Data Science II: Homework 5

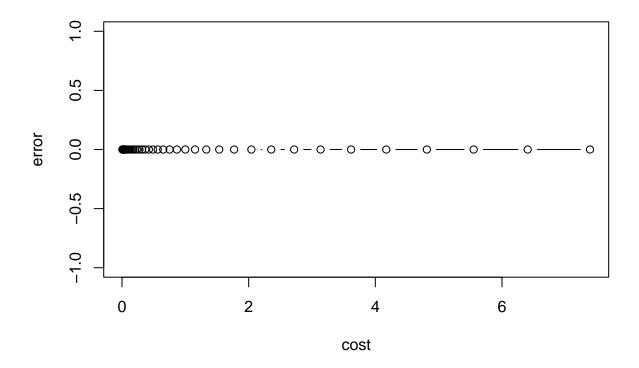
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Question 1: In this problem, we will apply support vector machines to predict whether a given car gets high or low gas mileage based on the dataset "auto.csv" (used in Homework 3; see Homework 3 for more details of the dataset). The response variable is mpg cat. The predictors are cylinders, displacement, horsepower, weight, acceleration, year, and origin. Split the dataset into two parts: training data (70%) and test data (30%).

```
auto = read.csv("auto.csv")
head(auto)
##
     cylinders displacement horsepower weight acceleration year origin mpg_cat
## 1
                         307
                                           3504
                                                         12.0
                                                                70
             8
                                     130
                                                                               low
## 2
             8
                         350
                                     165
                                           3693
                                                         11.5
                                                                70
                                                                         1
                                                                               low
## 3
             8
                         318
                                     150
                                           3436
                                                         11.0
                                                                70
                                                                         1
                                                                               low
## 4
             8
                         304
                                     150
                                           3433
                                                         12.0
                                                                70
                                                                               low
## 5
             8
                         302
                                     140
                                           3449
                                                         10.5
                                                                70
                                                                         1
                                                                               low
## 6
                         429
                                     198
                                           4341
                                                         10.0
                                                                70
set.seed(111111)
datSplit = initial_split(data = auto, prop = 0.7)
trainData = training(datSplit)
testData = testing(datSplit)
head(trainData)
     cylinders displacement horsepower weight acceleration year origin mpg_cat
##
## 1
             4
                         134
                                      95
                                           2515
                                                         14.8
                                                                78
                                                                               low
## 2
                         156
                                      92
                                           2585
                                                         14.5
                                                                82
                                                                         1
                                                                              high
## 3
             6
                         168
                                     120
                                           3820
                                                         16.7
                                                                76
                                                                         2
                                                                               low
                                                                79
## 4
             4
                         151
                                      90
                                           2670
                                                         16.0
                                                                         1
                                                                              high
## 5
             6
                         258
                                     110
                                           3632
                                                         18.0
                                                                74
                                                                         1
                                                                               low
## 6
                          98
                                      68
                                           2135
                                                         16.6
                                                                78
                                                                              high
trainData$mpg_cat = as.factor(trainData$mpg_cat)
testData$mpg_cat = as.factor(testData$mpg_cat)
```

(a): Fit a support vector classifier to the training data. What are the training and test error rates?

Performance of `svm'



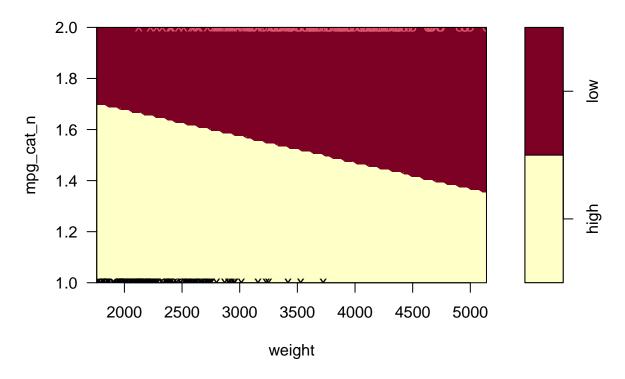
```
linear.tune$best.parameters

## cost
## 1 0.006737947

best.linear <- linear.tune$best.model
summary(best.linear)</pre>
```

```
## Call:
## best.svm(x = mpg_cat ~ ., data = trainData, cost = exp(seq(-5, 2,
       len = 50)), kernel = "linear", scale = TRUE)
##
##
## Parameters:
      SVM-Type: C-classification
   SVM-Kernel:
##
                linear
##
          cost: 0.006737947
##
## Number of Support Vectors: 108
##
   (53 55)
##
##
##
## Number of Classes: 2
##
## Levels:
## high low
pred.linear <- predict(best.linear, newdata = testData)</pre>
confusionMatrix(data = pred.linear,
reference = testData$mpg_cat)
## Confusion Matrix and Statistics
##
##
             Reference
## Prediction high low
##
         high
               59
##
         low
                 0 59
##
##
                  Accuracy: 1
##
                    95% CI: (0.9692, 1)
##
       No Information Rate: 0.5
##
       P-Value [Acc > NIR] : < 2.2e-16
##
##
                     Kappa: 1
##
   Mcnemar's Test P-Value : NA
##
##
##
               Sensitivity: 1.0
##
               Specificity: 1.0
##
            Pos Pred Value : 1.0
##
            Neg Pred Value: 1.0
                Prevalence: 0.5
##
            Detection Rate: 0.5
##
##
      Detection Prevalence: 0.5
##
         Balanced Accuracy: 1.0
##
##
          'Positive' Class : high
##
```

