KNN Project

This is an R Markdown (http://rmarkdown.rstudio.com) Notebook. When you execute code within the notebook, the results appear beneath the code.

Try executing this chunk by clicking the *Run* button within the chunk or by placing your cursor inside it and pressing *Cmd+Shift+Enter*.

```
# Packages needed to run this R notebook:
    - "class" for KNN algorithm
     - "e1071" for SVM algorithm
    - "rpart" for decision trees algorithm
     - "rpart.plot" for plotting decision tree
     - "randomForest" for random forest algorithm
 # Run this to install and load the necessary packages
ipak <- function(pkg) {</pre>
    new.pkg <- pkg[!(pkg %in% installed.packages()[, "Package"])]</pre>
    if (length(new.pkg))
        install.packages(new.pkg, dependencies = TRUE)
    sapply(pkg, require, character.only = TRUE)
}
packages <- c("class", "e1071", "rpart", "rpart.plot", "randomForest", "stringr")</pre>
ipak (packages)
## Loading required package: class
## Loading required package: e1071
## Loading required package: rpart
## Loading required package: rpart.plot
## Loading required package: randomForest
## randomForest 4.6-14
## Type rfNews() to see new features/changes/bug fixes.
```

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Loading required package: stringr

```
## class e1071 rpart rpart.plot randomForest
## TRUE TRUE TRUE TRUE
## stringr
## TRUE
```

```
## If you would like to calculate the time it takes to execute a chunk of code, pleas
e use the code below
sleep_for_a_minute <- function() { Sys.sleep(0) }
start_time <- Sys.time()
sleep_for_a_minute()
# Insert code you would like to measure
end_time <- Sys.time()
time <- end_time - start_time
time # Duration of the execution in seconds</pre>
```

```
## Time difference of 0.004997969 secs
```

```
# The first three chunks are to read the observations
# TRAINING SET
# First, we get the list of all files in the directory

list_files <- list.files(path = "digits/trainingDigits",all.files = TRUE) # We get the list of all files
paste0("There are ",length(list_files)," files in the directory")</pre>
```

```
## [1] "There are 1936 files in the directory"
```

```
list_files <- list_files[3:1936] # We remove the first two (they indicate the folders
)

list_files_path <- list() # We create a list with the relative filepath of files
for (item in list_files){
  name <- paste("digits/trainingDigits/",item,sep = "") #This is the relative path. D
epending on your operating system, you might need to enter the full path
  list_files_path[item] <- name
}
length(list_files_path) == length(list_files)</pre>
```

```
## [1] TRUE
```

[1] TRUE

length(test list files)

```
# TRAINING CLASS (Y)
# Here we create a list with the class of each object (the first number of each file
represents the class)

train_Y <- c()

for (f in list_files) {
    first_char <- substr(f, 1, 1)
        train_Y[f] <- first_char
}

train_Y <- Reduce('rbind', train_Y)
train_Y <- c(train_Y)
train_Y <- as.data.frame(train_Y)

train_Y = train_Y[,1]</pre>
```

```
# OBSERVATIONS (X)
# Now, we will read through the list of training files and put them on a list. Number
of observations: 1934

train_num_list <- list()

for (i in list_files_path) {
   read_txt <- scan(i,what="character", sep=NULL) #We read the txt as characters, not
   as numbers
   oneline <- as.numeric(stringr::str_split(paste(read_txt, collapse = ""), "")[[1]])
# We convert 32x32 to 1x1024
   train_num_list[[i]] <- oneline
}
print(length(train_num_list) == length(train_Y)) # We make sure that the number of ob
servations and the length of the classes match</pre>
```

```
# TEST SET
# First, we get the list of all files in the directory
test list files <- list.files(path = "digits/testDigits",all.files = TRUE)</pre>
```

```
## [1] 948
```

```
test_list_files <- test_list_files[3:length(test_list_files)]

test_list_files_path <- list()  # We create a list with the relative filepath of file
s
for (item in test_list_files){
  name <- paste("digits/testDigits/",item,sep = "")  #This is the relative path. Depen
ding on your operating system, you might need to enter the full path
  test_list_files_path[item] <- name
}

# name <- paste("digits/testDigits/",i,sep = "")
# print(test_list_files_path[946])</pre>
```

```
# TEST CLASS (test_y)
# Here we create a list with the class of each object (the number each file represent
s)

test_y <- list()

for (f in test_list_files) {
    first_char <- substr(f, 1, 1)
    test_y[f] <- first_char
}

test_y <- Reduce('rbind', test_y)
test_y <- c(test_y)
test_y <- as.data.frame(test_y)

test_y = as.data.frame(test_y[,1])

length(test_y) == length(test_list_files_path)</pre>
```

```
## [1] FALSE
```

