

CBM003 ADD/CHANGE FORM

Undergraduate Council
 New Course Course Change
 Core Category: NONE Effective Fall 2007

or

Graduate/Professional Studies Council
 New Course Course Change
 Effective Fall

RECEIVED OCT 13 2006

APPROVED JAN 24 2007

1. Department: ET College: TECH
2. Person Submitting Form: Rupa Iyer Telephone: 713-743-4076
3. Course Information on New/Revised course:
 - Instructional Area / Course Number / Long Course Title:
BTEC / 3301 / Principles Of Genomics/Proteomics And Bioinformatics
 - Instructional Area / Course Number / Short Course Title (30 characters max.)
BTEC / 3301 / GENOMICS/PROTEOMICS & BIOINFO
 - SCH: 3.00 Level: JR CIP Code: 2612010002 Lect Hrs: 3.0 Lab Hrs: 0
4. Justification for adding/changing course: To provide for new discipline areas
5. Was the proposed/revised course previously offered as a special topics course? Yes No
 If Yes, please complete:
 - Instructional Area / Course Number / Long Course Title:
 / /
 - Content ID: Start Date (yyyy3):
6. Is this course offered for undergraduate credit only? Yes No
7. Authorized Degree Program(s): BS, Biotechnology
 - Does this course affect major/minor requirements in the College/Department? Yes No
 - Does this course affect major/minor requirements in other Colleges/Departments? Yes No
 - Are special fees attached to this course? Yes No
 - Can the course be repeated for credit? Yes No
8. Grade Option: Letter (A, B, C ...) Instruction Type: lecture
9. If this form involves a change to an existing course, please obtain the following information from the course inventory: Instructional Area / Course Number / Long Course Title
 / /
 - Start Date (yyyy3): Content I.D.:
10. Proposed Catalog Description:
 Cr: (3-0) Prerequisites: BIOL 2333/2133, 3301, and ITEC 2334 Description (30 words max.): Overview of the fields of bioinformatics and genomics. Topics, tools, issues and current trends in these and related fields will be discussed.

11. Dean's Signature:  Date: 10/12/06
 Print/Type Name: Fred D. Lewallen

University of Houston
Proposed Course Outline for BTEC 3301, Introduction to
Genomics/Proteomics and Bioinformatics

Course Objectives: Students who successfully complete this course will be able to:

- Describe genomics, proteomics and related fields and their application in predication of structure and function
- Analyze gene expression and interpret its significance
- Familiar with topics, tools, issues and current trends in these and related fields.
- Recognize the utility of bioinformatics applications in proteomics data analysis

Course Outline

- 1. Prokaryotic genome**
 - a. Gene Structure
 - b. Gene Density
 - c. G- C Content
- 2. Eukaryotic Genome**
 - a. Gene Structure
 - b. Open Reading Frames
 - c. GC- content
 - d. Gene Expression
 - e. Transposition
 - f. Gene Density
- 3. Genomics**
 - a. Prokaryotic genomes and gene recognition
 - b. Eukaryotic genomes and gene recognition
- 4. Protein Folding**
 - a. Polypeptide**
 - i. Secondary Structure
 - ii. Tertiary and Primary structure
 - iii. Structure prediction
 - iv. Predicting RNA secondary structures.
- 5. Proteomics**
 - a. Protein Classification
 - b. Experimental Techniques
 - c. Inhibitors and Drug Design
 - d. Ligand Screening
 - e. X- ray crystal Structures
 - f. Empirical methods and prediction techniques
 - g. Posttranslational modification and prediction
- 6. Proteomics**
 - a. Protein Classification

- b. Experimental Techniques
- c. Inhibitors and Drug Design
- d. Ligand Screening
- e. X- ray crystal Structures
- f. Empirical methods and prediction techniques
- g. Posttranslational modification and prediction

7. Proteomics

- a. Protein Classification
- b. Experimental Techniques
- c. Inhibitors and Drug Design
- d. Ligand Screening
- e. X- ray crystal Structures
- f. Empirical methods and prediction techniques
- g. Posttranslational modification and prediction

8. Introduction to programming and data structures

- a. The basics
- b. Program control
- c. Readability
- d. Data structures
- e. Input and output

Recommended Text: Krane, D. E. and Raymer, M. L. (2003) Fundamental Concepts of Bioinformatics. Benjamin Cummings.