

# Main

*Fall2019-proj3-grp7*

In your final repo, there should be an R markdown file that organizes **all computational steps** for evaluating your proposed Facial Expression Recognition framework.

This file is currently a template for running evaluation experiments. You should update it according to your codes but following precisely the same structure.

```
if(!require("EBImage")){
  source("https://bioconductor.org/biocLite.R")
  biocLite("EBImage")
}
if(!require("R.matlab")){
  install.packages("R.matlab")
}
if(!require("readxl")){
  install.packages("readxl")
}

if(!require("dplyr")){
  install.packages("dplyr")
}
if(!require("readxl")){
  install.packages("readxl")
}

if(!require("ggplot2")){
  install.packages("ggplot2")
}

if(!require("caret")){
  install.packages("caret")
}

if(!require("OpenImageR")){
  install.packages("OpenImageR")
}

if(!require("FSelectorRcpp")){
  install.packages("FSelectorRcpp")
}

if(!require("mlr")){
  install.packages("mlr")
}

if(!require("kernlab")){
  install.packages("kernlab")
}
```

```

if(!require("gbm")){
  install.packages("gbm")
}

if(!require("class")){
  install.packages("class")
}
if(!require("MASS")){
  install.packages("MASS")
}
if(!require("e1071")){
  install.packages("e1071")
}
library(R.matlab)
library(readxl)
library(dplyr)
library(EBImage)
library(ggplot2)
library(caret)
library(OpenImageR)
library(FSelectorRcpp)
library(mlr)
library(kernlab)
library(gbm)
library(class)

library("e1071")
library("MASS")

```

**Step 0** set work directories, extract paths, summarize

```

set.seed(0)
# setwd("~/Desktop/5243/Project 3/fall2019-proj3-sec2--grp7-master/doc")
# here replace it with your own path or manually set it in RStudio to where this rmd file is located.
# use relative path for reproducibility
# setwd("../doc")

```

Provide directories for training images. Training images and Training fiducial points will be in different subfolders.

```

train_dir <- "../data/train_set/" # This will be modified for different data sets.
train_image_dir <- paste(train_dir, "images/", sep="")
train_pt_dir <- paste(train_dir, "points/", sep="")
train_label_path <- paste(train_dir, "label.csv", sep="")

```

**Step 1:** set up controls for evaluation experiments.

In this chunk, we have a set of controls for the evaluation experiments.

- (T/F) cross-validation on the training set

- (number) K, the number of CV folds
- (T/F) process features for training set
- (T/F) run evaluation on an independent test set
- (T/F) process features for test set

```
run.cv=TRUE # run cross-validation on the training set
K <- 5 # number of CV folds
run.feature.train=FALSE # process features for training set
run.test=TRUE # run evaluation on an independent test set
run.feature.test=TRUE # process features for test set
run.feature.test.test=FALSE # process features for test_test set
```

Using cross-validation or independent test set evaluation, we compare the performance of models with different specifications. In this Starter Code, we tune parameter k (number of neighbours) for KNN.

## Step 2: import data and train-test split

```
#train-test split
info <- read.csv(train_label_path)
n <- nrow(info)
n_train <- round(n*(4/5), 0)
train_idx <- sample(info$Index, n_train, replace = F)
test_idx <- setdiff(info$Index, train_idx)
```

If you choose to extract features from images, such as using Gabor filter, R memory will exhaust all images are read together. The solution is to repeat reading a smaller batch(e.g 100) and process them.

```
n_files <- length(list.files(train_image_dir))

image_list <- list()
for(i in 1:100){
  image_list[[i]] <- readImage(paste0(train_image_dir, sprintf("%04d", i), ".jpg"))
}
```

Fiducial points are stored in matlab format. In this step, we read them and store them in a list.

```
#function to read fiducial points
#input: index
#output: matrix of fiducial points corresponding to the index
readMat.matrix <- function(index){
  return(round(readMat(paste0(train_pt_dir, sprintf("%04d", index), ".mat"))[[1]],0))
}

#load fiducial points
fiducial_pt_list <- lapply(1:n_files, readMat.matrix)
save(fiducial_pt_list, file="../output/fiducial_pt_list.RData")
```

## Step 3: feature selection

feature.R should be the wrapper for all your feature engineering functions and options. The function feature( ) should have options that correspond to different scenarios for your project and produces an R

object that contains features and responses that are required by all the models you are going to evaluate later.