SVM classification plot



```
best.linear.model = linear.tune$best.model
test.pred = predict(best.linear.model, newdata = testData)
test.error = mean(test.pred != testData$mpg_cat)
cat("Test Data Error Rate:", test.error, "\n")
```

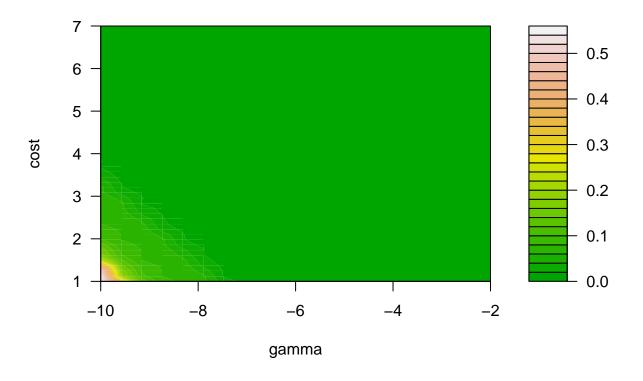
Test Data Error Rate: 0

```
train.pred = predict(best.linear.model, newdata = trainData)
train.error = mean(train.pred != trainData$mpg_cat)
cat("Training Data Error Rate:", train.error, "\n")
```

Training Data Error Rate: 0

(b): Fit a support vector machine with a radial kernel to the training data. What are the training and test error rates?

Performance of `svm'



```
radial.tune$best.parameters

## gamma cost
## 9 0.00131808 2.718282

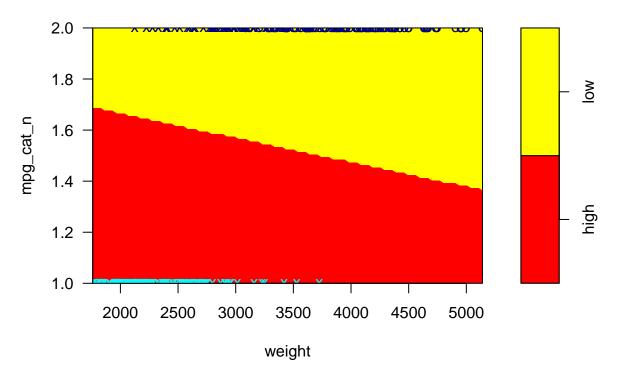
best.radial = radial.tune$best.model
summary(best.radial)

## Call:
## Call:
## best.svm(x = mpg_cat ~ ., data = trainData, gamma = exp(seq(-10,
```

```
-2, len = 20)), cost = exp(seq(1, 7, len = 50)), kernel = "radial")
##
##
##
## Parameters:
##
      SVM-Type: C-classification
   SVM-Kernel: radial
##
##
          cost: 2.718282
##
## Number of Support Vectors: 105
##
##
   (5253)
##
##
## Number of Classes: 2
##
## Levels:
## high low
pred.radial = predict(best.radial, newdata = testData)
confusionMatrix(data = pred.radial,
reference = testData$mpg_cat)
## Confusion Matrix and Statistics
##
##
             Reference
## Prediction high low
##
         high
               59
##
         low
                 0 59
##
##
                  Accuracy: 1
                    95% CI : (0.9692, 1)
##
##
       No Information Rate: 0.5
##
       P-Value [Acc > NIR] : < 2.2e-16
##
##
                     Kappa: 1
##
##
    Mcnemar's Test P-Value : NA
##
##
               Sensitivity: 1.0
##
               Specificity: 1.0
##
            Pos Pred Value : 1.0
##
            Neg Pred Value: 1.0
##
                Prevalence: 0.5
##
            Detection Rate: 0.5
      Detection Prevalence: 0.5
##
##
         Balanced Accuracy: 1.0
##
##
          'Positive' Class : high
##
plot(best.radial, trainData,
    mpg_cat_n ~ weight,
slice = list(
```

