## **Preliminary Reading List**

'Bioinformatics: A Practical Guide to the Analysis of Genes and Proteins', A. Baxevanis and B.F. F. Ouellette.

'Genomes 3', T.A. Brown

(2<sup>nd</sup> edition available online: <a href="http://www.ncbi.nlm.nih.gov/books/NBK21128/">http://www.ncbi.nlm.nih.gov/books/NBK21128/</a>)

For some of the practical sessions in R, we will make good use of the statistical programming environment, <a href="https://www.r-project.org">www.r-project.org</a>

Online course in R: https://www.coursera.org/course/rprog

A good book to get you started in R: 'Introduction to Scientific Programming and Simulation with R', Owen Jones, Robert Maillardet and Andrew Robinson

'Computational Genome Analysis', Richard Deonier, Simon Tavaré and Michael Waterman.

The following titles are recommended maths textbooks for those of you who are coming from a biological background and want to brush up on the mathematics:

'Applied Statistics for Engineers and Physical Scientists', Johannes Ledolter and Robert Hogg,

(Chapters 2, 3 and 4)

'Statistical Methods in Bioinformatics: An Introduction', Warren Ewens and Gregory Grant

(Chapters 1, 3, 8 and 9)

'Introductory Statistics with R', Peter Dalgaard, (Chapter 2)

'Biological Sequence Analysis' Richard Durbin et al. (first few chapters)

MIT have produced a 'statistics and probability primer' for computational biologists at: <a href="http://tinyurl.com/compbiostatsprimer">http://tinyurl.com/compbiostatsprimer</a>