- feature.R
- Input: list of images or fiducial point
- Output: an RData file that contains extracted features and corresponding responses

```
source("../lib/feature.R")
tm_feature_train <- NA
if(run.feature.train){
  ## Distance calculation
  tm_feature_train <- system.time(dat_train <- feature_train(fiducial_pt_list, train_idx))
  dat_train <- cbind(dat_train, as.factor(info$emotion_idx[train_idx]))
  colnames(dat_train)[dim(dat_train)[2]] <- "emotion_idx"
  dat_train <- as.data.frame(dat_train)
  colnames(dat_train) <- make.names(colnames(dat_train), unique=T)

  ## Normalize
  tm_feature_train <- tm_feature_train +
    system.time(dat_train_stand <- feature_normalization(dat_train[,c(-dim(dat_train)[2])]))
  dat_train_stand <- cbind(dat_train_stand, dat_train$emotion_idx)
  colnames(dat_train_stand)[dim(dat_train_stand)[2]] <- "emotion_idx"

  ## Feature selection
  tm_feature_train <- tm_feature_train +
    system.time(feature_name <- feature_selection(dat_train_stand, "emotion_idx"))
  dat_train_selected <- dat_train[,feature_name]

  ## Calculate size from all selected distance
  tm_feature_train <- tm_feature_train +
    system.time(dat_train_double <- feature_selection_size(dat_train_selected))

  ## Add manually selected feature
  tm_feature_train <- tm_feature_train +
    system.time(dat_train_ratio <- manually_feature(fiducial_pt_list, train_idx))

  dat_train_selected_stand <- cbind(dat_train_selected, dat_train_double, dat_train_ratio)
  dat_train_selected_stand <- feature_normalization(dat_train_selected_stand)
  dat_train_selected_stand <- cbind(dat_train_selected_stand, dat_train$emotion_idx)
  colnames(dat_train_selected_stand)[dim(dat_train_selected_stand)[2]] <- "emotion_idx"
}

tm_feature_test <- NA
if(run.feature.test){
  ## This is the result from test

  feature_name <- c("point.7.to.point.21", "point.10.to.point.13", "point.10.to.point.33",
    "point.11.to.point.49", "point.12.to.point.55", "point.14.to.point.18",
    "point.23.to.point.50", "point.34.to.point.46", "point.50.to.point.62",
    "point.59.to.point.62")

  ## Distance Feature
  tm_feature_test <- system.time(dat_test <- feature_train(fiducial_pt_list, test_idx))
```

```

dat_test <- cbind(dat_test, as.factor(info$emotion_idx[test_idx]))
colnames(dat_test)[dim(dat_test)[2]] <- "emotion_idx"
dat_test <- as.data.frame(dat_test)
colnames(dat_test)<-make.names(colnames(dat_test),unique=T)
dat_test_selected <- dat_test[,feature_name]

## Size Feature
tm_feature_test <- tm_feature_test +
  system.time(dat_test_double <- feature_selection_size(dat_test_selected))

## Add manually selected feature
tm_feature_test <- tm_feature_test +
  system.time(dat_test_ratio <- manually_feature(fiducial_pt_list, test_idx))

dat_test_selected_stand <- cbind(dat_test_selected,dat_test_double,dat_test_ratio)
dat_test_selected_stand <- feature_normalization(dat_test_selected_stand)
dat_test_selected_stand <- cbind(dat_test_selected_stand,dat_test$emotion_idx)
colnames(dat_test_selected_stand)[dim(dat_test_selected_stand)[2]] <- "emotion_idx"
}

tm_feature_test_test <- NA
if(run.feature.test.test){
  test_test_idx = c(1:2500)
  ## This is the result from test
  feature_name <- c("point.7.to.point.21","point.10.to.point.13","point.10.to.point.33",
    "point.11.to.point.49","point.12.to.point.55","point.14.to.point.18",
    "point.23.to.point.50","point.34.to.point.46","point.50.to.point.62",
    "point.59.to.point.62")

  ## Distance Feature
  tm_feature_test_test <- system.time(dat_test_test <- feature_train(fiducial_pt_list, test_test_idx))
  colnames(dat_test_test)<-make.names(colnames(dat_test_test),unique=T)
  dat_test_test_selected <- dat_test_test[,feature_name]

  ## Size Feature
  tm_feature_test_test <- tm_feature_test_test +
    system.time(dat_test_test_double <- feature_selection_size(dat_test_test_selected))

  ## Add manually selected feature
  tm_feature_test_test <- tm_feature_test_test +
    system.time(dat_test_test_ratio <- manually_feature(fiducial_pt_list, test_test_idx))

  dat_test_selected_stand_test <- cbind(dat_test_test_selected,dat_test_test_double,
    dat_test_test_ratio)
  dat_test_selected_stand_test <- feature_normalization(dat_test_selected_stand_test)
}

##save(dat_train_selected_stand, file="../output/feature_train.RData")
## Because feature train takes over 10 hours, we do not knit this part.
## The previous feature selection train is already included in the output file.
save(dat_test_selected_stand, file="../output/feature_test.RData")

```

```
##save(dat_test_selected_stand_test, file="../output/feature_test_test.RData")
```

### ### Step 4: Train a classification model with training features and responses

```
load("../output/feature_train.RData")
dat_train_selected <- dat_train_selected_ratio_stand55
dat_test_selected <- dat_test_selected_stand
```

Call the train models and test models from library:

- 1. KNN
- 2. LDA
- 3. SVM with radial kernel (improved model)
- 4. GBM with tree stumps (baseline Model)

#### 1. KNN

- Do model selection by choosing among different values of training model parameters.
- Choose the “best” parameter value
- Train accuracy:
- KNN: Run test on test images
- evaluation
- Summarize Running Time

