

朝陽科技大學 094學年度第2學期教學大綱  
Computational Biology 計算生物學

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| 當期課號   | 7245   | Course Number      | 7245  |
| 授課教師   | 陳永福  | Instructor         | CHEN,YUNG FU  |
| 中文課名   | 計算生物學  | Course Name        | Computational Biology   |
| 開課單位   | 資訊管理系碩士班二A   | Department         |   |
| 修習別    | 選修   | Required/Elective  | Elective  |
| 學分數    | 3  | Credits            | 3   |
| 課程目標   | 人類基因組是由大約三十億鹼基對(A, T, C, G)所組成。將這些資料的順序正確排列就是基因定序。而這只是一個開端而已，隨之而來的是更大量的分析資料，在這大量的資料裡，有用的資料卻是隱藏在許多雜訊之中，計算生物學領域就是要找出有用的資訊，其主要的重點是有關如何儲存、管理、傳輸、進而分析生物相關的大量資訊。   | Objectives         | 1.Sequence Alignment<br>2.Evolutionary Trees<br>3.Superstrings<br>4.Protein Structure<br>5.System Biology<br>6.String Matching<br>7.Superstructures<br>8.RNA Structures<br>9.Genome Rearrangement<br>10.Pattern Discovery<br>11.Divide-and-Conquer<br>12.Sorting by Reversal<br>13.Visual Display   |
| 教材     | Bioinformatics and Functional Genomics by J. Pevsner, Wiley-Liss 2003  | Teaching Materials | Bioinformatics and Functional Genomics by J. Pevsner, Wiley-Liss 2003   |
| 成績評量方式 | 1. 出席：10%<br>2. 作業：30%<br>3. 期中考試：30%<br>4. 期末報告：30%   | Grading            | 1.Attendance:10%<br>2.Homework: 30%<br>3.Mid-Term Exam:30%<br>4.Final Exam:30%  |
| 教師網頁   | <a href="http://www.csie.dyu.edu.tw/~yfbchen/">www.csie.dyu.edu.tw/~yfbchen/</a>   |                    |   |
| 教學內容   | Chap 0: Cell Function, Protein and DNA (細胞功能、蛋白質、DNA)、Transcription and Translation (基因之轉錄及轉譯)<br>Chap 1: Introduction to Bioinformatics (生物資訊學簡介)<br>Chap 2: Access to Sequence Data and Literature Information (基因序列資料和文獻資訊存取)<br>Chap 3: Pairwise Sequence Alignment (雙序列比對)<br>Chap 4: Basic Local Alignment Search Tool (BLAST序列比對工具)<br>Chap 5: Basic Local Alignment Search Tool (BLAST序列比對工具)<br>Chap 6: Bioinformatic Approaches to Gene Expression (利用生物資訊探討基因表現)<br>Chap 7: Gene Expression: Microarray Data Analysis (基因表現：微晶片陣列資料分析)<br>Chap 8: Protein Analysis and Proteomics (蛋白質分析及蛋白質體學)<br>Chap 9: Protein Structure (蛋白質結構)<br>Chap 10: Multiple Sequence Alignment (多序列比對)<br>Chap 11: Molecular Phylogeny and Evolution (分子種系和演化)<br>Chap 12: Completed Genomes and the Tree of Life (已完成之生物基因體和生命樹) | Syllabus           | Chap 0: Cell Function, Protein and DNA, Transcription and Translation<br>Chap 1: Introduction to Bioinformatics<br>Chap 2: Access to Sequence Data and Literature Information<br>Chap 3: Pairwise Sequence Alignment<br>Chap 4: Basic Local Alignment Search Tool<br>Chap 5: Basic Local Alignment Search Tool<br>Chap 6: Bioinformatic Approaches to Gene Expression<br>Chap 7: Gene Expression: Microarray Data Analysis<br>Chap 8: Protein Analysis and Proteomics<br>Chap 9: Protein Structure<br>Chap 10: Multiple Sequence Alignment<br>Chap 11: Molecular Phylogeny and Evolution<br>Chap 12: Completed Genomes and the Tree of Life |