

## WEEK 1 HOMEWORK

### INSTRUCTIONS

- Every learner should submit his/her own homework solutions. However, you are allowed to discuss the homework with each other (in fact, I encourage you to form groups and/or use the forums) – but everyone must submit his/her own solution; you may not copy someone else's solution.
- The homework will be peer-graded. In analytics modeling, there are often lots of different approaches that work well, and I want you to see not just your own, but also others.
- The homework grading scale reflects the fact that the primary purpose of homework is learning:

Rating	Meaning	Point value (out of 100)
4	All correct (perhaps except a few details) <u>with</u> a deeper solution than expected	100
3	Most or all correct	90
2	Not correct, but a reasonable attempt	75
1	Not correct, insufficient effort	50
0	Not submitted	0

### Question 2.1

Describe a situation or problem from your job, everyday life, current events, etc., for which a classification model would be appropriate. List some (up to 5) predictors that you might use.

### Question 2.2

The files `credit_card_data.txt` (without headers) and `credit_card_data-headers.txt` (with headers) contain a dataset with 654 data points, 6 continuous and 4 binary predictor variables. It has anonymized credit card applications with a binary response variable (last column) indicating if the application was positive or negative. The dataset is the “Credit Approval Data Set” from the UCI Machine Learning Repository (<https://archive.ics.uci.edu/ml/datasets/Credit+Approval>) without the categorical variables and without data points that have missing values.

1. Using the support vector machine function `ksvm` contained in the R package `kernlab`, find a good classifier for this data. Show the equation of your classifier, and how well it classifies the data points in the full data set. (Don't worry about test/validation data yet; we'll cover that topic soon.)

Notes on `ksvm`

- You can use `scaled=TRUE` to get `ksvm` to scale the data as part of calculating a classifier.

- The term  $\lambda$  we used in the SVM lesson to trade off the two components of correctness and margin is called C in ksvm. One of the challenges of this homework is to find a value of C that works well; for many values of C, almost all predictions will be “yes” or almost all predictions will be “no”.
- ksvm does not directly return the coefficients  $a_0$  and  $a_1 \dots a_m$ . Instead, you need to do the last step of the calculation yourself. Here’s an example of the steps to take (assuming your data is stored in a matrix called data):<sup>1</sup>

```
# call ksvm. Vanilladot is a simple linear kernel.
model <- ksvm(data[,1:10],data[,11],type="C-
svc",kernel="vanilladot",C=100,scaled=TRUE)
# calculate  $a_1 \dots a_m$ 
a <- colSums(model@xmatrix[[1]] * model@coef[[1]])
a
# calculate  $a_0$ 
a0 <- -model@b
a0
# see what the model predicts
pred <- predict(model,data[,1:10])
pred
# see what fraction of the model's predictions match the
actual classification
sum(pred == data[,11]) / nrow(data)
```

**Hint:** You might want to view the predictions your model makes; if C is too large or too small, they’ll almost all be the same (all zero or all one) and the predictive value of the model will be poor. Even finding the right order of magnitude for C might take a little trial-and-error.

**Note:** If you get the error “Error in vanilladot(length = 4, lambda = 0.5) : unused arguments (length = 4, lambda = 0.5)”, it means you need to convert data into matrix format:

```
model <-
ksvm(as.matrix(data[,1:10]),as.factor(data[,11]),type="C-
svc",kernel="vanilladot",C=100,scaled=TRUE)
```

2. You are welcome, but not required, to try other (nonlinear) kernels as well; we’re not covering them in this course, but they can sometimes be useful and might provide better predictions than vanilladot.
3. Using the k-nearest-neighbors classification function `kkn` contained in the R `kkn` package, suggest a good value of k, and show how well it classifies that data points in the full data set. Don’t forget to scale the data (`scale=TRUE` in `kkn`).

Notes on `kkn`

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<sup>1</sup> I know I said I wouldn’t give you exact R code to copy, because I want you to learn for yourself. In general, that’s definitely true – but in this case, because it’s your first R assignment and because the `ksvm` function leaves you in the middle of a mathematical calculation that we haven’t gotten into in this course, I’m giving you the code.

- You need to be a little careful. If you give it the whole data set to find the closest points to  $i$ , it'll use  $i$  itself (which is in the data set) as one of the nearest neighbors. A helpful feature of R is the index  $-i$ , which means "all indices except  $i$ ". For example, `data[-i, ]` is all the data except for the  $i$ th data point. For our data file where the first 10 columns are predictors and the 11<sup>th</sup> column is the response, `data[-i, 11]` is the response for all but the  $i$ th data point, and `data[-i, 1:10]` are the predictors for all but the  $i$ th data point.  
(There are other, easier ways to get around this problem, but I want you to get practice doing some basic data manipulation and extraction, and maybe some looping too.)
- **Note** that `kknn` will read the responses as continuous, and return the fraction of the  $k$  closest responses that are 1 (rather than the most common response, 1 or 0).