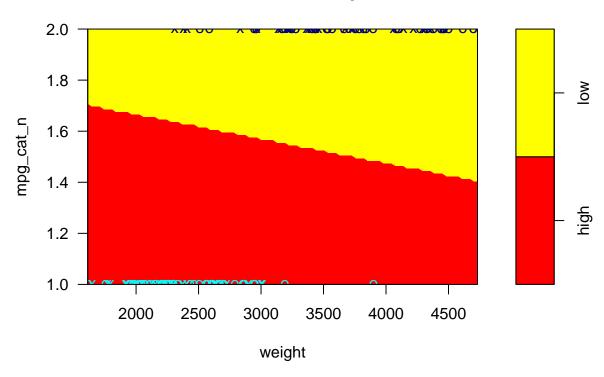
```
cylinders = median(trainData$cylinders, na.rm = TRUE),
    displacement = median(trainData$displacement, na.rm = TRUE),
    horsepower = median(trainData$horsepower, na.rm = TRUE),
    weight = median(trainData$weight, na.rm = TRUE),
    acceleration = median(trainData$acceleration, na.rm = TRUE),
    year = median(trainData$year, na.rm = TRUE),
    origin = median(trainData$origin, na.rm = TRUE)
),
grid = 100,
symbolPalette = c("cyan","darkblue"),
color.palette = heat.colors)
```

SVM classification plot



```
plot(best.radial, testData,
    mpg_cat_n ~ weight,
slice = list(
    cylinders = median(trainData$cylinders, na.rm = TRUE),
    displacement = median(trainData$displacement, na.rm = TRUE),
    horsepower = median(trainData$horsepower, na.rm = TRUE),
    weight = median(trainData$weight, na.rm = TRUE),
    acceleration = median(trainData$acceleration, na.rm = TRUE),
    year = median(trainData$year, na.rm = TRUE),
    origin = median(trainData$origin, na.rm = TRUE)
    ),
    grid = 100,
    symbolPalette = c("cyan", "darkblue"),
```

SVM classification plot



Question 2: In this problem, we perform hierarchical clustering on the states using the USArrests data in the ISLR package. For each of the 50 states in the United States, the dataset contains the number of arrests per 100,000 residents for each of three crimes: Assault, Murder, and Rape. The dataset also contains the percent of the population in each state living in urban areas, UrbanPop. The four variables will be used as features for clustering.

```
library(factoextra)
## Welcome! Want to learn more? See two factoextra-related books at https://goo.gl/ve3WBa
library(gridExtra)
```

##
Attaching package: 'gridExtra'

```
## The following object is masked from 'package:dplyr':
##
##
       combine
library(corrplot)
## corrplot 0.95 loaded
library(RColorBrewer)
library(gplots)
##
## Attaching package: 'gplots'
## The following object is masked from 'package:plotrix':
##
##
       plotCI
## The following object is masked from 'package:stats':
##
##
       lowess
library(jpeg)
data("USArrests")
str(USArrests)
## 'data.frame':
                  50 obs. of 4 variables:
## $ Murder : num 13.2 10 8.1 8.8 9 7.9 3.3 5.9 15.4 17.4 ...
## $ Assault : int 236 263 294 190 276 204 110 238 335 211 ...
## $ UrbanPop: int 58 48 80 50 91 78 77 72 80 60 ...
## $ Rape : num 21.2 44.5 31 19.5 40.6 38.7 11.1 15.8 31.9 25.8 ...
```

(a): Using hierarchical clustering with complete linkage and Euclidean distance, cluster the states. Cut the dendrogram at a height that results in three distinct clusters. Which states belong to which clusters?