#### 2. LDA

- Train the model with the entire training set using the selected model (model parameter) via cross-validation.

```
source("../lib/train_lda.R")
tm_train=NA
tm_train <- system.time(fit_train <- train(dat_train_selected, par_best))
save(fit_train, file="../output/fit_train.RData")
```

- Train Accuracy:

```
source("../lib/test_lda.R")
tm_test=NA
if(run.test){
  pred_train <- test(fit_train, dat_train_selected)
}

accu <- mean(dat_train_selected$emotion_idx == pred_train)
accu
```

```
## [1] 0.533
```

- LDA: Run test on test images

```
source("../lib/test_lda.R")
tm_test=NA
if(run.test){
  load(file="../output/fit_train.RData")
  tm_test <- system.time(pred <- test(fit_train, dat_test_selected))
}
```

- evaluation

```
accu <- mean(dat_test_selected$emotion_idx == pred)
cat("The accuracy of model:", "is", accu*100, "%.\n")
```

```
## The accuracy of model: is 47.6 %.
```

```
library(caret)
confusionMatrix(pred, dat_test_selected$emotion_idx)
```

```
## Confusion Matrix and Statistics
```

```
##
##           Reference
## Prediction  1  2  3  4  5  6  7  8  9 10 11 12 13 14 15 16 17 18 19 20 21
##           1 15  0  2  0  1  1  3  0  0  1  0  1  1  0  0  0  0  1  1  0  1
##           2  0 17  0  0  0  0  0  2  0  0  0  0  0  0  0  0  0  0  0  0  0
##           3  5  0 11  3  0  1  0  0  0  2  0  0  2  0  0  0  0  0  0  0  0
##           4  0  0  0 10  0  0  0  0  0  2  2  0  2  0  0  0  0  0  0  1  0
##           5  0  0  0  0 15  0  0  2  0  0  0  0  0  0  0  0  2  0  0  0  0
##           6  5  0  2  1  1 13  0  0  1  6  4  7  8  1  1  1  3  0  0  2  2
##           7  0  0  0  0  0  0 13  0  1  0  0  0  0  1  1  0  1  2  1  2  0
##           8  0  2  0  0  0  0  0 15  0  0  0  0  0  0  0  0  1  0  0  1  0
##           9  0  7  0  0  0  0  0  1 12  0  0  0  0  0  0  0  0  0  0  0  1
##          10  0  0  2  2  0  0  0  0  0 12  3  1  2  0  0  1  0  0  1  0  2
##          11  0  0  0  0  0  0  0  0  0  0  8  1  0  0  0  0  0  0  0  0  0
##          12  1  0  0  1  0  2  0  0  0  1  3 10  4  0  0  0  0  0  0  0  0
##          13  0  0  1  6  0  1  0  0  0  2  1  3  3  0  1  0  0  0  0  0  0
##          14  0  0  0  0  0  2  0  0  0  0  0  1  0 13  4  0  1  0  0  1  3
##          15  0  0  0  0  1  0  0  0  0  0  0  0  0  5 10  0  0  0  1  1  0
##          16  0  0  2  0  0  0  0  0  0  0  0  0  0  0  0 14  3  2  0  0  0
##          17  0  0  0  0  0  0  1  1  0  0  0  0  0  0  2  0  9  2  1  2  2
##          18  0  0  0  0  2  0  0  0  0  0  0  0  0  1  0  1  6 10  0  1  0
##          19  0  0  0  0  0  0  2  0  0  0  0  0  0  0  2  0  1  3  5  3  1
##          20  0  1  0  0  0  0  0  0  1  0  0  0  0  1  0  1  2  1  1  6  0
##          21  0  1  0  0  0  1  4  0  1  0  0  0  0  0  1  0  1  2  3  3 12
##          22  0  0  0  2  0  0  0  0  0  0  2  0  0  0  0  1  1  0  0  1  0
##
##           Reference
## Prediction 22
##           1  0
##           2  1
```

```

##      3  6
##      4  0
##      5  0
##      6  8
##      7  0
##      8  0
##      9  0
##     10  0
##     11  0
##     12  0
##     13  0
##     14  0
##     15  0
##     16  2
##     17  0
##     18  1
##     19  1
##     20  0
##     21  2
##     22  5
##
## Overall Statistics
##
##           Accuracy : 0.476
##           95% CI : (0.4315, 0.5208)
##       No Information Rate : 0.062
##       P-Value [Acc > NIR] : < 2.2e-16
##
##           Kappa : 0.4512
##
## Mcnemar's Test P-Value : NA
##
## Statistics by Class:
##
##           Class: 1 Class: 2 Class: 3 Class: 4 Class: 5 Class: 6
## Sensitivity      0.5769  0.6071  0.5500  0.4000  0.7500  0.6190
## Specificity      0.9726  0.9936  0.9604  0.9853  0.9917  0.8894
## Pos Pred Value   0.5357  0.8500  0.3667  0.5882  0.7895  0.1970
## Neg Pred Value   0.9767  0.9771  0.9809  0.9689  0.9896  0.9816
## Prevalence       0.0520  0.0560  0.0400  0.0500  0.0400  0.0420
## Detection Rate   0.0300  0.0340  0.0220  0.0200  0.0300  0.0260
## Detection Prevalence 0.0560  0.0400  0.0600  0.0340  0.0380  0.1320
## Balanced Accuracy 0.7747  0.8004  0.7552  0.6926  0.8708  0.7542
##
##           Class: 7 Class: 8 Class: 9 Class: 10 Class: 11
## Sensitivity      0.5652  0.7143  0.7500  0.4615  0.3478
## Specificity      0.9811  0.9916  0.9814  0.9705  0.9979
## Pos Pred Value   0.5909  0.7895  0.5714  0.4615  0.8889
## Neg Pred Value   0.9791  0.9875  0.9916  0.9705  0.9695
## Prevalence       0.0460  0.0420  0.0320  0.0520  0.0460
## Detection Rate   0.0260  0.0300  0.0240  0.0240  0.0160
## Detection Prevalence 0.0440  0.0380  0.0420  0.0520  0.0180
## Balanced Accuracy 0.7732  0.8530  0.8657  0.7160  0.6729
##
##           Class: 12 Class: 13 Class: 14 Class: 15 Class: 16
## Sensitivity      0.4167  0.1364  0.5909  0.4545  0.7368

