Final Project

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Contents

```
#Data Read-in & Clean-up
##Training set
```

First, read in all the train files. Each files_{name}[i] will display a file route directed to one photo.

```
##setup code
files_buildings = mixedsort(
  sort(
    list.files("~/Data Science/Project/seg_train/buildings",
               pattern = "*.jpg", all.files = T, full.names = T, no.. = T, )))
files_forest = mixedsort(
  sort(
    list.files("~/Data Science/Project/seg train/forest",
               pattern = "*.jpg", all.files = T, full.names = T, no.. = T)))
files_glacier = mixedsort(
  sort(
    list.files("~/Data Science/Project/seg_train/glacier",
               pattern = "*.jpg", all.files = T, full.names = T, no.. = T)))
files_mountain = mixedsort(
  sort(
    list.files("~/Data Science/Project/seg_train/mountain",
               pattern = "*.jpg", all.files = T, full.names = T, no.. = T)))
files_sea = mixedsort(
  sort(
    list.files("~/Data Science/Project/seg_train/sea",
               pattern = "*.jpg", all.files = T, full.names = T, no.. = T)))
files_street = mixedsort(
  sort(
    list.files("~/Data Science/Project/seg_train/street",
               pattern = "*.jpg", all.files = T, full.names = T, no.. = T)))
```

Since our laptops are unable to handle all the photos, we decided to randomly select 200 photos from each category's training set.

```
##randomly selected 200 photos from each category (with set.seed(1)).
set.seed(1)
f_buildings = sample(files_buildings, 200)
f_forest = sample(files_forest, 200)
f_glacier = sample(files_glacier, 200)
```

```
f_mountain = sample(files_mountain, 200)
f_sea = sample(files_sea, 200)
f street = sample(files street, 200)
##combine all six files and load the image
files = c(f_buildings, f_forest, f_glacier, f_mountain, f_sea, f_street)
image list = lapply(files, load.image)
image_list[[1]] #what the info of the first image looks like
## Image. Width: 150 pix Height: 150 pix Depth: 1 Colour channels: 3
##get the classname from the directory
class_list = basename(dirname(files))
table(class_list)
## class_list
## buildings
               forest glacier mountain sea
                                                      street
##
        200
                200
                            200
                                     200
                                                200
                                                         200
remove(f_buildings, f_forest, f_glacier, f_mountain, f_sea, f_street,
      files_buildings, files_forest, files_glacier,
      files_mountain, files_sea, files_street, files)
```

For each i in image_list[[i]], image_list[[i]] contains info of one image. To expand these info/pixels, we need to transform each of them to a vector. Use img_matrix to set up the empty matrix with numbers of columns to be 67500 (150*150). For each image, the value of each pixel will be stored in each of the columns.

Use badimage to count for all images that are not 150*150 pixels, and these images will be discarded from further analyses.

Use the for loop to change each list to a vector. Details in comments. Use img_df to create a new data frame from the matrix.

```
#build the structure of an empty matrix
img_matrix = matrix(ncol = 67500, nrow = length(image_list))
badimage = NULL

for(i in 1:length(image_list)){
   if(length(as.vector(image_list[[i]])) == 67500) {
      img_matrix[i,] = c(as.vector(image_list[[i]]))
      #fill in each row with one observation's pixels
      #do this for each observation unless photo does not meet requirement
}
else{badimage = append(badimage, i)}
#if image does not meet requirement, record which i it is
#so when appending class to each observation,
#will use this badimage to get rid of the class of those not met the requirement
}
img_df = as.data.frame(img_matrix[-badimage,])
```

```
print(dim(img_df)) #dimension of the training set
## [1] 1195 67500
```

The number of rows is not exactly 1200 because there are several images that do not meet the 150*150 size criteria, so they were discarded from further analysis.