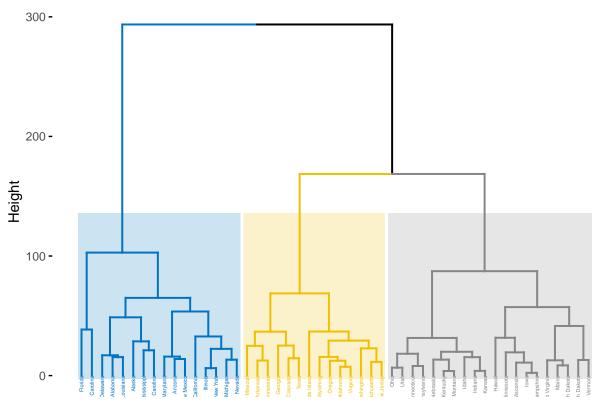
```
hc.complete <- hclust(dist(USArrests), method = "complete")
hc.average <- hclust(dist(USArrests), method = "average")
hc.single <- hclust(dist(USArrests), method = "single")
hc.centroid <- hclust(dist(USArrests), method = "centroid")

fviz_dend(hc.complete, k = 3,
    cex = 0.3,
    palette = "jco", # color scheme; other palettes:"npg", "aaas"...
    color_labels_by_k = TRUE,
    rect = TRUE, # whether to add a rectangle around groups.
    rect_fill = TRUE,
    rect_border = "jco",</pre>
```

labels_track_height = 2.5)

```
## Warning: The '<scale>' argument of 'guides()' cannot be 'FALSE'. Use "none" instead as
## of ggplot2 3.3.4.
## i The deprecated feature was likely used in the factoextra package.
## Please report the issue at <a href="https://github.com/kassambara/factoextra/issues">https://github.com/kassambara/factoextra/issues</a>.
## This warning is displayed once every 8 hours.
## Call 'lifecycle::last_lifecycle_warnings()' to see where this warning was
## generated.
```

Cluster Dendrogram



ind4.complete <- cutree(hc.complete, 3)
USArrests[ind4.complete == 1,]</pre>

##		Murder	Assault	UrbanPop	Rape
##	Alabama	13.2	236	58	21.2
##	Alaska	10.0	263	48	44.5
##	Arizona	8.1	294	80	31.0
##	California	9.0	276	91	40.6
##	Delaware	5.9	238	72	15.8
##	Florida	15.4	335	80	31.9
##	Illinois	10.4	249	83	24.0
##	Louisiana	15.4	249	66	22.2
##	Maryland	11.3	300	67	27.8
##	Michigan	12.1	255	74	35.1
##	Mississippi	16.1	259	44	17.1
##	Nevada	12.2	252	81	46.0
##	New Mexico	11.4	285	70	32.1

```
## New York
                     11.1
                               254
                                         86 26.1
## North Carolina
                     13.0
                               337
                                         45 16.1
                                         48 22.5
## South Carolina
                     14.4
                               279
USArrests[ind4.complete == 2,]
##
                  Murder Assault UrbanPop Rape
## Arkansas
                             190
                     8.8
                                        50 19.5
## Colorado
                     7.9
                             204
                                        78 38.7
                                        60 25.8
## Georgia
                    17.4
                             211
                                        85 16.3
## Massachusetts
                     4.4
                             149
## Missouri
                     9.0
                             178
                                        70 28.2
## New Jersey
                     7.4
                             159
                                        89 18.8
## Oklahoma
                     6.6
                             151
                                        68 20.0
## Oregon
                     4.9
                             159
                                        67 29.3
## Rhode Island
                     3.4
                                        87 8.3
                             174
## Tennessee
                    13.2
                             188
                                        59 26.9
## Texas
                    12.7
                             201
                                        80 25.5
## Virginia
                     8.5
                             156
                                        63 20.7
## Washington
                     4.0
                                        73 26.2
                             145
## Wyoming
                     6.8
                                        60 15.6
                             161
USArrests[ind4.complete == 3,]
```