library(class)

knn df <- as.data.frame(knn1)</pre>

knn1 <- knn(train=X train, test=X test, cl=train Y, k=3)</pre>

```
# OBSERVATIONS (test_x)
# Now, we will read through the list of training files and put them on a list. Number
of observations: 1934
test num list <- list()</pre>
for (i in test list files path) {
  read test txt <- scan(i, what="character", sep=NULL) #We read the txt as characters,
not as numbers
 oneline <- as.numeric(stringr::str split(paste(read test txt, collapse = ""), "")[[</pre>
1]]) # We convert 32x32 to 1x1024
 test num list[[i]] <- oneline</pre>
\verb|print(length(test_num_list)| == \ length(test_y)) \ \# \ \textit{We make sure that the number of obse}
rvations and the length of the classes match
## [1] FALSE
# We get the X matrix for our training data as a data frame
X_train <- as.data.frame(do.call(rbind, unname(train num list)))</pre>
# We get the X of our test data that we would like to predict, as a data frame
X test <- as.data.frame(do.call(rbind, unname(test num list)))</pre>
# KNN algorithm and we convert to a list with predictions
```

```
# We have to compare knn outcome with test_y, so we know how precise the knn is

results <- cbind.data.frame(knn_df[,1], test_y[,1])

count_right <- 0

# We run through all the predicted values and
# compare them with the real values. If correct, we add 1 to count_right

for (row in 1:nrow(results)) {
   knn_pred <- results[row,1]
   test_obs <- results[row,2]
   if (knn_pred == test_obs) {
      count_right <- count_right + 1
   }
   }
   count_right</pre>
```

```
## [1] 932
```

```
knn_accuracy <- count_right/nrow(results)
knn_accuracy</pre>
```

```
## [1] 0.9852008
```

