



六、教學進度(Syllabi)

2008/11/12	Protein Structure Prediction and Anslsysis	張培均
2008/11/19	期中考	張培均
2008/11/26	Intermolecular Interactions and Biological Pathways	張培均
2008/12/3	Assessing Pairwise Sequence Similarity: BLAST and FASTA	張培均
2008/12/10	Creation and Analysis of Protein Multiple Sequence Alignment	張培均
2008/12/17	Sequence Assembly and Finishing Methods	張培均
2008/12/24	Phylogenetic Analysis	張培均
2008/12/31	Computational Approaches in Comparative Genomics	張培均
2009/1/7	Proteomics and Protein Identification	張培均
2009/1/14	期末考	張培均

七、評量方式(Evaluation)

期中報告：50% 期末報告：50%

八、講義位址(<http://>)

九、教育目標

重新查詢