Add class assignment to the data frame

```
#append class and remove excessive values
new_class = class_list[-badimage]
img_df$class = as.factor(new_class)

remove(img_matrix)
```

##Test Data

The procedure should be similar.

```
#everything the same for test data
files_buildings = mixedsort(
  sort(
   list.files("~/Data Science/Project/seg_test/buildings",
               pattern = "*.jpg", all.files = T, full.names = T, no.. = T)))
files_forest = mixedsort(
  sort(
    list.files("~/Data Science/Project/seg_test/forest",
               pattern = "*.jpg", all.files = T, full.names = T, no.. = T)))
files_glacier = mixedsort(
  sort(
   list.files("~/Data Science/Project/seg_test/glacier",
               pattern = "*.jpg", all.files = T, full.names = T, no.. = T)))
files_mountain = mixedsort(
  sort(
   list.files("~/Data Science/Project/seg_test/mountain",
               pattern = "*.jpg", all.files = T, full.names = T, no.. = T)))
files_sea = mixedsort(
  sort(
   list.files("~/Data Science/Project/seg_test/sea",
               pattern = "*.jpg", all.files = T, full.names = T, no.. = T)))
files street = mixedsort(
  sort(
    list.files("~/Data Science/Project/seg_test/street",
               pattern = "*.jpg", all.files = T, full.names = T, no.. = T)))
f_buildings = sample(files_buildings, 200)
f_forest = sample(files_forest, 200)
f_glacier = sample(files_glacier, 200)
f_mountain = sample(files_mountain, 200)
f_sea = sample(files_sea, 200)
f_street = sample(files_street, 200)
```

```
files = c(f_buildings, f_forest, f_glacier, f_mountain, f_sea, f_street)
image_list = lapply(files, load.image)
##class
class_list = basename(dirname(files))
remove(f_buildings, f_forest, f_glacier, f_mountain, f_sea, f_street,
       files_buildings, files_forest, files_glacier,
       files_mountain, files_sea, files_street, files)
img_matrix = matrix(ncol = 67500, nrow = length(image_list))
badimage = NULL
for(i in 1:length(image list)){
  if(length(as.vector(image list[[i]])) == 67500) {
    img_matrix[i,] = c(as.vector(image_list[[i]]))
    #fill in each row with one observation's pixels
    #do this for each observation unless photo does not meet requirement
 }
 else{badimage = append(badimage, i)}
  #if image does not meet requirement, record which i it is
 #so when appending class to each observation,
  #will use this badimage to get rid of the class of those not met the requirement
test_df = as.data.frame(img_matrix[-badimage,])
new class = class list[-badimage]
test_df$class = as.factor(new_class)
remove(img_matrix, image_list)
#PCA
##PCA on the training set
Run the prcomp command to construct the PCA
colclass = ncol(img_df)
img_pca = prcomp(img_df[, -colclass], center = T, scale = T)
```

See how much variance each PCs explains, so first by computing the cumulative sum of variance explained, and output the number of PCs that expalins 75, 80, and 90 percent of total variance. THe img_cumsum contians the variance each pc explains. Each pc?? is the index that contains variance closet to 75, 80, or 90 percent.

```
#get the cumulative variance explained
pr.var = img_pca$sdev^2
pve=pr.var/sum(pr.var)

img_cumsum = cumsum(pve)
img_cumsum[1:10]
```

```
## [1] 0.2152665 0.3474630 0.4010042 0.4335586 0.4630449 0.4867607 0.5085308
## [8] 0.5231124 0.5364468 0.5476634

pc75 = which.min(abs(img_cumsum - 0.75))
pc80 = which.min(abs(img_cumsum - 0.80))
pc90 = which.min(abs(img_cumsum - 0.90))
pc75
## [1] 99
pc80
## [1] 170
pc90
## [1] 422
```

Build the training set data frame based on PCA results

```
#build dataframe
df75 = data.frame(img_df$class, img_pca$x[,1:pc75])
colnames(df75)[1] = "class"

print(dim(df75)) #dimension on the train dataset
## [1] 1195 100
```