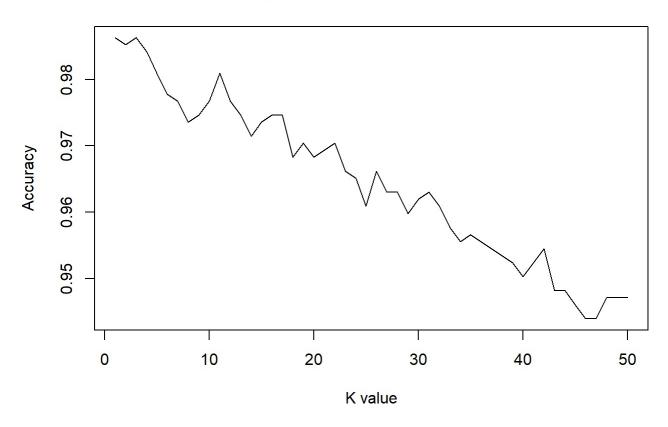
We create a confusion matrix with the predicted values and real values.
confMat <- table(data=results[,1], reference=results[,2])
confMat</pre>

```
##
   reference
    0 1 2
           3 4 5 6 7 8
## data
##
   0 87
      0 0 0 0 0 0 0 0
   1 0 96 0 0 0 0 0 3 1
##
   2 0 0 92 0 0 0 0 0 0
##
   3 0 0 0 84 0 1 0 0 1 1
##
##
  4 0 0 0 0 114 0 0 0 0
  5 0 0 0 0 0 105 0 0 0 1
##
   6 0 0 0 0 0 2 87 0 1 0
##
  7 0 1 0 0 0 0 96 0
##
                         1
##
  8 0 0 0 0 0 0 0 86
                         0
  9 0 0 0 1 0 0 0 0 85
##
```

```
# We compare the accuracy of the different values of k. It will take several minutes
to compute
library(class)
accuracy list <- c()</pre>
for (k in 1:50) {
  knn2 <- knn(train=X train, test=X test, cl=train Y, k, use.all=FALSE)
  knn df2 <- as.data.frame(knn2)</pre>
  results2 <- cbind.data.frame(knn_df2[,1],test_y[,1])</pre>
  count right2 <- 0
  for (row in 1:nrow(results2)){
   knn pred1 <- results2[row,1]</pre>
   test obs1 <- results2[row,2]</pre>
    if (knn pred1 == test obs1) {
      count right2 <- count right2 +1</pre>
  }
  accuracy <- count right2/nrow(results2)</pre>
  accuracy list[k] <- accuracy</pre>
print(accuracy list)
## [1] 0.9862579 0.9852008 0.9862579 0.9841438 0.9809725 0.9778013 0.9767442
## [8] 0.9735729 0.9746300 0.9767442 0.9809725 0.9767442 0.9746300 0.9714588
## [15] 0.9735729 0.9746300 0.9746300 0.9682875 0.9704017 0.9682875 0.9693446
## [22] 0.9704017 0.9661734 0.9651163 0.9608879 0.9661734 0.9630021 0.9630021
## [29] 0.9598309 0.9619450 0.9630021 0.9608879 0.9577167 0.9556025 0.9566596
## [36] 0.9556025 0.9545455 0.9534884 0.9524313 0.9503171 0.9524313 0.9545455
## [43] 0.9482030 0.9482030 0.9460888 0.9439746 0.9439746 0.9471459 0.9471459
## [50] 0.9471459
print(summary(accuracy list))
     Min. 1st Qu. Median Mean 3rd Qu.
## 0.9440 0.9545 0.9641 0.9644 0.9746 0.9863
# We see above that the optimal K value is 1 which gives us an accuracy of nearly 99%
# We plot the accuracy of the different k values
plot(accuracy_list[1:50], ylab="Accuracy", xlab="K value", type="1", main="Accuracy o
```

f KNN algorithm by K values")

