# Introduction to Genome-wide Association Studies (GWAS)

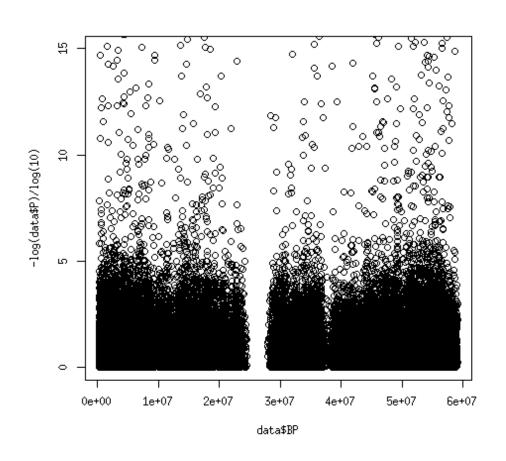
**Practical** 

Part 1: discovery

Teresa Ferreira and Nilufer Rahmioglu

Wellcome Centre for Human Genetics/Big Data Institute

### chr19-example data

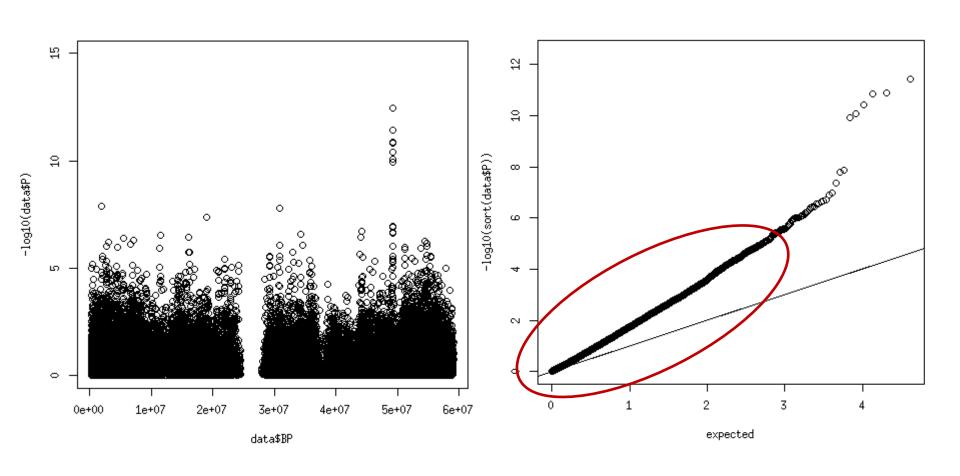


#### QC

- Missingness per indiv > 0.02
- Missingness per marker > 0.01
- Heterozygosity <0, >0.2
- HWE p<1x10<sup>-04</sup>
- MAF < 1%

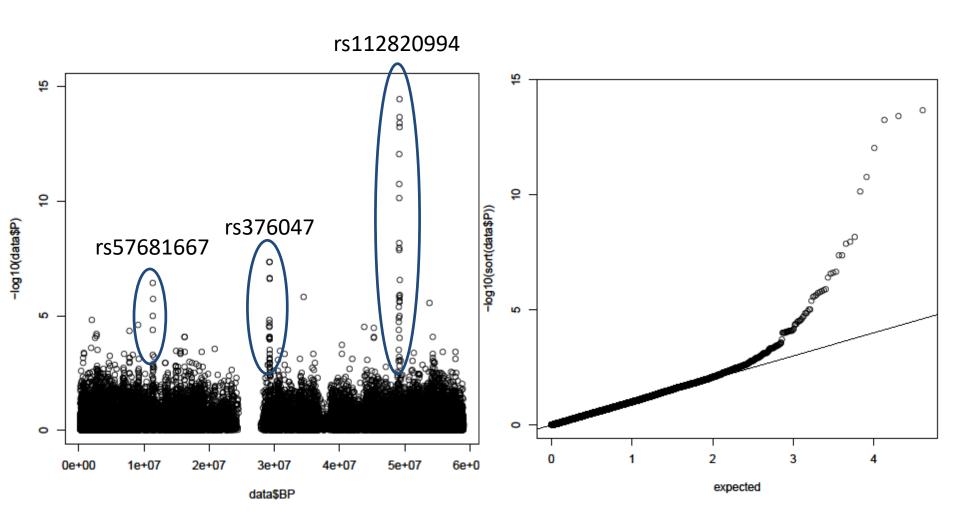
## chr19-example data

#### Cleaned data

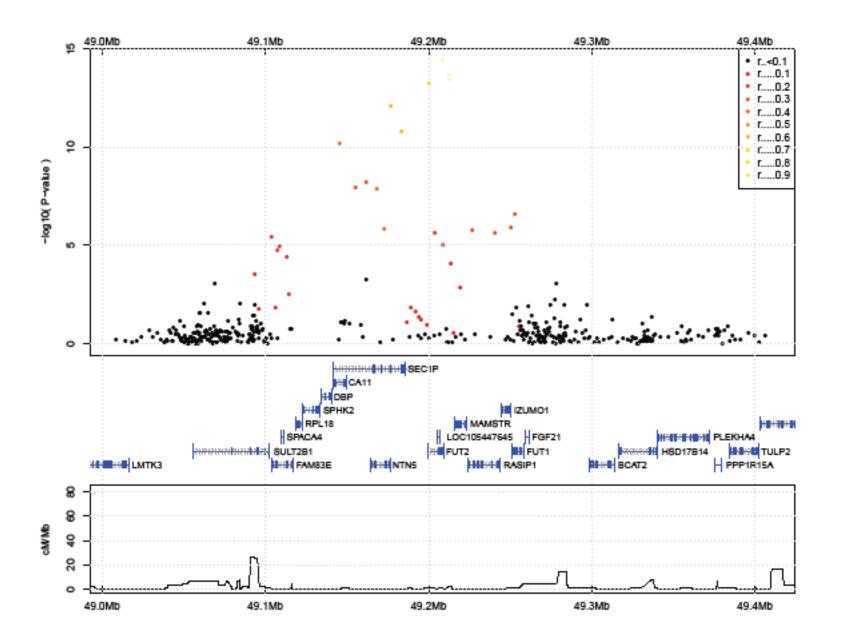


Population stratification?

## Adjusting for PCs



#### rs112820994



#### rs112820994

