

V. Outline of Course Content

- A. Understanding gene structure and function
 - 1. Storage and expression of genetic information
 - 2. Structure and function of genomes
 - 3. Molecular structure of eukaryotic and prokaryotic genes

- B. Data search and sequence comparisons
 - 1. Sequence alignment methods
 - 2. Database searching methods
 - 3. Multiple sequence alignments

- C. Structure Prediction
 - 1. Basic concepts
 - 2. Methods for secondary structure prediction
 - 3. Tertiary and quaternary structure
 - 4. Protein folding models and structure prediction

- D. Substitution patterns and phylogenetics
 - 1. Substitution patterns
 - 2. Molecular phylogenetics

- E. Genomics
 - 1. Prokaryotic genomes and gene recognition
 - 2. Eukaryotic genomes and gene recognition
 - 3. Gene expression analysis and microarrays

- F. Proteomics
 - 1. Proteome, Protein Classification and techniques
 - 2. Post-translational modification and other predictions
 - 3. Ligand screening and drug design

VI. Educational Goals and Learning Outcomes

A. Educational Goals

Students will:

- 1. Develop an ability to apply principles and generalizations already learned about science and technology to new problems and situations. (GE-RVCC 1; NJ GE 3)
- 2. Learn terms and facts of bioinformatics.
- 3. Learn concepts and theories of bioinformatics. (GE-RVCC 3, 7; NJ GE 2)
- 4. Develop an ability to synthesize and integrate information and ideas. (GE-RVCC 3, 7; NJ GE 2)
- 5. Analyze and interpret data (information) and draw logical conclusions. (GE-RVCC 3, 7; NJ GE 2)
- 6. Apply mathematical and computing skills to the solution of problems. (G.E. 7)
- 7. Improve computer skills. (GE-RVCC 3; NJ GE 4)
- 8. Develop an informed understanding of the role of science and technology in society. (GE-RVCC 5; NJ GE 9)

B. Learning Outcomes

Students will be able to:

- 1. Describe the field of bioinformatics.

2. Describe the fundamental concepts of bioinformatics and its applications to genomics and proteomics.
3. Define the structure and applications of genetic, nucleic acid and protein databases.
4. Search, retrieve and format information from biological databases.
5. Perform a variety of bioinformatics analyses using biological databases.

VII. Modes of Teaching and Learning

- A. Lecture/discussion
- B. Laboratory in an Internet equipped computer lab
- C. Internet resources
- D. Handouts prepared from the latest articles relevant to Bioinformatics
- E. Scientific articles and news
- F. Guest speakers

VIII. Papers, Examinations, and other Assessment Instruments

- A. Unit exercises
- B. Small-group projects
- C. Individual projects
- D. Participation in class/small group discussion

IX. Grade Determinants

- A. Unit exercises
- B. Small-group projects
- C. Individual projects

X. Texts and Materials

- A. Suggested Textbook
 1. Krane, D.E. and Raymer, M.L. (2003) *Fundamental Concepts of Bioinformatics*. Benjamin Cummings.
- B. Supplementary Texts
 2. Baxevanis, A.D. and Ouellette, B.F. (2001) *Bioinformatics: A Practical Guide to the Analysis of Genes and Proteins, 2nd Ed.* John Wiley and Sons, Inc.
- C. Current event articles
- D. Web resources
- E. Computer-based resources

(Please Note: The course outline is intended only as a guide to course content and resources. Do not purchase textbooks based on this outline. The RVCC Bookstore is the sole resource for the most up-to-date information about textbooks.)

XI. Resources

The library has the resources necessary for students to complete assignments necessary for this course.

XII. Honors Options

This is not an honors course.