```



## Specificity	0.9748	0.9686	0.9749	0.9833	0.9813
## Pos Pred Value	0.4545	0.1667	0.5200	0.5556	0.6087
## Neg Pred Value	0.9707	0.9606	0.9811	0.9751	0.9895
## Prevalence	0.0480	0.0440	0.0440	0.0440	0.0380
## Detection Rate	0.0200	0.0060	0.0260	0.0200	0.0280
## Detection Prevalence	0.0440	0.0360	0.0500	0.0360	0.0460
## Balanced Accuracy	0.6957	0.5525	0.7829	0.7189	0.8591
##	Class: 17	Class: 18	Class: 19	Class: 20	Class: 21
## Sensitivity	0.2903	0.4348	0.3571	0.2500	0.5000
## Specificity	0.9765	0.9748	0.9733	0.9832	0.9601
## Pos Pred Value	0.4500	0.4545	0.2778	0.4286	0.3871
## Neg Pred Value	0.9542	0.9728	0.9813	0.9630	0.9744
## Prevalence	0.0620	0.0460	0.0280	0.0480	0.0480
## Detection Rate	0.0180	0.0200	0.0100	0.0120	0.0240
## Detection Prevalence	0.0400	0.0440	0.0360	0.0280	0.0620
## Balanced Accuracy	0.6334	0.7048	0.6652	0.6166	0.7300
##	Class: 22				
## Sensitivity	0.1923				
## Specificity	0.9852				
## Pos Pred Value	0.4167				
## Neg Pred Value	0.9570				
## Prevalence	0.0520				
## Detection Rate	0.0100				
## Detection Prevalence	0.0240				
## Balanced Accuracy	0.5888				

Note that the accuracy is not high but is better than that of random guess(4.5%).

- Summarize Running Time Prediction performance matters, so does the running times for constructing features and for training the model, especially when the computation resource is limited.

```
cat("Time for training model=", tm_train[1], "s \n")
```

```
## Time for training model= 0.077 s
```

```
cat("Time for testing model=", tm_test[1], "s \n")
```

```
## Time for testing model= 0.005 s
```

### 3. SVM (*improved model*)

- Tune the SVM model with cross-validation:

```
## must have selected features first
tm_train=NA
tm_train <- system.time(tuned_parameters <- tune.svm(emotion_idx~.,
  data = dat_train_selected,
  gamma = 10^(-5:-1),
  cost = c(30,35,40),
  tunecontrol = tune.control(cross = 12)
))
summary(tuned_parameters)
```

```
##
## Parameter tuning of 'svm':
##
## - sampling method: 12-fold cross validation
##
## - best parameters:
##   gamma cost
##   0.001    35
##
## - best performance: 0.5394843
##
## - Detailed performance results:
##   gamma cost    error dispersion
## 1  1e-05    30 0.7414899 0.01624426
## 2  1e-04    30 0.5805437 0.03553754
## 3  1e-03    30 0.5409873 0.02619165
## 4  1e-02    30 0.5654865 0.03189845
## 5  1e-01    30 0.6474581 0.04071884
## 6  1e-05    35 0.7314918 0.01769932
## 7  1e-04    35 0.5765307 0.03410334
## 8  1e-03    35 0.5394843 0.02718324
## 9  1e-02    35 0.5714896 0.02705938
## 10 1e-01    35 0.6474581 0.04071884
## 11 1e-05    40 0.7164797 0.02016448
## 12 1e-04    40 0.5735156 0.03079428
## 13 1e-03    40 0.5414893 0.02932813
## 14 1e-02    40 0.5759986 0.02377225
## 15 1e-01    40 0.6474581 0.04071884
```

