KNN Model

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```
# import datasets
wine <- read.csv("C:/Users/xinya/Downloads/Cornell Classes/STSCI 3740/final project/wine-quality-white-</pre>
```

Fitting a KNN Model

mean(knn.pred != testing\$quality)

})

print(knn.errors)

```
# normalize the data using z-score
normalize <- function(x) {</pre>
  return((x - mean(x)) / sd(x))
}
all_columns <- names(wine)</pre>
columns_to_normalize <- all_columns[all_columns != "quality" & sapply(wine, is.numeric)]</pre>
wine norm <- wine
wine_norm[columns_to_normalize] <- lapply(wine[columns_to_normalize], normalize)</pre>
# change type of wine to white=1, red=2
wine_norm$type <- as.numeric(factor(wine_norm$type))</pre>
\# split the dataset into train/test
set.seed(1)
index <- sample(1:nrow(wine_norm), size=nrow(wine_norm)*0.7, rep=FALSE)</pre>
training <- wine_norm[index, ]</pre>
testing <- wine_norm[-index, ]</pre>
training_X <- training %>% select(-quality)
testing_X <- testing %>% select(-quality)
# try different values of k from 1 to 20
k.values <- 1:20
knn.errors <- sapply(k.values, function(k) {</pre>
  knn.pred <- knn(training_X, testing_X, training$quality, k=k)</pre>
```

[1] 0.3989744 0.4794872 0.4707692 0.4574359 0.4482051 0.4456410 0.4507692

```
## [8] 0.4528205 0.4451282 0.4528205 0.4364103 0.4420513 0.4317949 0.4379487
## [15] 0.4358974 0.4358974 0.4461538 0.4415385 0.4389744 0.4405128
```

The value of k that seems to perform the best on this data is k=1.

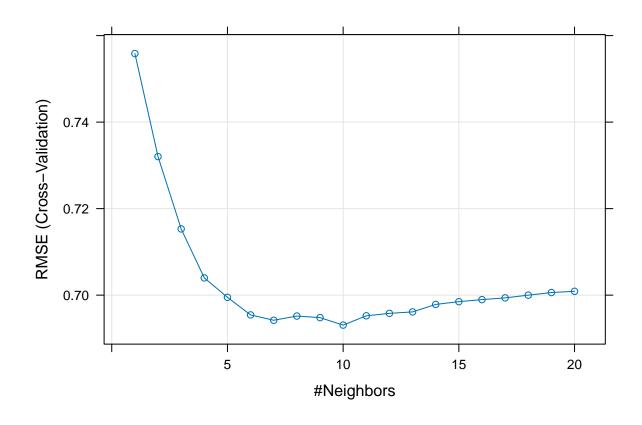
Choose the optimal k-value using Cross Validation

```
set.seed(1)
# 10-fold cross validation
control <- trainControl(method = "cv", number = 10)</pre>
knn_cv <- train(
 quality ~ .,
 data = wine_norm,
 method = "knn",
 trControl = control,
  tuneGrid = expand.grid(k = 1:20)
knn cv
## k-Nearest Neighbors
##
## 6497 samples
##
     12 predictor
##
## No pre-processing
## Resampling: Cross-Validated (10 fold)
## Summary of sample sizes: 5847, 5846, 5847, 5847, 5847, 5848, ...
## Resampling results across tuning parameters:
##
```

```
##
    k
        RMSE
                  Rsquared
                             MAE
##
     1 0.7558513 0.3795189 0.4239693
     2 0.7320306 0.3612061 0.4939238
##
##
     3 0.7153067 0.3654421 0.5125768
##
     4 0.7039897 0.3718375 0.5170525
##
     5 0.6994976 0.3727748 0.5246780
##
     6 0.6954288 0.3756623 0.5260905
##
     7 0.6941926 0.3755005 0.5286038
##
     8 0.6951634 0.3725416 0.5314023
##
     9 0.6948018 0.3720174 0.5344289
##
    10 0.6930709 0.3743660 0.5342596
##
    11 0.6952228 0.3699285 0.5365115
##
    12 0.6957883 0.3683444
                            0.5382253
##
    13 0.6961228 0.3676329 0.5401778
##
    14 0.6978546 0.3643808 0.5417053
##
    15 0.6984920 0.3632550 0.5426139
##
    16 0.6989681 0.3623560 0.5431027
##
    17 0.6993684 0.3614560 0.5446430
##
    18 0.6999960 0.3602518 0.5456382
##
    19 0.7005973 0.3591466 0.5467203
```

```
## 20 0.7008890 0.3587130 0.5473792 ## ## RMSE was used to select the optimal model using the smallest value. ## The final value used for the model was k = 10.
```

```
plot(knn_cv)
```



Fitting the best model

```
model <- knn(training_X, testing_X, training$quality, k=10)

confusion_matrix <- table(coPredicted = model, Actual = testing$quality)
print(confusion_matrix)</pre>
```

```
##
                Actual
##
   coPredicted
                    3
                             5
##
                    0
                         0
                             0
                                  0
                                                0
                         2
                             5
##
                    0
                                  0
                       32 392 186
##
               5
                    4
                                     16
##
               6
                       33 219 547 155
               7
                    0
                            19 123
                                    144
                                          31
                                                1
##
##
               8
                    0
                         0
                             0
                                  5
                                       2
                                            2
                                                1
               9
                             0
                                  0
                                       0
                                            0
                                                0
##
                    0
                         0
```