Accuracy of KNN algorithm by K values



Now we will apply the Support Vector Machine (SVR) for classification and compare the results

```
library(e1071)
svm_model <- svm(train_Y ~ ., data=X_train, scale=FALSE)
summary(svm_model)</pre>
```

```
##
## Call:
## svm(formula = train Y ~ ., data = X train, scale = FALSE)
##
##
## Parameters:
##
     SVM-Type: C-classification
## SVM-Kernel: radial
         cost: 1
##
##
       gamma: 0.0009765625
## Number of Support Vectors: 1139
##
   ( 78 130 101 124 109 122 87 93 135 160 )
##
##
##
## Number of Classes: 10
##
## Levels:
## 0 1 2 3 4 5 6 7 8 9
```

```
# We get the predicted values for the test data
pred_svm <- predict(svm_model, X_test)
summary(pred_svm)</pre>
```

```
## 0 1 2 3 4 5 6 7 8 9
## 88 99 87 80 111 108 87 97 96 93
```

```
# We calculate the accuracy of the algorithm
rightguess <- 0

pred <- as.data.frame(pred_svm)

results_svm <- cbind.data.frame(pred,test_y)

results_svm <- as.data.frame(results_svm)

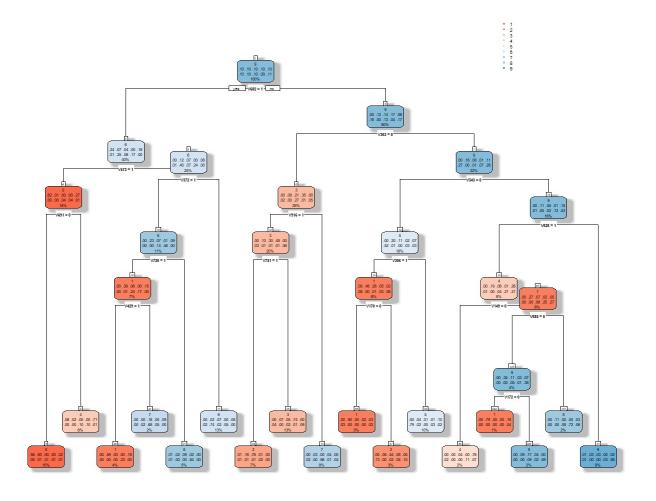
for (row in (1:nrow(results_svm))) {
   svm_pred <- results_svm[row,1]
   svm_test <- results_svm[row,2]
   if (svm_pred==svm_test) {
      rightguess <- rightguess+1
   }
   }
}

accuracy_svm <- rightguess/946
accuracy_svm</pre>
```

```
## [1] 0.9735729
confMat <- table(data=results_svm[,1], reference=results_svm[,2])</pre>
confMat
##
     reference
## data 0 1 2 3 4 5 6 7 8 9
## 0 87 0 0 0 1 0 0 0 0
    1 0 96 0 0 1 0 0 0 1 1
##
   2 0 0 87 0 0 0 0 0 0
##
   3 0 0 0 79 0 0 0 0 1
##
   4 0 0 0 0 109 0 1 1 0 0
##
## 5 0 0 0 1 0 107 0 0 0
## 6 0 0 0 0 0 0 86 0 1 0
## 7 0 1 0 0 0 0 95 0 1
## 8 0 0 4 1 2 0 0 0 89 0
## 9 0 0 1 4 1 1 0 0 86
summary(confMat)
## Number of cases in table: 946
## Number of factors: 2
## Test for independence of all factors:
## Chisq = 8027, df = 81, p-value = 0
# We try now the Decission Tree algorithm
library(rpart)
library(rpart.plot)
tree model <- rpart(train Y ~ ., method="class", data=X train)</pre>
```

```
plot <- rpart.plot(tree_model, box.palette="RdBu", shadow.col="gray", nn=TRUE)</pre>
```

confMat



```
# We calculate the accuracy of the Decission Tree predictions

prediction_dt <- predict(object=tree_model,newdata=X_test,type="class")

results_dt <- cbind.data.frame(prediction_dt,test_y)
rightcount_dt <- 0

for (item in 1:nrow(results_dt)) {
   dt_pred <- results_dt[item,1]
   dt_true <- results_dt[item,2]
   if (dt_pred==dt_true) {
      rightcount_dt = rightcount_dt+1
   }
}

accuracy_dt <- rightcount_dt/nrow(results_dt)
accuracy_dt</pre>
```

```
## [1] 0.7547569

confMat <- table(data=results_dt[,1], reference=results_dt[,2])</pre>
```

```
##
    reference
## data 0 1 2 3 4 5 6 7 8 9
   082 0 0 0 1 0 0 2 0 0
   1 0 47 0 0 14 2 0 0 19 2
   2 0 25 60 4 0 4 0 0 6 2
##
##
   3 0 10 4 74 1 2 0 1 1 6
   4 1 1 0 0 70 0 1 5 2 1
##
##
   5 0 2 3 1 9 89 2 0 0 1
   6 2 2 5 0 14 1 84 0 8 0
##
   7 0 3 9 2 2 7 0 86 1 4
##
   8 2 6 6 0 0 1 0 1 49 0
##
   9 0 1 5 4 3 2 0 1 5 73
```

```
# We train the Random Forest algorithm
library(randomForest)
number_of_trees <- 200  # I tried with 200, change this value to try different tree n
umbers
rf_model <- randomForest(data=X_train, y=train_Y, x=X_train, ntree=number_of_trees)
prediction_rf <- predict(object=rf_model, newdata=X_test, type="class")</pre>
```

```
# We calculate the accuracy of Random Forest
results_rf <- cbind.data.frame(prediction_rf,test_y)

rightcount_rf <- 0
for (item in 1:nrow(results_rf)) {
    rf_pred <- results_rf[item,1]
    rf_true <- results_rf[item,2]
    if (rf_pred==rf_true) {
        rightcount_rf = rightcount_rf+1
    }
}

accuracy_rf <- rightcount_rf/nrow(results_rf)
accuracy_rf</pre>
```

```
## [1] 0.9809725
```