- Train the model

```
source("../lib/train_svm.R")
par_best=NULL
fit_train_final_svm <- train(dat_train_selected, tuned_parameters$best.parameters)
save(fit_train_final_svm, file="../output/fit_train_final.RData")
```

- Train accuracy:

```
source("../lib/test_svm.R")
load("../output/fit_train_final.RData")

if(run.test){
  pred_train <- test(fit_train_final_svm, dat_train_selected)
}

accu <- mean(dat_train_selected$emotion_idx == pred_train)
accu
```

```
## [1] 0.5725
```

- SVM: Run test on test images

```
source("../lib/test_svm.R")
tm_test=NA
if(run.test){
  load(file="../output/fit_train.RData")
  tm_test <- system.time(pred <- test(fit_train_final_svm, dat_test_selected))
}
```

\*\*\*\*\* SVM: Run test\_test on test images

```
source("../lib/test_svm.R")
tm_test=NA
if(run.test){
  load(file="../output/fit_train_final.RData")
  tm_test <- system.time(pred <- test(fit_train_final_svm, dat_test_selected))
}
```

- evaluation

```
accu <- mean(dat_test_selected$emotion_idx == pred)
cat("The accuracy of model:", "is", accu*100, "%.\n")
```

## The accuracy of model: is 55 %.

```
library(caret)
confusionMatrix(pred, dat_test_selected$emotion_idx)
```

## Confusion Matrix and Statistics

```
##
##           Reference
## Prediction  1  2  3  4  5  6  7  8  9 10 11 12 13 14 15 16 17 18 19 20 21
##           1 20  0  2  0  0  0  1  0  0  0  0  0  1  0  0  0  0  1  0  0  0
##           2  0 23  0  0  0  0  0  1  1  0  0  0  0  0  0  0  0  0  1  0  0
##           3  2  0 13  3  0  3  0  0  0  1  1  0  1  0  0  0  0  0  1  0  0
##           4  0  0  2 15  0  0  0  0  0  5  4  2  8  0  0  0  0  0  0  0  0
##           5  0  0  0  0 17  0  0  0  0  0  0  0  0  0  0  0  4  1  0  0  0
##           6  0  0  0  1  0  7  1  0  0  1  1  0  3  0  0  2  0  0  0  0  0
##           7  1  0  0  0  0  0  9  0  0  0  0  0  0  0  0  0  0  0  0  0  0
##           8  0  2  0  0  0  0  1 17  0  0  0  0  0  0  0  0  2  0  0  1  0
##           9  0  3  0  0  0  0  0  1 14  0  0  0  0  0  0  0  0  0  0  1  1
##          10  0  0  1  4  0  0  0  0  0 15  4  2  2  0  1  1  0  0  1  0  2
##          11  0  0  0  0  0  1  0  0  0  1  9  1  1  0  0  0  0  0  0  0  0
##          12  1  0  0  1  0  5  0  0  0  0  2 13  0  1  0  0  0  0  0  0  0
##          13  2  0  1  1  0  2  0  0  0  3  2  3  4  0  0  0  0  0  0  1
##          14  0  0  0  0  0  3  0  0  0  0  0  1  0 17  3  0  1  0  1  2  5
##          15  0  0  0  0  1  0  0  0  0  0  0  0  1  3 16  0  0  0  2  0  0
##          16  0  0  1  0  0  0  0  0  0  0  0  1  0  0  0 14  1  1  0  0  1
##          17  0  0  0  0  0  0  2  2  0  0  0  0  0  1  0  0 13  1  0  2  2
##          18  0  0  0  0  2  0  1  0  0  0  0  0  0  0  0  1  4 13  1  0  0
##          19  0  0  0  0  0  0  2  0  0  0  0  0  0  0  2  0  1  3  2  3  0
##          20  0  0  0  0  0  0  0  0  1  0  0  0  0  0  0  0  5  2  2  8  2
##          21  0  0  0  0  0  0  6  0  0  0  0  1  0  0  0  1  0  1  3  4 10
```

```

##          22  0  0  0  0  0  0  0  0  0  0  0  0  0  1  0  0  0  0  0  0  3  0
##          Reference
## Prediction 22
##          1  0
##          2  1
##          3  3
##          4  1
##          5  0
##          6  1
##          7  0
##          8  0
##          9  0
##         10  3
##         11  2
##         12  0
##         13  0
##         14  1
##         15  0
##         16  3
##         17  1
##         18  0
##         19  2
##         20  1
##         21  1
##         22  6
##
## Overall Statistics
##
##          Accuracy : 0.55
##          95% CI : (0.5052, 0.5942)
##    No Information Rate : 0.062
##    P-Value [Acc > NIR] : < 2.2e-16
##
##          Kappa : 0.5283
##
## McNemar's Test P-Value : NA
##
## Statistics by Class:
##
##          Class: 1 Class: 2 Class: 3 Class: 4 Class: 5 Class: 6
## Sensitivity      0.7692  0.8214  0.6500  0.6000  0.8500  0.3333
## Specificity      0.9895  0.9915  0.9688  0.9537  0.9896  0.9791
## Pos Pred Value   0.8000  0.8519  0.4643  0.4054  0.7727  0.4118
## Neg Pred Value   0.9874  0.9894  0.9852  0.9784  0.9937  0.9710
## Prevalence       0.0520  0.0560  0.0400  0.0500  0.0400  0.0420
## Detection Rate   0.0400  0.0460  0.0260  0.0300  0.0340  0.0140
## Detection Prevalence 0.0500  0.0540  0.0560  0.0740  0.0440  0.0340
## Balanced Accuracy 0.8793  0.9065  0.8094  0.7768  0.9198  0.6562
##
##          Class: 7 Class: 8 Class: 9 Class: 10 Class: 11
## Sensitivity      0.3913  0.8095  0.8750  0.5769  0.3913
## Specificity      0.9979  0.9875  0.9876  0.9557  0.9874
## Pos Pred Value   0.9000  0.7391  0.7000  0.4167  0.6000
## Neg Pred Value   0.9714  0.9916  0.9958  0.9763  0.9711
## Prevalence       0.0460  0.0420  0.0320  0.0520  0.0460

