

Objective:

Looking to volume and complexity of biological data, advanced programming skills are required for researchers in order to get the most out of their data analyses. This course will provide the expanded programming skills necessary to develop software that can exploit the complex information landscape of bioinformatics.

Theory:**UNIT I**

Introduction to Molecular Biology, Central Dogma, Sequencing technologies, etc., Introduction to different programming languages used in biological sciences, Perl, Python, R, Ruby etc.

UNIT II

Unix/Linux shell programming, Perl: Introduction, Scalar Data, Arrays and list Data, Control Structures, Regular Expressions; Subroutines, File handles and file tests; Function; formats, Directory Access, Process management, Other data Transformation; Finding a sub string, Extracting and replacing a sub string. Bio-Perl: Introduction, Architecture, Uses of bioperl, Introduction to python/ Bio-Python

UNIT III

R: open-source language and environment for statistical computing and graphics.

UNIT IV

Bioconductor - open-source software for the analysis of biomedical and genomic data, mainly R packages. Analysis of microarray data, Next Generation Sequencing (NGS) data etc.

UNIT V

Parallel programming/ Computing – MapReduce, Interfaces with other languages: C, Java, Perl, Python, XML, etc. – Omega Project (www.omegahat.org), Formatting Data: sorting, Transliteration Database manipulation: DBM Databases and DBM Hashes, Opening and closing DBM Hashes, Fixed-Length Random –Access Databases, Variable-Length (Text) Databases, Win32 Databases Interfaces.

Practical:

1. Unix/Linux shell scripting
2. Active perl-
3. R, Map reduce

Reference Books:

1. The R Project for Statistical Computing [<http://www.R-project.org/>].
2. Perl.org [<http://www.perl.org/>].
3. Python Programming Language [<http://www.python.org/>].
4. Ruby Programming Language [<http://www.ruby-lang.org/>].
5. Tisdall J (2001) Beginning perl for bioinformatics. O'Reilly Media, Inc.
6. Tisdall JD (2003) Mastering perl for bioinformatics. O'Reilly Media, Inc.
7. Kinser J (2008) Python for bioinformatics. Jones & Bartlett Publishers.
8. Model M (2009) Bioinformatics programming using python. O'Reilly Media, Inc.
9. JRuby [<http://jruby.codehaus.org/>].
10. Randal L. Schwartz, Tom Phoenix, Brian d foy. 2008. Learning Perl. O-Reilly.
11. Sriram Srinivasan. 1997. Advanced Perl Programming. O-Reilly.
12. Tim Bunce and Alligator Descartes. 2000. Programming the perl DBI. O-Reilly.