, p = 0.05, list = FALSE) # 5% of the original datas
Data_prep_5 = Data_prep_4[subset,]
# NOTES: randomly choose 5% of the whole dataset.
# because of collinearity issue, we decided to take out these two variables.
Data_prep_6 = Data_prep_5 %>% select(., -publish_day, -LDA_04)
# the whole dataset after preprocessing
News_Data = Data_prep_6
# training and test dataset split
set.seed(4)
trRows = createDataPartition(News_Data$popularity, p = 0.7, list = FALSE)
###training data###
News Data train = News Data[trRows,]
###test data###
News_Data_test = News_Data[-trRows,]
# training + test data
x_total = model.matrix(popularity~., News_Data)[,-1]
y_total = News_Data$popularity
# training data with 1390 observations
x_train = model.matrix(popularity~., News_Data_train)[,-1]
y_train = News_Data_train$popularity
# test data with 594 observations
x_test = model.matrix(popularity~., News_Data_test)[,-1]
y_test = News_Data_test$popularity
# NOTES: 70% training + 30% test
```

Explorataty Analysis

```
# correlation coefficient matrix
corrplot::corrplot(cor(x_total), tl.cex = 0.45)
```

K means clustering

Hierarchical clustering

PCA

```
pca = prcomp(x_total_scaled)
pca$rotation
```

```
corrplot(pca$rotation %*% diag(pca$sdev), tl.cex = 0.45)
```

```
# scree plot
fviz_eig(pca, addlabels = TRUE)
```

```
fviz_contrib(pca, choice = 'var', axes = 1)
```

2 Model Selection all using 'caret' package

Logistic Regression without regularization

```
# logistic regression (glm) without regularization
# train model with training data using 'caret' package
set.seed(4)
glm.fit = train(x = x_train, y = y_train,
```

Logistic Regression with regularization

```
# logistic regression (glmn) with regularization (elastic)
# train model with training data using 'caret' package
glmnGrid = expand.grid(.alpha = seq(0, 1, length = 6),
                       .lambda = exp(seq(-10, -1, length = 20)))
set.seed(4)
glmn.fit = train(x = x_train, y = y_train,
                 method = 'glmnet',
                 tuneGrid = glmnGrid,
                 metric = 'Accuracy',
                 trControl = trainControl(method = 'cv', number = 10, classProbs = TRUE))
# tuning result for alpha, lambda
glmn_tune = glmn.fit$bestTune
glmn_tune
# plot of the tuning result
\#plot(glmn.fit, xTrans = function(x) log(x), highlight = TRUE)
ggplot(glmn.fit, highlight = TRUE) + scale_x_log10()
```

Linear Discriminant Analysis LDA

Quadratic Discriminant Analysis QDA

Naive Bayes

```
# Naive Bayes (nb)
# train model with training data using 'caret' package
# library("klaR")
### takes too long to run, very lagging ###
set.seed(4)
nbGrid = expand.grid(usekernel = TRUE,
                  fL = 1,
                  adjust = 10)
nb.fit = train(x = x_train[, 1:2], y = y_train,
            preProcess = c('center', 'scale'),
            method = 'nb',
            tuneGrid = nbGrid,
            metric = 'Accuracy',
            trControl = trainControl(method = 'cv', number = 10, classProbs = TRUE))
dim(x_train)
# NOTES: because of the '0' values in the dataset, trainig for this model didn't work.
#plot(nb.fit)
```

Classification Tree

```
# tuning result for cp
ct_tune = ct.fit$bestTune
ct_tune
```

```
# plot of the tuning result
ggplot(ct.fit, highlight = TRUE)

# plot the tree
rpart.plot(ct.fit$finalModel)
```

Random Forest

```
# Random Forest (rf)
# train model with training data using 'caret' package
### takes about 70 min to run ###
###################################
rf.grid = expand.grid(mtry = 1:47,
                     splitrule = 'gini',
                     min.node.size = 1:10)
set.seed(4)
rf.fit = train(popularity~., News_Data_train,
              method = 'ranger',
              tuneGrid = rf.grid,
              metric = 'Accuracy',
              trControl = trainControl(method = 'cv', number = 10, classProbs = TRUE))
# save the object
#saveRDS(rf.fit, file = './rf.fit.rda')
# read in the object
rf.fit = readRDS('/Users/junyuanzheng/Desktop/Data Science II/Final_Project/rf.fit.rda')
# tuning result for mtry, min.node.size
rf_tune = rf.fit$bestTune
rf_tune
# plot of the tuning result
ggplot(rf.fit, highlight = TRUE)
```

#rf.fit resample

Boosting

SVM with linear kernel

```
# save the object
#saveRDS(svml.fit, file = './svml.fit.rda')
# read in the object
svml.fit = readRDS('/Users/junyuanzheng/Desktop/Data Science II/Final_Project/svml.fit.rda')
#summary(svml.fit)
ggplot(svml.fit, highlight = TRUE)
#best.svml.fit = svml.fit$bestTune # cost =
```

SVM with radial kernel

```
# save the object
#saveRDS(svmr.fit, file = './svmr.fit.rda')
# read in the object
svmr.fit = readRDS('/Users/junyuanzheng/Desktop/Data Science II/Final_Project/svmr.fit.rda')
#summary(svmr.fit)
ggplot(svmr.fit, highlight = TRUE)
#best.svmr.fit = svmr.fit$bestTune # sigma = 0.00309, C = 8.209
```

Summary

Accuracy comparison:

```
# parallel plot of accuracy
parallelplot(resamp, metric = 'Accuracy', lwd = 3, lty = 1, alpha = 0.7)

