

1. Cystic Fibrosis dataset

On the class website you will find the file `cystfibr.txt` which contains a set of measurements on a set of individuals with cystic fibrosis. Save the file locally onto your computer.

- (a) Take a look at this file using a text editor like Wordpad or Notepad. Read the data into R using `read.table()`; call the resulting dataframe `cf`.
Check that the data have been formatted correctly by typing:

```
> head(cf)
```

- (b) Obtain the following:
- (i) the number of individuals in the dataset;
 - (ii) the number of variables measured on each individual;
 - (iii) a vector of the names of the variables measured on each individual;
 - (iv) the mean, median, standard deviation and range of each of the variables (use `apply()`);
- (c) Create the following data frames and for each one calculate the mean of each of the variables
- (i) a new data frame containing just individuals older than 15
 - (ii) a new data frame containing just individuals with `bmp` in the interval `[70,90]`
 - (iii) a new data frame containing just individuals with either `FEV1` greater than 30 or `RV` greater than 300.

2. Data Frames

Load the `MASS` library and take a look at the dataset called `survey`.

```
> library(MASS)
> head(survey)
```

You can look at the documentation:

```
> ?survey
```

and get a brief summary of each variable:

```
> summary(survey)
```

- (a) Find the mean pulse rate of the students. What goes wrong here?

The vector of pulses stored in `survey` contains some entries which are labelled `NA`. This is used in R to represent **missing data**.

- (b) Try looking at the documentation for `mean()` to see how to get around this.
- (c) The ages are recorded as fractions representing a number of months. Change that columns of the data frame so that it contains whole years (the `floor()` command may be useful).

(d) Find the mean pulse rate for students under 20.

Subsetting. Suppose I want to obtain the records of students who are over 190 cm tall. Since data frames allow me to subset just like a matrix, I might just think of typing:

```
> survey[survey$Height > 190, ]
```

What goes wrong here? Try instead the following:

```
> subset(survey, Height > 190)
```

The subset function ignores missing values, which is usually the behaviour we would prefer. We can also select only some of the fields of the data frame if we prefer:

```
> subset(survey, Height > 190, select = c("Pulse", "Clap"))
```

Recall that the & operator does a point-wise logical ‘and’ comparison.

```
> subset(survey, (Pulse > 70) & (Smoke == "Heavy"))
```

Similarly, | is for ‘or’, and ! for ‘not’.

```
> with(survey, (Pulse > 70) | (Pulse < 45))
> !(survey$Age > 30)
```

(e) Find the mean age of students who write with their right hand.

(f) What proportion of left handers do not clap with their left hand on top?

(g) Using the plot() command, plot the pulse of the subjects against their age.

Try subtracting 10 from the age and taking the logarithm (using the function log()), to obtain a slightly clearer picture.

3. Factors

Take a look at the birthwt data from the MASS package.

(a) How is race stored in these data? Is this sensible?

(b) Turn this into a factor with level names as indicated in the documentation. To do this, look at the documentation for the function factor().

(c) Use the table() command to count the number of babies of each race.

(d) Try the following command:

```
> tab = with(birthwt, table(smoke, low))
```

What do the results in tab suggest?

4. ***Gaussian Random Walk.** Write a function which simulates a Gaussian random walk with n steps. (In other words, $X_0 = 0$ and $X_i - X_{i-1} \sim N(0, 1)$ independently.)

Generate and plot the walk for n = 1000 (use type="l").

Try to vectorise your code (if it is already vectorised, try constructing a naïve version with a for() loop). Compare the speed of the vectorised and unvectorised versions by generating a random walk of length 50,000