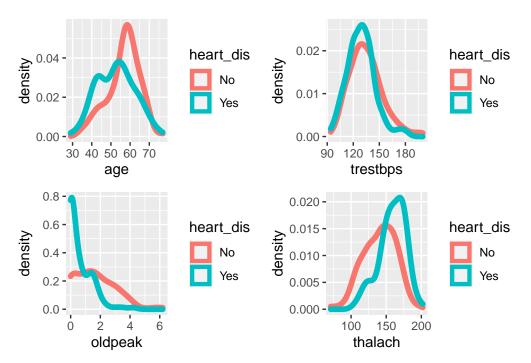
STAA577: HW3

- 36 points
- Due: see Canvas for due dates
- Submit your HW by uploading a PDF or DOC file to Canvas. I recommend using the HW3_yourname.Rmd file as a template to submit your answers.
- Include all of your R code in an appendix, unless explicitly asked to show it. Your final document may include R output and graphics, but all code should be in the appendix.
- 1. (2 points) Textbook (An Introduction to Statistical Learning with Applications in R, by James, Witten, Hastie, and Tibshirani) problem: Exercises 4.7, Conceptual Question #2.
- 2. (3 points) Textbook (An Introduction to Statistical Learning with Applications in R, by James, Witten, Hastie, and Tibshirani) problem: Exercises 4.7, Conceptual Question #3.
- 3. Recall the heart data from HW2. It contains information about 303 patients and whether or not they have heart disease. The variable target is 1 if the patient has heart disease and 0 if they do not. The data are split into a training and test set on Canvas. You can read more about the variables in the data here: https://www.kaggle.com/datasets/redwankarimsony/heart-disease-data. Be sure to treat the variables sex, cp, fps, exang and restecg as factors in both the training and test data sets.
 - (a) (4 points) Use the training data to find a multiple logistic regression model using age, sex, cp, trestbps, thalach, exang, oldpeak, ca, and thal as predictors. Use a Bayes classification rule and report both the prediction training accuracy the prediction test accuracy.
 - (b) (1 point) For plotting purposes, create a variable called heart_dis which is set to "yes" for patients with heart disease and "no" for patients without heart disease. Add this variable to your training data set.

```
heart_dis <- rep("Yes", dim(trainingdata)[1])
heart_dis[trainingdata$target == 0] <- "No"
trainingdata$heart_dis <- heart_dis</pre>
```

(c) (5 points) Using ggplot and geom_density(), create four plots, one for each of the continuous predictors: age, trestbps, oldpeak, and thalach showing the density of the predictor for each of the two classes of heart disease status. Within each plot, use the color option in geom_density() to create two density plots, one for each of the heart disease groups. Save each figure as an individual graph, then use the + function in the patchwork package to arrange the four graphs into a single image (or you can use facetting). Here is an example of what your final plot should look like:



- (d) (4 points) Use the training data to fit an LDA model using age, sex, cp, trestbps, thalach, exang, oldpeak, ca, and thal as predictors. Use a Bayes classification rule and report both the prediction training accuracy the prediction. test accuracy.
- (e) (3 points) Repeat the previous problem using QDA.
- (f) (2 points) Examine documentation for the knn() function in the class} package. In order to use this function, you will need to create two new data sets (one for training, one for testing) that contain only the predictorsage, sex, cp, trestbps, thalach, exang, oldpeak, ca, andthal. Create these two data sets. I recommend using a pipe and theselect()' function.
- (g) (2 points) KNN predicts an observation's class by identifying the observations that are nearest to it. As a result, the scale of the variables matters. To control for differences in units and scale, we need to standardize our numerical variables. Use the scale() function to center and standardize the four numerical variables (age, trestbps, oldpeak, thalach) in both the training and test data sets.
- (h) (4 points) Use the knn() function to use K-nearest-neighbors with k = 14 to classify the test data. Report only your **test accuracy**.
- 4. (6 points) Using the Boston data set, fit classification models in order to predict whether a given suburb has a crime rate above or below the median (use the code below). The testing and training data sets are provided in the code. Make sure to not include a crime rate as a predictor. Explore logistic regression, LDA, and KNN models using various subsets of the predictors. Describe your findings (at least 3 different takes is enough). Feel free to be creative.

```
library(MASS) ## data is in here
set.seed(10) ## reproducible

## take a look
head(Boston)

## create training and testing data (More on this in Week 4)
trn_samples <- sample(1:dim(Boston)[1], 440, replace=FALSE)</pre>
```

```
training_Boston <- Boston[trn_samples,]
testing_Boston <- Boston[-trn_samples,]

## create a response variable
training_Boston$crimMedian <- training_Boston$crim > median(training_Boston$crim)
testing_Boston$crimMedian <- testing_Boston$crim > median(training_Boston$crim)
```