```
accuracy <- sum(diag(confusion_matrix)) / sum(confusion_matrix)</pre>
accuracy
## [1] 0.5574359
```

Apply PCA to reduce diensionality

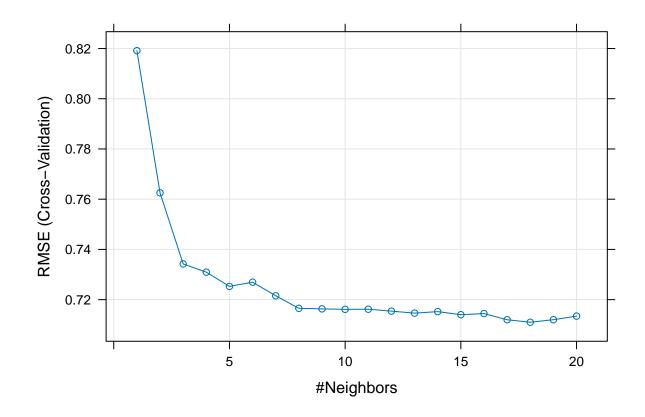
```
# apply PCA before fitting KNN
new_wine <- wine</pre>
new_wine$type <- as.numeric(factor(new_wine$type))</pre>
target <- new_wine$quality</pre>
predictors <- new_wine %>% select(-quality)
# standardize data
scaled_data <- scale(predictors)</pre>
# Perform PCA
pca_result <- prcomp(scaled_data, center = TRUE, scale. = TRUE)</pre>
summary(pca_result)
## Importance of components:
##
                              PC1
                                     PC2
                                             PC3
                                                    PC4
                                                            PC5
                                                                     PC6
                                                                            PC7
## Standard deviation
                           1.9518 1.5902 1.2496 0.9853 0.85077 0.78329 0.7324
## Proportion of Variance 0.3175 0.2107 0.1301 0.0809 0.06032 0.05113 0.0447
## Cumulative Proportion 0.3175 0.5282 0.6583 0.7392 0.79952 0.85065 0.8953
                               PC8
                                       PC9
                                             PC10
                                                      PC11
                                                              PC12
## Standard deviation
                          0.70921 0.59368 0.5068 0.34552 0.15539
## Proportion of Variance 0.04192 0.02937 0.0214 0.00995 0.00201
## Cumulative Proportion 0.93727 0.96664 0.9880 0.99799 1.00000
Decided to choose the first eight PCs
# reduce dimensionality
# choose the first eight PCs
```

```
pca_data <- pca_result$x[, 1:8]</pre>
```

Fit KNN Model on PCA-reduced data

```
# apply KNN on PCA-reduced data
set.seed(1)
index_2 <- sample(1:nrow(pca_data), size=nrow(pca_data)*0.7, rep=FALSE)</pre>
training_data <- pca_data[index_2, ]</pre>
testing_data <- pca_data[-index_2, ]</pre>
training_targt <- target[index_2]</pre>
testing_target <- target[-index_2]</pre>
```

```
# 10-fold cross validation to find the best k
control_new <- trainControl(method = "cv", number = 10)</pre>
knn cv 2 <- train(
 training_data,
 training_targt,
 method = "knn",
 trControl = control_new,
 tuneGrid = expand.grid(k = 1:20)
knn_cv_2
## k-Nearest Neighbors
##
## 4547 samples
     8 predictor
##
##
## No pre-processing
## Resampling: Cross-Validated (10 fold)
## Summary of sample sizes: 4091, 4091, 4093, 4092, 4092, 4091, ...
## Resampling results across tuning parameters:
##
##
        RMSE
                   Rsquared
    k
                              MAE
##
     1 0.8191349 0.2974479 0.4807663
##
     2 0.7625294 0.3141066 0.5264729
##
     3 0.7342127 0.3306869 0.5353884
##
     4 0.7309495 0.3219460 0.5462665
##
     5 0.7252672 0.3256003 0.5506567
##
     6 0.7269512 0.3199774 0.5561480
##
     7 0.7215278 0.3253284 0.5557852
##
     8 0.7165359 0.3322161 0.5540389
##
     9 0.7163207 0.3311995 0.5561248
    10 0.7161339 0.3301913 0.5570248
##
##
    11 0.7161817 0.3296976 0.5592316
##
    12 0.7154242 0.3309020 0.5601571
##
    13 0.7146273 0.3320483 0.5608825
    14 0.7152282 0.3306918 0.5617482
##
##
    15 0.7140067 0.3327959 0.5607729
##
    16 0.7144465 0.3320431 0.5621654
    17 0.7119925 0.3367552 0.5608877
##
##
    18 0.7109959 0.3386110
                              0.5605507
##
    19 0.7120115 0.3367728 0.5614468
##
    20 0.7134455 0.3339659 0.5633644
##
## RMSE was used to select the optimal model using the smallest value.
## The final value used for the model was k = 18.
plot(knn_cv_2)
```



```
knn_model <- knn(train = training_data, test = testing_data, cl = training_targt, k = 17)
confusion_matrix <- table(Predicted = knn_model, Actual = testing_target)
print(confusion_matrix)</pre>
```

```
##
              Actual
                 3
## Predicted
                      4
                          5
                               6
                                        8
                                             9
##
            3
                 0
                      0
                          0
                                             0
##
             4
                 0
                      1
                          2
                                    0
                                             0
            5
                 5
##
                    34 367 200
                                  14
                                             0
            6
                 3
                    30 247
                                             2
##
                             566 176
                                       25
            7
##
                 0
                      2
                         19
                              94
                                 126
                                       31
                                             1
            8
##
                 0
                      0
                          0
                               0
                                        0
                                             0
                          0
                               0
                                    0
                                        0
                                             0
```

```
accuracy <- sum(diag(confusion_matrix)) / sum(confusion_matrix)
accuracy</pre>
```

[1] 0.5435897