##On testing set

```
#project the pcs onto the test data
pcatest = predict(img_pca, newdata = test_df)
pcatest = as.data.frame(pcatest)

#format the test data
test75 = pcatest[, 1:pc75]
test75 = data.frame(test_df$class, test75)
colnames(test75)[1] = "class"
dim(test75) # dimension of the new test data
## [1] 1195 100
```

Therefore, we now have the projected data frame

#kNN

Normalize the variables

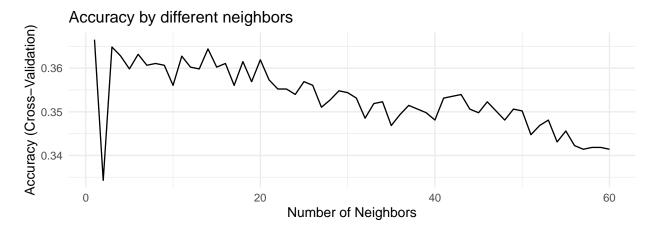
```
#normalizing function
normalize <- function(x) {
return ((as.numeric(x) - min(as.numeric(x))) / (max(as.numeric(x)) - min(as.numeric(x)))) }
#normalize the train dataset followed by the test dataset
normdf = apply(df75[, -1], 2, normalize)
normdf = data.frame(df75$class, normdf)
colnames(normdf)[1] = "class"
normdf$class = as.factor(normdf$class)
normtest = apply(test75[,-1], 2, normalize)</pre>
```

```
normtest = data.frame(test75$class, normtest)
colnames(normtest)[1] = "class"
```

Use 10-fold CV to assess the optimal number of neighbors on the entire data (training + testing)

The variable **kuse** is used to store the number of k that provides the highest accuracy. The following plot show the different accuracy by different number of neighbors.

```
ggplot() +
  geom_line(aes(x = 1:60, y = knnfit$results[,2])) +
  labs(x = "Number of Neighbors", y = "Accuracy (Cross-Validation)",
        title = "Accuracy by different neighbors") +
  theme_minimal()
```



Now perform knn.

```
knntrain = knn3Train(normdf[,-1], normtest[,-1], cl = normdf[,1], k = kuse, prob = T)
knntab = table(knntrain, normtest$class)
kable(knntab)
```