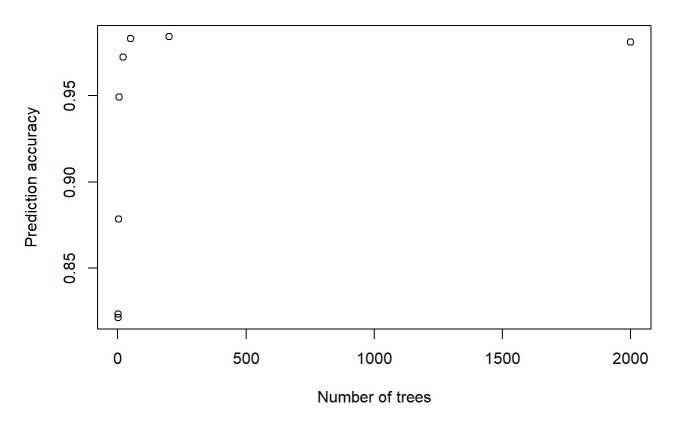
```
# Accuracy of random forest for different number of trees
accuracy_rf_list <- c()</pre>
x \leftarrow c(1,2,3,5,20,50,200,2000)
for (i in 1:length(x)){
  rf model <- randomForest(data=X train, y=train Y, x=X train, ntree=x[i])</pre>
  prediction rf <- predict(object=rf model, newdata=X test, type="class")</pre>
  results rf <- cbind.data.frame(prediction rf, test y)</pre>
  rightcount rf <- 0
  for (item in 1:nrow(results rf)){
   rf_pred <- results_rf[item,1]</pre>
   rf true <- results rf[item,2]</pre>
    if (rf_pred==rf_true) {
      rightcount rf = rightcount rf+1
    }
  }
  accuracy rf <- rightcount rf/nrow(results rf)</pre>
  accuracy_rf_list[i] = accuracy_rf
```

```
accuracy_rf_list
```

```
## [1] 0.8213531 0.8234672 0.8784355 0.9492600 0.9725159 0.9830867 0.9841438
## [8] 0.9809725
```

```
\label{lem:plot} $$\operatorname{plot}(y=\operatorname{accuracy\_rf\_list},\ x=x,\ xlab="Number of trees",\ ylab="Prediction accuracy",\ main="Random Forest Accuracy")$
```

Random Forest Accuracy



```
# Naive Bayes algorithm
nb_model <- naiveBayes(x=X_train, y=train_Y, laplace = 0.01)</pre>
```

```
# Predictions of new data and we calculate accuracy

prediction_nb <- predict(object=nb_model, newdata=X_test,
    type = c("class"))

prediction_rf <- predict(object=rf_model, newdata=X_test, type="class")

results_nb <- cbind.data.frame(prediction_nb,test_y)

rightcount_nb <- 0

for (item in 1:nrow(results_nb)) {
    nb_pred <- results_nb[item,1]
    nb_true <- results_nb[item,2]
    if (nb_pred==nb_true) {
        rightcount_nb = rightcount_nb+1
    }
}

accuracy_nb <- rightcount_nb/nrow(results_nb)
accuracy_nb</pre>
```

```
## [1] 0.8657505
confMat <- table(data=results_nb[,1], reference=results_nb[,2])</pre>
confMat
##
     reference
## data 0 1 2 3 4 5 6 7 8 9
   086000100000
##
    1 0 72 0 0 1 0 0 0 1
   2 0 2 81 0 0 0 0 0 0
##
   3 0 0 1 73 0 4 0 0 0 14
##
##
   4 0 0 0 0 92 0 0 0 1
   5 0 0 0 0 0 93 0 0 0 0
##
   6 0 1 0 0 4 1 86 0 1 0
##
##
   7 0 0 0 1 1 0 0 93 0 2
## 8 1 20 10 9 13 7 1 3 90 18
   9 0 2 0 2 2 3 0 0 0 53
##
```