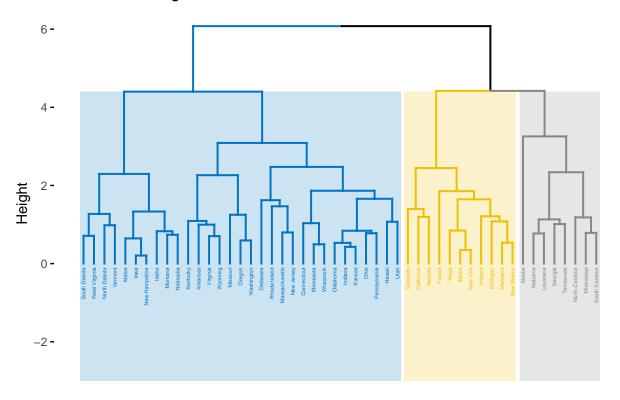
##	Murder	Assault	${\tt UrbanPop}$	Rape
## Connecticut	3.3	110	77	11.1
## Hawaii	5.3	46	83	20.2
## Idaho	2.6	120	54	14.2
## Indiana	7.2	113	65	21.0
## Iowa	2.2	56	57	11.3
## Kansas	6.0	115	66	18.0
## Kentucky	9.7	109	52	16.3
## Maine	2.1	83	51	7.8
## Minnesota	2.7	72	66	14.9
## Montana	6.0	109	53	16.4
## Nebraska	4.3	102	62	16.5
## New Hampshire	2.1	57	56	9.5
## North Dakota	0.8	45	44	7.3
## Ohio	7.3	120	75	21.4
## Pennsylvania	6.3	106	72	14.9
## South Dakota	3.8	86	45	12.8
## Utah	3.2	120	80	22.9
## Vermont	2.2	48	32	11.2
## West Virginia	5.7	81	39	9.3
## Wisconsin	2.6	53	66	10.8

(b): Hierarchically cluster the states using complete linkage and Euclidean distance, after scaling the variables to have standard deviation one. Does scaling the variables change the clustering results? In your opinion, should the variables be scaled before the inter-observation dissimilarities are computed?

Yes, clustering with scaling affects the number of states in each cluster-there are more states is cluster 3 when scaling compared to when there is not scaling. Yes, the variables be scaled before the inter-observation

dissimilarities are computed.

Cluster Dendrogram



```
ind4.scaled <- cutree(hc.complete.scaled, 3)
USArrests[ind4.scaled == 1, ]</pre>
```

##		Murder	${\tt Assault}$	UrbanPop	Rape
##	Alabama	13.2	236	58	21.2
##	Alaska	10.0	263	48	44.5
##	Georgia	17.4	211	60	25.8
##	Louisiana	15.4	249	66	22.2
##	Mississippi	16.1	259	44	17.1
##	North Carolina	13.0	337	45	16.1

```
## South Carolina 14.4 279 48 22.5
## Tennessee 13.2 188 59 26.9
```

USArrests[ind4.scaled == 2,]

##		${\tt Murder}$	${\tt Assault}$	UrbanPop	Rape
##	Arizona	8.1	294	80	31.0
##	California	9.0	276	91	40.6
##	Colorado	7.9	204	78	38.7
##	Florida	15.4	335	80	31.9
##	Illinois	10.4	249	83	24.0
##	Maryland	11.3	300	67	27.8
##	Michigan	12.1	255	74	35.1
##	Nevada	12.2	252	81	46.0
##	New Mexico	11.4	285	70	32.1
##	New York	11.1	254	86	26.1
##	Texas	12.7	201	80	25.5

USArrests[ind4.scaled == 3,]

##		Mundon	\	IIwhanDan	Dono
##	Arkansas	8.8	190	UrbanPop 50	19.5
##	Connecticut	3.3	110	77	
##	Delaware	5.9	238	72	15.8
##	Hawaii	5.3	236 46	83	20.2
##	Idaho	2.6	120	54	14.2
##	Indiana	7.2	113	65	21.0
##	Indiana Iowa	2.2	56	57	11.3
##	Kansas	6.0	115	66	18.0
##	Kentucky	9.7	109	52	16.3
##	Maine	2.1	83	51	7.8
##	Massachusetts	4.4	149	85	16.3
##	Minnesota	2.7	72	66	14.9
##	Missouri	9.0	178	70	28.2
##	Montana	6.0	109	53	16.4
##	Nebraska	4.3	102	62	16.5
##	New Hampshire	2.1	57	56	9.5
##	New Jersey	7.4	159	89	18.8
##	North Dakota	0.8	45	44	7.3
##	Ohio	7.3	120	75	21.4
##	Oklahoma	6.6	151	68	20.0
##	Oregon	4.9	159	67	29.3
##	Pennsylvania	6.3	106	72	14.9
##	Rhode Island	3.4	174	87	8.3
##	South Dakota	3.8	86	45	12.8
##	Utah	3.2	120	80	22.9
##	Vermont	2.2	48	32	11.2
##	Virginia	8.5	156	63	20.7
##	Washington	4.0	145	73	26.2
##	West Virginia	5.7	81	39	9.3
##	Wisconsin	2.6	53	66	10.8
##	Wyoming	6.8	161	60	15.6