Statistical computing MATH10093 Computer lab 6

Finn Lindgren

6/3/2019

Summary

In this lab session you will develop code for cross validated model selection for basic climate estimation in a part of Scotland. You will not hand in anything, but you should keep your code script file for later use.

- 1. Complete lab 5 by reading the solutions!
- 2. Initialise lab 6: Open RStudio and either
 - (a) Open the relevant existing project for the lab, if you have one, or
 - (b) Create a new project for the folder where you want to store the lab files.
 - (c) Copy the TMINallobs.csv and TMINalltest.csv into the project folder, or a subdirectory data/ of that folder.
 - (d) Download and save the lab06code.R file to the same folder, and read the data (leave out data/ if you didn't use a subfolder for the data):

- 3. Write a function cvk_define_splits(N, K) that takes inputs
 - N: The number of data points in the Training&Validation set
 - K: The number of "splits" or "folds" for cross validation

The output should be a numeric vector of length N, containing indices $1, \ldots, K$ defining a random splitting into K subsets. The size of each

subset, N_k , should be equal to N/K if N is divisible by K. If N is not divisible by K they should have nearly equal size, for example

$$N_k = \lceil kN/K \rceil - \lceil (k-1)N/K \rceil, \quad k = 1, \dots, K.$$

The floor() and ceiling() functions implement $\lfloor \cdot \rfloor$ and $\lceil \cdot \rceil$. To ensure that all $N_k \geq 1$, we require $K \leq N$.

Theory task: Prove that this choice of N_1, \ldots, N_K sums to N.

Note: this function should be called only once in any cross validation procedure, since different random numbers will be generated each time it is called! The random numbers can be reset using the set.seed() function, but since that is a deterministic operation, that function should normally also be called only once (it is useful to call it once at the beginning of an R script to make the results repeatable and debugging easier).

Optional: Make the code give an error and helpful error message if K > N (with the stop() function).

The output should look something like this (with some random permutation)

```
# Should have 3 ones, 2 twos, 3 threes, and 2 fours:
cvk_define_splits(10, 4)

## [1] 3 4 3 4 1 1 1 2 3 2

# Error message on invalid inputs:
cvk_define_splits(N = 4, K = 10)

## Error in cvk_define_splits(N = 4, K = 10): K-fold cross validation
cannot have K > N
```

4. Write a function cvk_split(data, splits, k) that takes inputs

data: A data.frame with one observation per row

splits: The index vector defining the K data subsets, as produced by cvk_define_splits

k: The index $k \in 1, 2, ..., K$ for the subset to be used as validation set

The output should be a list with two named elements:

train: A data frame with all rows of data in subsets $j \neq k$

valid: A data frame with all rows of data in the kth subset

Test the function:

```
mysplits <- cvk_define_splits(nrow(TMINallobs), 10)
data1 <- cvk_split(TMINallobs, mysplits, 1)</pre>
```

Compare the full data with the subsample:

```
# In R, to view the first rows of the data frames:
head(TMINallobs)
head(data1$train)
```

```
# In Rmd or Rnw, with library(xtable) and code chunk option results="asis":
print(xtable(head(TMINallobs)), size = "\\scriptsize")
print(xtable(head(data1$train)), size = "\\scriptsize")
```

5. Look at the file lab06code.R and familiarise yourself with the function train_and_validate() and the other functions, that are all related to Coursework A. Check that the following gives the expected output (if you recall the approximate scores from Q1 in Coursework A):

6. Write a function cvk_do_all(data, splits, formula) that calls train_and_validate() for each of the K splits and collects the results. The inputs should be:

```
data: The data frame to be split into Training and Validation subsets splits: The splitting information, e.g. as produced by cvk_define_splits formula: A model formula suitable for lm()
```

The output should be a data.frame with K cross validated scores in each column for columns named SE, DS, and Brier. Hints:

To find K from the splits parameter, use $K \leftarrow \max(\text{splits})$ There are many ways to iterate over the k-values. One option is a for-loop:

```
results <- ... # preallocate storage for the results
for (k in 1:K) {
    # do something and store in results, appropriately indexed by k
}
# Post-process the results</pre>
```

Test your function with e.g. this:

7. The following function (copy the code from the end of lab06code.R) is meant to construct a formula with max_order harmonic functions. Unfortunately, it fails to handle the case max_order = 0, when no harmonic functions should be included. Modify the code to fix this problem. Hint: One option is to use if (...) {...} else {...}.

```
# See the code for the function:
make_formula
# Try the function
```

make_formula(2)
make_formula(0)

8. Use cvk_do_all() to obtain cross validated prediction scores for models of the type in Quesion 2 of Coursework A.

Evaluate cross-validated scores for a range of models with $\mathtt{max_order} \in \{0,1,2,3,4,5\}$. Compare the scores for successive models and attempt to detect when adding harmonics no longer has a clear effect. Do the three different types of scores agree? Also compare with the scores for your favourite model when estimating with all of TMINallobs and validating with TMINalltest.

Note: If you calculate the basic variance estimate from Lecture 6 (with $\frac{1}{K(K-1)}$, corrected in the pdf after the lecture) it seems either the variance is underestimated, or there is bias in the score estimates; likely due to the dependence between the scores and between cross-validation folds, as discussed in the lecture. A more conservative estimate might instead be to use $\frac{1}{K-1}$, which could be motivated as the approximate upper bound if the cross-validated scores were perfectly correlated. However, even this might not reflect the true uncertainty.

Hint: Part of your code might look something like this:

```
S0 <- cvk_do_all(TMINallobs, the_splits, make_formula(0))
S1 <- cvk_do_all(TMINallobs, the_splits, make_formula(1))
S1 - S0
```

9. Optional: Do another analysis of the same models, but with a leave-station-out cross validation approach. Write a cvk_define_splits_lso(groups) function to define the data splitting based on the values in groups, which should be TMINallobs\$ID when you call the function. Hints:

Consider a combination of as.factor() and as.numeric(). By using this function, the rest of the code can remain intact!