```

## Detection Rate	0.0180	0.0340	0.0280	0.0300	0.0180
## Detection Prevalence	0.0200	0.0460	0.0400	0.0720	0.0300
## Balanced Accuracy	0.6946	0.8985	0.9313	0.7663	0.6894
##	Class: 12	Class: 13	Class: 14	Class: 15	Class: 16
## Sensitivity	0.5417	0.1818	0.7727	0.7273	0.7368
## Specificity	0.9790	0.9686	0.9644	0.9854	0.9834
## Pos Pred Value	0.5652	0.2105	0.5000	0.6957	0.6364
## Neg Pred Value	0.9769	0.9626	0.9893	0.9874	0.9895
## Prevalence	0.0480	0.0440	0.0440	0.0440	0.0380
## Detection Rate	0.0260	0.0080	0.0340	0.0320	0.0280
## Detection Prevalence	0.0460	0.0380	0.0680	0.0460	0.0440
## Balanced Accuracy	0.7603	0.5752	0.8686	0.8563	0.8601
##	Class: 17	Class: 18	Class: 19	Class: 20	Class: 21
## Sensitivity	0.4194	0.5652	0.1429	0.3333	0.4167
## Specificity	0.9765	0.9811	0.9733	0.9727	0.9643
## Pos Pred Value	0.5417	0.5909	0.1333	0.3810	0.3704
## Neg Pred Value	0.9622	0.9791	0.9753	0.9666	0.9704
## Prevalence	0.0620	0.0460	0.0280	0.0480	0.0480
## Detection Rate	0.0260	0.0260	0.0040	0.0160	0.0200
## Detection Prevalence	0.0480	0.0440	0.0300	0.0420	0.0540
## Balanced Accuracy	0.6980	0.7732	0.5581	0.6530	0.6905
##	Class: 22				
## Sensitivity	0.2308				
## Specificity	0.9916				
## Pos Pred Value	0.6000				
## Neg Pred Value	0.9592				
## Prevalence	0.0520				
## Detection Rate	0.0120				
## Detection Prevalence	0.0200				
## Balanced Accuracy	0.6112				

## Summarize Running Time

Prediction performance matters, so does the running times for constructing features and for training the model, especially when the computation resource is limited.

```
#cat("Time for constructing training features=", tm_feature_train[1], "s \n")
#cat("Time for constructing testing features=", tm_feature_test[1], "s \n")
cat("Time for training model=", tm_train[1], "s \n")
```

```
## Time for training model= 259.809 s
```

```
cat("Time for testing model=", tm_test[1], "s \n")
```

```
## Time for testing model= 0.119 s
```

## 4. GBM (*Baseline Model*)

- Tune GBM.

```

hyper_grid <- expand.grid(
  shrinkage = c(.001, .01),
  interaction.depth = c(1, 3),
  n.minobsinnode = c(5, 10),
  bag.fraction = c(.65, .8),
  optimal_trees = 0,           # a place to dump results
  min_RMSE = 0                 # a place to dump results
)

# randomize data
random_index <- sample(1:nrow(dat_train_selected), nrow(dat_train_selected))
random_train <- dat_train_selected[random_index, ]

# grid search
for(i in 1:nrow(hyper_grid)) {

  # reproducibility
  set.seed(123)

  # train model
  gbm.tune <- gbm(
    formula = emotion_idx~.,
    distribution = "multinomial",
    data = random_train,
    n.trees = 100,
    interaction.depth = hyper_grid$interaction.depth[i],
    shrinkage = hyper_grid$shrinkage[i],
    n.minobsinnode = hyper_grid$n.minobsinnode[i],
    bag.fraction = hyper_grid$bag.fraction[i],
    train.fraction = .75,
    n.cores = NULL, # will use all cores by default
    verbose = FALSE
  )

  # add min training error and trees to grid
  hyper_grid$optimal_trees[i] <- which.min(gbm.tune$valid.error)
  hyper_grid$min_RMSE[i] <- sqrt(min(gbm.tune$valid.error))
}

hyper_grid %>%
  dplyr::arrange(min_RMSE) %>%
  head(10)

