

Exercises, STK4040, week 42

October 11, 2007

Exercise 1

Consider the linear model $\mathbf{y} = \mathbf{X}\beta + \varepsilon$, where $\varepsilon \sim N_n(\mathbf{0}, \sigma^2 \mathbf{I}_n)$. We assume that the $n \times (p + 1)$ matrix \mathbf{X} has full rank $p + 1 < n$, and that the first column of \mathbf{X} is $\mathbf{1}_n$.

The least squares estimate of β is $\hat{\beta} = (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{y}$. The predicted response values (A.K.A. fitted values) are $\hat{\mathbf{y}} = \mathbf{X} \hat{\beta}$. Let $\mathbf{H} = \mathbf{X}(\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T$. Then the predicted response values can be written as $\hat{\mathbf{y}} = \mathbf{H} \mathbf{y}$. (\mathbf{H} is often called the ‘hat matrix’, because it transforms \mathbf{y} into $\hat{\mathbf{y}}$.) The residuals are given by $\hat{\varepsilon} = \mathbf{y} - \hat{\mathbf{y}} = (\mathbf{I} - \mathbf{H}) \mathbf{y}$. Define $\mathbf{M} = \mathbf{I} - \mathbf{H}$. Then $\hat{\varepsilon} = \mathbf{M} \mathbf{y}$.

Prove the following:

1. \mathbf{H} is idempotent (i.e., $\mathbf{H}\mathbf{H} = \mathbf{H}$) and symmetric.
2. \mathbf{M} is idempotent and symmetric.
3. $\mathbf{M}\mathbf{X} = \mathbf{0}$, and $\mathbf{M}\mathbf{1}_n = \mathbf{0}$.
4. $\hat{\varepsilon} = \mathbf{M}\varepsilon$, $\mathbf{X}^T \hat{\varepsilon} = \mathbf{0}$, and $\sum_{i=1}^n \hat{\varepsilon}_i = 0$.
5. $\hat{\mathbf{y}}^T \hat{\varepsilon} = 0$.
6. $\mathbf{1}_n^T \hat{\varepsilon} = 0$.

Exercise 2

Given the linear model $\mathbf{y} = \mathbf{X}_1 \beta + \mathbf{X}_2 \varphi + \varepsilon$, where $\varepsilon \sim N_n(\mathbf{0}, \sigma^2 \mathbf{I}_n)$, and $\mathbf{X} = (\mathbf{X}_1, \mathbf{X}_2)$ has full rank.

Show that $\hat{\beta}$ can be calculated by first regressing \mathbf{y} and \mathbf{X}_1 onto \mathbf{X}_2 , and then regress the residuals of \mathbf{y} onto the residuals of \mathbf{X}_1 .

Exercise 3

The file `dataset_42.txt` contains sensory data from an experiment in which 105 persons tasted 6 different cheeses. Each person has given each cheese an integer score between 1 and 9, denoting how well they liked the cheese (9 is best and 1 is worst).

The file contains a variable `Code`, giving the sex of the person (A for female and B for male), and one variable for each cheese: `A_Cow_Full_fat`, `C_Cow_Full_fat`, `D_Cow_Low_fat`, `E_Cow_Low_fat`, `G_Buffalo_Full_fat`, and `I_Buffalo_Full_fat`. The cheeses A, C, D, and E are made from cow milk, while G and I are made from buffalo milk. The cheeses A, C, G, and I have regular fat content, while D and E are low fat cheeses.

Do a principal component analysis of the liking scores. How many components are important? Plot loadings and scores, and see if you can find any patterns. Tip: the data can be read in and set up in R like this:

```
tmp <- read.table("dataset_42.txt")
names(tmp)
Code <- tmp$Code
X <- as.matrix(tmp[,2:7])
```

Plot the scores with codes and/or colours to denote sex. Do you see a pattern now? Interpret the loadings. (Tip: You can plot the codes by using the argument `pch = as.character(Code)` in the plot function, and colours can be added with the argument `col = as.numeric(Code)`.)