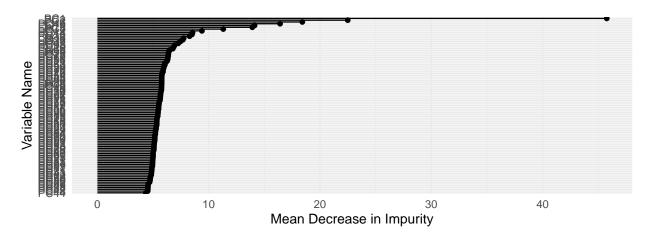
	buildings	forest	glacier	mountain	sea	street
buildings	16	0	1	1	3	15
forest	56	145	14	10	28	69
glacier	28	6	65	22	35	22
mountain	56	31	57	105	55	30
sea	31	7	41	44	61	19
street	13	11	19	16	18	45

```
Accuracy:
sum(diag(knntab))/sum(knntab)
## [1] 0.3656904
#Random Forest
hyper_grid = expand.grid(
       mtry=c(1:15),
        node_size=c(1:15),
        sampe_size=c(0.55, 0.632, 0.7, 0.8),
        00B_err=0
start.time = Sys.time()
for(i in 1:nrow(hyper_grid)) {
       ranger_rf = ranger(
                formula=class~.,
                data=df75,
                num.trees=2000,
                mtry=hyper_grid$mtry[i],
                min.node.size=hyper_grid$node_size[i],
                sample.fraction=hyper_grid$sampe_size[i],
                seed=1
        )
       hyper_grid$00B_err[i] = ranger_rf$prediction.error
end.time = Sys.time()
time.taken = end.time - start.time
time.taken
## Time difference of 1.663089 hours
hyper_grid %>%
dplyr::arrange(00B_err) %>%
head(10)
##
     mtry node_size sampe_size OOB_err
## 1
      10
                  7
                        0.800 0.4276151
## 2
                  15
                        0.550 0.4284519
        11
## 3
        7
                  1
                        0.550 0.4292887
                         0.632 0.4292887
## 4
        11
                  4
                  10
                          0.632 0.4309623
## 5
        11
```

```
## 6 7
                  4 0.700 0.4309623
## 7
        9
                  9
                        0.632 0.4317992
## 8
       11
                 14
                       0.632 0.4317992
## 9
       5
                       0.700 0.4317992
                  5
                  3
## 10
       14
                         0.800 0.4317992
which.min(hyper_grid$00B_err) # 775
## [1] 775
hyper_grid[775,]
      mtry node_size sampe_size OOB_err
## 775 10
                  7
                          0.8 0.4276151
ranger_rf = ranger(
               formula=class~.,
               data=df75,
               num.trees=2000,
               mtry=10,
               min.node.size=7,
               sample.fraction=0.8,
               seed=1,
               importance='impurity'
ranger_rf
## Ranger result
##
## Call:
## ranger(formula = class ~ ., data = df75, num.trees = 2000, mtry = 10, min.node.size =
##
## Type:
                                   Classification
## Number of trees:
                                   2000
                                   1195
## Sample size:
## Number of independent variables: 99
                                   10
## Mtry:
## Target node size:
## Variable importance mode:
                                   impurity
## Splitrule:
                                   gini
## 00B prediction error:
                                   42.76 %
ranger_rf$confusion.matrix
##
             predicted
             buildings forest glacier mountain sea street
## true
##
    buildings
                    114
                           19
                                   15
                                            12 9
    forest
                     10
                           158
                                    1
                                             6
                                                3
                                                      21
##
                                  121
                                            23 21
                                                      15
##
    glacier
                     12
                           6
##
    mountain
                     14
                           10
                                   32
                                           109 29
                                                       4
##
                           16
                                   50
    sea
                     18
                                            44 59
                                                      13
## street
                     29
                           26
                                   13
                                           6 3
                                                      123
```

```
imp = ranger_rf$variable.importance
```

```
imp = as.data.frame(imp)
head(imp)
##
            imp
## PC1 45.76979
## PC2 22.49475
## PC3 13.91928
## PC4 11.31050
## PC5 7.76092
## PC6 18.41533
imp$varnames = rownames(imp)
rownames(imp) = NULL
head(imp)
##
          imp varnames
## 1 45.76979
                   PC1
                   PC2
## 2 22.49475
## 3 13.91928
                   PC3
## 4 11.31050
                   PC4
## 5 7.76092
                   PC5
## 6 18.41533
                   PC6
ggplot(imp,
       aes(x=reorder(varnames, imp),
           y=imp)) +
        geom_point() +
        geom_segment(aes(x=varnames, xend=varnames, y=0, yend=imp)) +
        ylab('Mean Decrease in Impurity') +
        xlab('Variable Name') +
        coord_flip() +
        theme_minimal()
```



```
pred_RF = predict(ranger_rf, test75)
```

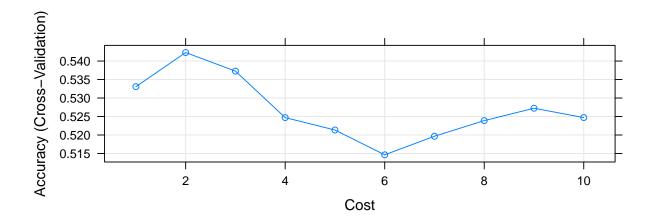
```
table(pred_RF$predictions, test75$class)
##
               buildings forest glacier mountain sea street
##
     buildings
                       62
                               5
                                        4
                                                 5 15
                                                            13
##
     forest
                       26
                             158
                                        7
                                                10
                                                    20
                                                            31
##
     glacier
                       37
                               6
                                                32
                                                    58
                                                            34
                                      133
                                                     39
                                                             6
##
     mountain
                       25
                              11
                                       19
                                               125
##
     sea
                       16
                               5
                                       21
                                                23
                                                    58
                                                             8
     street
                       34
                              15
                                       13
                                                 3
                                                    10
                                                           108
prop.table(table(pred_RF$predictions == test75$class))
##
##
       FALSE
                   TRUE
## 0.4610879 0.5389121
```