```

```

##      shrinkage interaction.depth n.minobsinnode bag.fraction optimal_trees
## 1      0.010             3             10          0.65           100
## 2      0.010             3             10          0.80           100
## 3      0.010             3              5          0.65           100
## 4      0.010             3              5          0.80           100
## 5      0.010             1             10          0.65           100
## 6      0.010             1              5          0.65           100
## 7      0.010             1             10          0.80           100
## 8      0.010             1              5          0.80           100
## 9      0.001             3             10          0.65           100

```

```
## 10      0.001          3          5          0.65          100
##      min_RMSE
## 1  1.487001
## 2  1.491959
## 3  1.492670
## 4  1.497072
## 5  1.581901
## 6  1.583199
## 7  1.583916
## 8  1.584736
## 9  1.690817
## 10 1.692194
```

- Train the model with the entire training set using the selected model (model parameter) via cross-validation.

```
source("../lib/train_gbm.R")
tm_train=NA
tm_train <- system.time(fit_train_baseline <- train(dat_train_selected, par = NULL))
save(fit_train_baseline, file="../output/fit_train_baseline_final.RData")
```

- Train Error:

```
source("../lib/test_gbm.R")
load("../output/fit_train_baseline_final.RData")

tm_test=NA
if(run.test){
  tm_test <- system.time(pred_train <- test(fit_train_baseline, dat_train_selected))
}

labels = colnames(pred_train)[apply(pred_train, 1, which.max)]
accu <- mean(dat_train_selected$emotion_idx == labels)
accu
```

```
## [1] 0.7705
```

- GBM: Run test on test images

```
source("../lib/test_gbm.R")
tm_test=NA
if(run.test){
  load(file="../output/fit_train.RData")
  tm_test <- system.time(pred <- test(fit_train_baseline, dat_test_selected))
}
```

- GBM: Run test\_test on test images

```
source("../lib/test_gbm.R")
tm_test_test=NA
if(run.feature.test.test){
  load(file="../output/fit_train_baseline_final.RData")
  tm_test <- system.time(pred <- test(fit_train_baseline, dat_test_selected))
}
```

