Homework 9: Support Vector Machines

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Possible Solutions to Selected Questions

Important: This document contains R code solutions and example answers for problems posed in the DS740: Data Mining homework. We are intentionally sharing this document with learners who have already completed the associated homework. We want you to be able to review and troubleshoot your code, as well as ask questions to prepare you for later assessments. By using this document, you are accepting responsibility **not** to share it with anyone else, including other students in the course or the program who might not have completed the homework yet. By upholding this agreement, you are helping us use this tool with learners in future terms.

From Problem 1:

Question:

Which method do you expect will be better for categorizing the regions of species of oak trees: A support vector classifier or logistic regression? Explain.

Possible Answer: A support vector classifier, because there is little overlap between the groups.

Question:

Use caret to build a support vector classifier (a SVM with a linear kernel) to categorize the trees' regions based on their (unstandardized) logSize and logRange. Test the following values of cost: .001, .01, .1, 1, 5, 10, 100. Because of this data set's small size, use leave-one-out cross-validation. Enter your R code below.

Possible Answer:

Question:

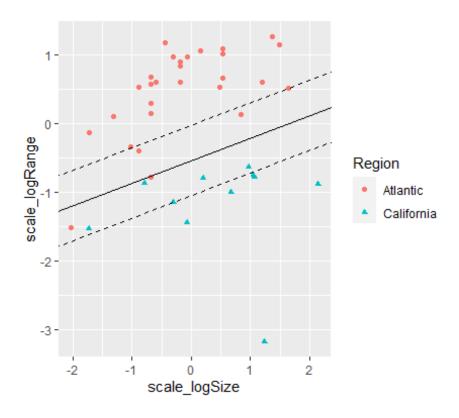
Make a graph showing the data points (similar to question 1), optimal hyperplane (line), and its margins. Include a legend.

- Use different colors and/or plotting characters to show the Regions of the points.
- You may use either the standardized or the unstandardized log(acorn size) and log(range).

Use *Insert -> Image* to upload your plot to this question on Canvas.

Possible Answer:

```
oak <- oak %>%
  tibble::rownames_to_column("Row") %>%
  mutate(is_SV = Row %in% attr(fit_oak$finalModel, "SVindex"))
b = attr(fit_oak$finalModel, "b")
coefs = attr(fit_oak$finalModel, "coef")[[1]]
oak SV <- oak %>%
  filter(is SV) %>%
  select(c(scale_logSize, scale_logRange)) %>%
  as.matrix()
#head(oak_SV) # Columns should be in the order x, y
              # relative to the graph
w = colSums(coefs * oak_SV) # beta_1, ... beta_p
oak %>%
  gf_point(scale_logRange ~ scale_logSize,
           color =~ Region,
           pch =~ Region) %>%
  gf_abline(intercept = b/w[2], slope = -w[1]/w[2]) %>%
  gf_abline(intercept = (b+1)/w[2], slope = -w[1]/w[2], lty = 2) %>%
  gf_abline(intercept = (b-1)/w[2], slope = -w[1]/w[2], lty = 2)
```



Question:

In question 3, we used caret to perform cross-validation for model selection (picking the best value of the cost parameter). In this question, we will use a *for* loop wrapper to perform a second layer of cross-validation. This will allow us to honestly assess the accuracy of our model-selection process.

- Set the random seed to 9.
- Create a vector to store the predicted regions (it should have length = the number of rows in the data set).
- Create vectors groups and cv_groups to perform 10-fold CV (like we did before learning about caret).
- Create a *for* loop to iterate through the folds of the outer layer of CV. Inside the *for* loop:
 - Create the variables groupii, train_set, and test_set, like we did for CV before learning about caret.
 - Use caret to perform an inner layer of LOOCV. caret should fit a support vector classifier and choose among costs of .001, .01, .1, 1, 5, 10, 100. This code can be the same as you used in question 3, except that it should use the training set from the outer layer of CV instead of the entire data set.
 - Use the model from caret to predict the regions of the data in the test set from the outer layer of CV.

Enter your R code below.

Possible Answer:

```
set.seed(9)
n = dim(oak)[1]
ngroups = 10 # 10-fold outer CV
groups = rep(1:ngroups, length = n)
cv_groups = sample(groups, n)
ctrl = trainControl(method = "LOOCV")
preds = vector(length = n)
best cost = numeric(length = ngroups)
for(ii in 1:ngroups){
 groupii = (cv groups == ii)
 train set = oak[!groupii, ]
 test_set = oak[groupii, ]
 data_used = train_set
 fit = train(Region ~ logSize + logRange,
              data = data_used,
              method = "svmLinear",
              tuneGrid = expand.grid(C = c(.001, .01, .1, 1, 5, 10, 100)),
              preProcess = c("center", "scale"),
              trControl = ctrl)
 best cost[ii] = fit$bestTune[[1]]
 preds[groupii] = predict(fit, newdata = test set)
}
best_cost
## [1] 100.0 0.1 1.0 1.0 1.0 1.0 1.0 1.0 0.1
```

From Problem 2:

Question:

Set the random seed to 9. Use caret to perform 10-fold cross-validation to compare different values of cost and sigma for a radial support vector machine. Use the same values of cost as listed previously: .001, .01, .1, 1, 5, 10, 100. Use sigma = 0.5, 1, 2, 3, and 4.

- Model the binary gas mileage variable as a function of all the other variables that are in Auto after question 10.
- Ask caret to model the probability that each point belongs to each category. (For purposes of this homework, it's OK if the model fails to converge.)

Enter your R code below.

Possible Answer:

Question:

Which combination of parameters gave the highest cross-validation accuracy?

```
fit_radial$bestTune
```

Question:

What was the cross-validation accuracy of the best model? Enter your answer to 4 decimal places.

```
fit_radial$results %>%
  filter(sigma == 1 & C == 1)
```

Question:

Use the best model to predict the probability that the following car would have **high** gas mileage:

1977 Chrysler Sunbeam

Cylinders: 4

Engine displacement: 132.5 cubic inches

Horsepower: 155 **Weight**: 2,910 lbs

Acceleration: 8.3 seconds **Origin**: American (1)

Enter your answer to 4 decimal places.

Question

Make a grid of example data points with

- weight = seq(min(Autoweight), max(Autoweight), length = 100)
- cylinders = 4
- origin = 1
- all other predictors set equal to their medians.

Predict the probability of having high gas mileage for each data point in the grid. Include your R code below.