# box plot of accuracy
bwplot(resamp, metric = 'Accuracy')
```

Kappa comparison:

```
# parallel plot of accuracy
parallelplot(resamp, metric = 'Kappa', lwd = 3, lty = 1, alpha = 0.7)

# box plot of accuracy
bwplot(resamp, metric = 'Kappa')
```

Model comparison are based mainly on Accuracy and Kappa.

As a result, SVM with radial kernel, Random Forest, and Boosting appear to have a better performance on the training dataset using Cross Validation.

Therefore, we are going to perform these three models on the test dataset.

3 Estimation on the test dataset

```
# SVM with radial kernel on test dataset
svmr.pred_test = predict(svmr.fit, newdata = News_Data_test[, -43])
svmr_Accuracy_test = mean(svmr.pred_test == News_Data_test[, 43])
svmr_Accuracy_test
confusionMatrix(data = svmr.pred_test,
                reference = News_Data_test[, 43],
                positive = 'popular')
# Random Forest on test dataset
rf.pred test = predict(rf.fit, newdata = News Data test[, -43])
rf_Accuracy_test = mean(rf.pred_test == News_Data_test[, 43])
rf_Accuracy_test
confusionMatrix(data = rf.pred_test,
                reference = News_Data_test[, 43],
                positive = 'popular')
# Boosting on test dataset
gbm.pred_test = predict(gbm.fit, newdata = News_Data_test[, -43])
gbm_Accuracy_test = mean(gbm.pred_test == News_Data_test[, 43])
gbm_Accuracy_test
confusionMatrix(data = gbm.pred_test,
                reference = News_Data_test[, 43],
                positive = 'popular')
```

4 Check important variables

Variable importance from random forest

```
# variable importance using Permutation
# refit the model using 'ranger' with the tuned hyperparameter
set.seed(4)
rf_VI.per = ranger(popularity~., News_Data_train,
                      mtry = 2, splitrule = 'gini',
                      min.node.size = 2,
                      importance = 'permutation',
                      scale.permutation.importance = TRUE)
# barplot
par(mai = c(0.5, 1.2, 0.5, 0.3))
barplot(sort(ranger::importance(rf_VI.per), decreasing = FALSE),
        las = 2, horiz = TRUE, cex.names = 0.4,
        col = colorRampPalette(colors = c('red', 'cyan', 'blue'))(50), main = 'Variable Importance by P
        space = 0.3)
# variable importance using Impurity
# refit the model using 'ranger' with the tuned hyperparameter
set.seed(4)
rf_VI.imp = ranger(popularity~., News_Data_train,
                   mtry = 2, splitrule = 'gini',
                   min.node.size = 2,
                   importance = 'impurity')
```

Variable importance from Boosting

```
# variable importance from Boosting
par(mai = c(0.5, 1.2, 0.5, 0.3))
summary(gbm.fit$finalModel, las = 2, cBars = 42, cex.names = 0.4, space = 0.3)
```

NOTES:By looking at the variable importance plot from fitted RF and Boosting model, they all agree that 'kw_avg_avg' to be an important vaiable relative to the others. Therefore, we are going to focus on this variable in the following Partial Dependence Plots (PDP).

Partial Dependence Plots (PDP) of 'kw_avg_avg'

Individual Conditional Expectation (ICE) of 'kw_avg_avg'

```
autoplot(train = News_Data_train, alpha = 0.02, center = TRUE) +
  ggtitle('Random Forest, centered') +
  theme(axis.text = element_text(size = 6))
ice1.gbm = gbm.fit %>%
  partial(pred.var = 'kw_avg_avg',
          grid.resolution = 100,
          ice = TRUE,
          prob = TRUE) %>%
  autoplot(train = News_Data_train, alpha = 0.02) +
  ggtitle('Boosting, non-centered') +
  theme(axis.text = element_text(size = 6))
ice2.gbm = gbm.fit %>%
  partial(pred.var = 'kw_avg_avg',
          grid.resolution = 100,
          ice = TRUE,
          prob = TRUE) %>%
  autoplot(train = News_Data_train, alpha = 0.02, center = TRUE) +
  ggtitle('Boosting, centered') +
  theme(axis.text = element_text(size = 6))
grid.arrange(ice1.rf, ice2.rf, ice1.gbm, ice2.gbm)
```

Partial Dependence Plots (PDP) of 'self_reference_min_shares'

Individual Conditional Expectation (ICE) of 'self_reference_min_shares'

```
theme(axis.text = element_text(size = 6))
ice2_2.rf = rf.fit %>%
  partial(pred.var = 'self_reference_min_shares',
          grid.resolution = 100,
          ice = TRUE,
          prob = TRUE) %>%
  autoplot(train = News Data train, alpha = 0.02, center = TRUE) +
  ggtitle('Random Forest, centered') +
  theme(axis.text = element_text(size = 6))
ice1_2.gbm = gbm.fit %>%
  partial(pred.var = 'self_reference_min_shares',
          grid.resolution = 100,
          ice = TRUE,
          prob = TRUE) %>%
  autoplot(train = News_Data_train, alpha = 0.02) +
  ggtitle('Boosting, non-centered') +
  theme(axis.text = element_text(size = 6))
ice2_2.gbm = gbm.fit %>%
  partial(pred.var = 'self_reference_min_shares',
          grid.resolution = 100,
          ice = TRUE,
          prob = TRUE) %>%
  autoplot(train = News_Data_train, alpha = 0.02, center = TRUE) +
  ggtitle('Boosting, centered') +
  theme(axis.text = element_text(size = 6))
grid.arrange(ice1_2.rf, ice2_2.rf, ice1_2.gbm, ice2_2.gbm)
```

LIME