#SVM

Use 10-fold CV to assess the optimal cost for the fitting process. Here, the radial basis kernel is chosen based on its better performance.

```
line.svm75 <- train(x=df75[,-1], y=df75[,1],
                 method = "svmRadial",
                 trControl =trainControl(method = "cv", number = 10),
                  tuneGrid =expand.grid(C =c(1:10), sigma = 0.01))
line.svm75
## Support Vector Machines with Radial Basis Function Kernel
##
## 1195 samples
    99 predictor
##
      6 classes: 'buildings', 'forest', 'qlacier', 'mountain', 'sea', 'street'
##
##
## No pre-processing
## Resampling: Cross-Validated (10 fold)
## Summary of sample sizes: 1076, 1076, 1075, 1075, 1076, 1075, ...
## Resampling results across tuning parameters:
##
##
       Accuracy
                   Kappa
##
      1 0.5330742 0.4397105
##
      2 0.5423039 0.4507796
##
      3 0.5372759 0.4447529
      4 0.5247059 0.4296965
##
##
      5 0.5213515 0.4256573
      6 0.5146359 0.4175993
##
##
      7 0.5196639 0.4236122
      8 0.5238866 0.4286705
##
##
      9 0.5272339 0.4326825
##
     10 0.5247199 0.4296665
##
## Tuning parameter 'sigma' was held constant at a value of 0.01
```

```
## Accuracy was used to select the optimal model using the largest value. ## The final values used for the model were sigma = 0.01 and C = 2. plot(line.svm75)
```



The final fitted model is decided using the radial basis kernel when cost is 2 and gamma is 0.01, which is confirmed in the previous CV plot.

```
# Conduct the multi-class SVM on the training set
img_svm75 <- svm(x=df75[,-1], y=df75[,1], type="C-classification",</pre>
                 kernel="radial", cost = 2, gamma = 0.01 )
summary(img_svm75)
##
## Call:
## svm.default(x = df75[, -1], y = df75[, 1], type = "C-classification",
##
       kernel = "radial", gamma = 0.01, cost = 2)
##
##
## Parameters:
##
      SVM-Type: C-classification
   SVM-Kernel: radial
##
##
          cost: 2
##
## Number of Support Vectors: 1141
##
    ( 200 164 192 186 200 199 )
##
##
## Number of Classes: 6
##
## Levels:
## buildings forest glacier mountain sea street
# Prediction on test data and save the confusion matrix in sum_table
pred_75 <- predict(img_svm75, newdata=test75[,-1], type="class")</pre>
```

```
svm_table <- table(pred_75, test75$class)</pre>
# Calculate the misclassification rate
matrix=as.matrix(table(pred_75, test75$class))
diag=diag(matrix)
n=sum(matrix)
accuracy= sum(diag)/n
accuracy
## [1] 0.5305439
svm_table
##
## pred_75     buildings forest glacier mountain sea street
## buildings
                   67
                          5
                                 16
                                         8 18
                                                    26
    forest
##
                    20
                         150
                                 2
                                          4 15
                                                    24
                                         25 40
##
    glacier
                    19
                          4
                                 115
                                                    19
##
    mountain
                    33
                          14
                                  21
                                         118 39
                                                    12
##
   sea
                    31
                          12
                                  33
                                         39 77
                                                   12
                    30
                          15
                                                   107
##
    street
                                  10
                                         4 11
```