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, annaffy, 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.