Bioconductor

Summary
Bioconductor is a set of scripts written in the R statistical programming language, useful for calculations on and visualization of biological data.

Available on
- Fourierseq
- Phylocluster (with modules)
  - affy, affydata, affyPLM, annafy, annotate, Biobase, Biostrings, DynDoc, gcrma, genefilter, geneplotter, hgu95av2.db, limma, marray, matchprobes, multtest, ROC, von, xtable, affyQCReport, makecdfenv

User documentation
- To get started using Bioconductor, check out the Bioconductor Getting Started page. For an overview, go to the main Bioconductor page.

Helpful tips
- Start by learning R, which is quite different from most structured programming languages. It is similar to SAS.