## HW 02

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## Question 1

## Read Data

```
df <- read.csv(file = 'gradAdmit.csv', header = T)</pre>
```

```
1(a)
 set.seed(400)
 n = nrow(df)
 # Get 20% for test set
 sample = sample.int(n = n, size = floor(.2*n), replace = F)
 train_wrong_index = df[-sample,]
 test = df[sample,]
 train = train_wrong_index
 rownames(train) = 1:nrow(train_wrong_index)
```

```
1(b)
 # Libraries
 library(e1071)
```

```
library(pROC)
```

```
## Type 'citation("pROC")' for a citation.
```

```
##
## Attaching package: 'pROC'
```

```
## The following objects are masked from 'package:stats':
##
##
       cov, smooth, var
```

```
library(caret)
```

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

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

```
library(hash)
```

```
Cross validation function
```

kernel = "radial",

cost = 1,

nfolds = 5

## hash-2.2.6.2 provided by Decision Patterns

GridSearch\_Custom <- function(nfolds = 5,</pre>

```
degree = 3,
                               gamma = 1/3,
                               coef0 = 0) {
  # Get folds
  folds = createFolds(1:nrow(train),
                      k = nfolds)
  temp\_acc\_list = c()
  temp_auc_list = c()
  for (i in 1:nfolds){
    # Get data for folds
    training = train[-folds[[i]],]
    validation = df[folds[[i]],]
    # Train temp SVM
    svm_temp <- svm(factor(admit) ~ gre + gpa + rank,</pre>
                    data = training,
                    scale = T,
                    probability = T,
                    # default radial
                    kernel = kernel,
                    # cost of constraints violation (default: 1)
                    cost = cost,
                    # needed for: polynomial (default: 3)
                    degree = degree,
                    # needed for all except linear (default: 1/(data dimension))
                    gamma = gamma,
                    # needed for: polynomial, sigmoid (default: 0)
                    coef0 = coef0
          )
    # Test temp SVM
    pred_temp <- predict(svm_temp,</pre>
                         validation,
                          decision.values = F,
                          probability = F)
    # Get temp accuracy and AUC
    acc_temp <- mean(pred_temp==validation$admit)</pre>
    roc_obj <- roc(as.numeric(validation$admit),</pre>
                   as.numeric(pred_temp),
                   levels = c(0, 1),
                   direction = "<")
    auc_temp <- auc(roc_obj)[1]</pre>
    # Print temp values
    # cat("Acc: ", acc_temp, "\n")
    # cat("AUC: ", auc_temp, "\n")
    # print("----")
    \#temp\_acc\_list = append(x = temp\_acc\_list, values = acc\_temp)
    temp_auc_list = append(x = temp_auc_list, values = auc_temp)
  }
  return(mean(temp_auc_list))
}
# Set parameters
```

```
kernels = c('linear', 'polynomial', 'radial', 'sigmoid')
 cost <- c(0.001, 0.1, 1)
 gamma <- c(0.001, 0.1, 1)
 coef0 <- c(1, 10, 100)
 degree <- c(3, 4, 5)
 # Set parameters testing set
 # cost <- c(1)
 # degree <- c(1)
 \# gamma <- c(1)
 # coef0 <- c(1)
 # Init best parameter storage
 best_params = hash()
 best_params[["k"]] <- 'radial'</pre>
 best_params[["c"]] <- 1</pre>
 best_params[["g"]] <- 1/3
 best_params[["c0"]] <- 0</pre>
 best_params[["d"]] <- 3</pre>
 best_params[["auc"]] <- 0</pre>
 for (k in kernels) {
   for (c in cost) {
      if (k == 'linear') {
        auc <- GridSearch_Custom(nfolds = nfolds,</pre>
                                     kernel = k,
                                    cost = c)
        if (auc > best_params[["auc"]]) {
          best_params[["k"]] <- k</pre>
          best_params[["c"]] <- c</pre>
          best_params[["g"]] <- 1/3
          best_params[["c0"]] <- 0</pre>
          best_params[["d"]] <- 3</pre>
          best_params[["auc"]] <- auc</pre>
        }
      } else {
        for (g in gamma) {
          if (k == 'radial') {
             auc <- GridSearch_Custom(nfolds = nfolds,</pre>
                                         kernel = k,
                                         cost = c,
                                         gamma = g)
             if (auc > best_params[["auc"]]) {
               best_params[["k"]] <- k</pre>
               best_params[["c"]] <- c</pre>
               best_params[["g"]] <- g</pre>
               best_params[["c0"]] <- 0</pre>
               best_params[["d"]] <- 3</pre>
               best_params[["auc"]] <- auc</pre>
          } else {
             for (c0 in coef0) {
               if (k == 'sigmoid') {
                 auc <- GridSearch_Custom(nfolds = nfolds,</pre>
                                              kernel = k,
                                              cost = c,
                                              gamma = g,
                                              coef0 = c0)
                 if (auc > best_params[["auc"]]) {
                   best_params[["k"]] <- k</pre>
                   best_params[["c"]] <- c</pre>
                   best_params[["g"]] <- g</pre>
                   best_params[["c0"]] <- c0</pre>
                   best_params[["d"]] <- 3</pre>
                   best_params[["auc"]] <- auc</pre>
               } else {
                 for (d in degree) {
                   auc <- GridSearch_Custom(nfolds = nfolds,</pre>
                                                kernel = k,
                                                cost = c,
                                                gamma = g,
                                                coef0 = c0,
                                                degree = d)
                   if (auc > best_params[["auc"]]) {
                      best_params[["k"]] <- k</pre>
                      best_params[["c"]] <- c</pre>
                      best_params[["g"]] <- g</pre>
                     best_params[["c0"]] <- c0</pre>
                     best_params[["d"]] <- d</pre>
                      best_params[["auc"]] <- auc</pre>
                   }
                 }
              }
            }
          }
        }
      }
   }
 }
The model with kernel polynomial performed the best. The hyper parameters are as follows:
```

- coef0: 1 - degree: 5

```
1(c)
```

- cost: 1 - gamma: 1

```
# Train final SVM
svm_final <- svm(factor(admit) ~ gre + gpa + rank,</pre>
                 data = train,
                 scale = T,
                 probability = T,
                 kernel = best_params[["k"]],
                 cost = best_params[["c"]],
                 degree = best_params[["d"]],
                 gamma = best_params[["g"]],
                 coef0 = best_params[["c0"]]
                 )
# Get final prediction
pred_final <- predict(svm_final,</pre>
                      decision.values = F,
                      probability = F)
# Get accuracy of final model
acc_final <- mean(pred_final==test$admit)</pre>
print(acc_final)
```

```
The final accuracy of my model is 0.6125
```

## [1] 0.6125