- evaluation

```
labels = colnames(pred)[apply(pred, 1, which.max)]
accu <- mean(dat_test_selected$emotion_idx == labels)
cat("The accuracy of model:", "is", accu*100, "%.\n")
```

```
## The accuracy of model: is 59.6 %.
```

```
library(caret)
confusionMatrix(as.factor(labels), dat_test_selected$emotion_idx)
```

```
## Warning in confusionMatrix.default(as.factor(labels),
## dat_test_selected$emotion_idx): Levels are not in the same order for
## reference and data. Refactoring data to match.
```

```
## Confusion Matrix and Statistics
```

```
##
##           Reference
## Prediction  1  2  3  4  5  6  7  8  9 10 11 12 13 14 15 16 17 18 19 20 21
##           1 22  0  3  1  0  1  0  0  0  1  1  2  2  0  0  1  1  0  0  0  0
##           2  0 19  0  0  0  0  0  1  4  0  0  0  0  0  0  0  0  0  1  0  0
##           3  1  0 14  2  0  1  0  0  0  4  3  0  1  0  0  0  0  0  1  0  0
##           4  0  0  0 17  0  0  0  0  0  2  1  1  5  0  0  0  0  0  0  0  0
##           5  0  0  0  0 18  0  0  0  0  0  0  0  0  0  0  0  1  1  0  0  0
##           6  1  0  0  0  0 10  0  0  1  2  1  2  1  0  0  0  0  0  0  0  0
##           7  1  0  0  0  0  0 14  0  0  0  0  0  0  0  1  0  1  1  1  1  1
##           8  0  1  0  0  0  0  2 16  0  0  0  0  0  0  0  0  1  0  0  1  0
##           9  0  7  0  0  0  0  0  1 11  0  0  0  0  0  0  0  0  0  0  1  2
##          10  0  0  1  4  0  0  0  0  0 16  2  2  2  0  0  1  0  0  1  0  1
##          11  0  0  1  0  0  1  0  0  0  0 14  3  2  0  0  0  0  0  0  0  0
##          12  1  0  0  0  0  4  0  0  0  0  1 11  1  0  0  0  0  0  0  0  0
##          13  0  0  0  1  0  1  0  0  0  0  0  0  6  0  0  0  0  0  0  0  1
##          14  0  0  0  0  0  2  0  0  0  0  0  0  0 17  3  0  1  0  1  1  2
##          15  0  0  0  0  1  0  0  0  0  0  0  0  1  3 15  0  1  0  2  0  0
##          16  0  0  1  0  0  0  0  0  0  0  0  1  0  0  0 15  2  2  0  0  0
##          17  0  0  0  0  1  0  0  1  0  0  0  0  0  1  0  0 15  1  0  1  0
##          18  0  0  0  0  0  0  1  0  0  0  0  0  0  0  1  1  5 13  0  0  0
##          19  0  0  0  0  0  0  2  0  0  0  0  0  0  0  1  1  0  3  5  2  2
##          20  0  1  0  0  0  0  1  2  0  0  0  0  1  1  0  0  2  2  1 13  5
##          21  0  0  0  0  0  1  3  0  0  0  0  0  0  0  1  0  1  0  1  4 10
##          22  0  0  0  0  0  0  0  0  0  1  0  2  0  0  0  0  0  0  0  0  0
##
##           Reference
## Prediction 22
##           1  1
##           2  0
##           3  3
##           4  0
##           5  0
##           6  0
##           7  0
##           8  0
##           9  1
```



```

##      10  1
##      11  1
##      12  1
##      13  0
##      14  0
##      15  0
##      16  2
##      17  1
##      18  0
##      19  3
##      20  0
##      21  5
##      22  7
##
## Overall Statistics
##
##           Accuracy : 0.596
##           95% CI : (0.5515, 0.6393)
##       No Information Rate : 0.062
##       P-Value [Acc > NIR] : < 2.2e-16
##
##           Kappa : 0.5766
##
## Mcnemar's Test P-Value : NA
##
## Statistics by Class:
##
##           Class: 1 Class: 2 Class: 3 Class: 4 Class: 5 Class: 6
## Sensitivity      0.8462  0.6786  0.7000  0.6800  0.9000  0.4762
## Specificity      0.9705  0.9873  0.9667  0.9811  0.9958  0.9833
## Pos Pred Value   0.6111  0.7600  0.4667  0.6538  0.9000  0.5556
## Neg Pred Value   0.9914  0.9811  0.9872  0.9831  0.9958  0.9772
## Prevalence       0.0520  0.0560  0.0400  0.0500  0.0400  0.0420
## Detection Rate   0.0440  0.0380  0.0280  0.0340  0.0360  0.0200
## Detection Prevalence 0.0720  0.0500  0.0600  0.0520  0.0400  0.0360
## Balanced Accuracy 0.9083  0.8329  0.8333  0.8305  0.9479  0.7297
##
##           Class: 7 Class: 8 Class: 9 Class: 10 Class: 11
## Sensitivity      0.6087  0.7619  0.6875  0.6154  0.6087
## Specificity      0.9853  0.9896  0.9752  0.9684  0.9832
## Pos Pred Value   0.6667  0.7619  0.4783  0.5161  0.6364
## Neg Pred Value   0.9812  0.9896  0.9895  0.9787  0.9812
## Prevalence       0.0460  0.0420  0.0320  0.0520  0.0460
## Detection Rate   0.0280  0.0320  0.0220  0.0320  0.0280
## Detection Prevalence 0.0420  0.0420  0.0460  0.0620  0.0440
## Balanced Accuracy 0.7970  0.8757  0.8314  0.7919  0.7960
##
##           Class: 12 Class: 13 Class: 14 Class: 15 Class: 16
## Sensitivity      0.4583  0.2727  0.7727  0.6818  0.7895
## Specificity      0.9832  0.9937  0.9791  0.9833  0.9834
## Pos Pred Value   0.5789  0.6667  0.6296  0.6522  0.6522
## Neg Pred Value   0.9730  0.9674  0.9894  0.9853  0.9916
## Prevalence       0.0480  0.0440  0.0440  0.0440  0.0380
## Detection Rate   0.0220  0.0120  0.0340  0.0300  0.0300
## Detection Prevalence 0.0380  0.0180  0.0540  0.0460  0.0460
## Balanced Accuracy 0.7208  0.6332  0.8759  0.8325  0.8864

```

##	Class: 17	Class: 18	Class: 19	Class: 20	Class: 21
## Sensitivity	0.4839	0.5652	0.3571	0.5417	0.4167
## Specificity	0.9872	0.9832	0.9712	0.9664	0.9664
## Pos Pred Value	0.7143	0.6190	0.2632	0.4483	0.3846
## Neg Pred Value	0.9666	0.9791	0.9813	0.9766	0.9705
## Prevalence	0.0620	0.0460	0.0280	0.0480	0.0480
## Detection Rate	0.0300	0.0260	0.0100	0.0260	0.0200
## Detection Prevalence	0.0420	0.0420	0.0380	0.0580	0.0520
## Balanced Accuracy	0.7355	0.7742	0.6642	0.7540	0.6915
##	Class: 22				
## Sensitivity	0.2692				
## Specificity	0.9937				
## Pos Pred Value	0.7000				
## Neg Pred Value	0.9612				
## Prevalence	0.0520				
## Detection Rate	0.0140				
## Detection Prevalence	0.0200				
## Balanced Accuracy	0.6315				

###Reference

- Du, S., Tao, Y., & Martinez, A. M. (2014). Compound facial expressions of emotion. Proceedings of the National Academy of Sciences, 111(15), E1454-E1462.