

Computing for Statistical Genetics

Session 1: Introduction

Before you start: the class site is

`http://faculty.washington.edu/kenrice/sisg`

- it will be helpful to bookmark this link. We also recommend logging on to the internet *before* you start your R session.

1. The file `http://faculty.washington.edu/kenrice/sisg/example-pheno.csv` has information on blood pressure and related variables. Read in the file and summarize the variables.
2. The file `http://faculty.washington.edu/kenrice/sisg/example-pheno.txt` has the same data as above. Download and read it in and check that it is the same data as in Q1.

Hint: note that these files are not in identical formats! You will need to use different code to read them in.