

R Programming: Worksheet 4

By the end of today you should be able to plot one and two-dimensional data; generate plots from scratch; add points, lines, blocks and legends;

1. Time Series

Load the data `beav1` from the `MASS` package, and look at the field `temp` (ignore the other fields for now).

- Using the function `ma3()` which you constructed on Sheet 3, calculate the moving average of length 3 for the `temp` data.
- Plot the original temperature data [use `plot()` with the option `type="l"`, as we did for the random walk on sheet 1.]
- Try plotting the moving average of length 3 on the same graph. First plot the original data as usual. Then, use the `points()` function to add on the second vector:

```
> n <- length(beav1$temp)
> points(2:(n - 1), ma3(beav1$temp), type = "l", col = 2)
```

- Write a second function `plotMA` which takes two vector arguments, `x` and `k`, and (i) calculates the moving averages of length `k` for each entry in `k`; (ii) plots them on top of each other as above.
Ideally, each line should be a different colour.

2. ChickWeight Data

Take a look at the `ChickWeight` data.

```
> ?ChickWeight
```

Do some exploration of the different fields and their qualities. For parts (a)-(e), use the base R function only (i.e. without loading the `lattice` package). If you want to unattach a package, use the following command:

```
> detach(package:lattice)
```

- Do a basic scatter plot of the logarithm of chick-weight against age. What are the limitations of this?
- Use colour to distinguish between the different diets.
- What is the range of log-weights and ages? [I'll let you guess the command for this one.]
- Using this, construct a new (blank) plot window with these ranges, and add x- and y-axes.
- Now plot an individual line for each chick's measurements, with colour corresponding to the diet they were given.
Do you see any further shortcomings in this plot?
- Can you achieve the same plot using `lattice`?
- Export your final plot to a PDF file.

3. Lynxes and Hares

I have sent you a file called `hares.dat`; it contains data on the annual number of lynx and hare pelts recovered by the Hudson's Bay Company in Canada. Save the file somewhere appropriate and read it into R as a data frame.

Produce a single plot of these data, consisting of a line for each species (which should be different colours). Make sure that:

- (a) the y -axis starts at 0.
- (b) both axes are correctly labelled and your plot has a title;
- (c) you provide a legend to label your series.

4. Binning Data*

Write (from scratch) your own function to produce histograms, using `cut()` and `table()`.