

Tentative Course Schedule

Physics 323/623: Computational Biophysics Lab

Time: Tuesday, 3:00-4:50 pm

Date	Topic	Reading Assignment (Read prior to coming to lab; it is expected that students will have completed lecture readings, as well)	Laboratory Assignments Due (due by the beginning of lab on due date)	Leader
Tues Aug 29	Lab 1: Physical models: amino acid and protein structure	*amino acid web sites; Chapter 3, section 3.10; *Richardson 1981 review		JSF
Tues Sep 5	Lab 2: Visualization of proteins: VMD and PDB	Chapter 3, section 3.10; *Richardson 1981 review; *VMD and PDB web pages; *Eglin xtal paper	Lab 1 due	JSF
Tues Sep 12	Lab 2/3: Visualization of proteins revisited	*Eglin xtal and NMR papers; Richardson 1994		JSF
Tues Sep 19	Lab 3: Visualization of proteins revisited (cont)	*Eglin xtal and NMR papers; Richardson 1994	Lab 2 due	JSF
Tues Sep 26	Lab 4: Introduction to MatLab and molecular forces	MatLab book (read Chapter 1 and bring to class); Jackson Chapter 2 (2.1- 2.14)	Lab 3 due	JSF
Tues Oct 3	Lab 4: Molecular forces (MatLab) (cont)	MatLab book (read Chapter 1 and bring to class); Jackson Chapter 2 (2.1- 2.14)		JSF
Tues Oct 10	(cancelled—work on lecture take-home midterm)			
Tues Oct 17	Lab 5: Helix-coil transition; distances; dihedrals (MatLab)	Chapter 3 (all) (Bring MatLab book to class)	Lab 4 due	FRS
Tues Oct 24	How to read a scientific paper	*Eglin research paper	Lab 5 due	JSF/FRS
Tues Oct 31	Lab 6: Introduction to DEAC cluster, Linux operating system; editing and queuing	*DEAC cluster web pages	~2 page paper: eglin c background, including sequence, structure, function, molecular dynamics	FRS
Tues Nov 7	Lab 7: Dynamics in explicit solvent: setting up the eglin simulations	*MD introduction paper; *NAMD web pages	~2 page paper: MD review of work on eglin c and CI2	FRS
Tues Nov 14	Lab 7 (cont): NAMD trajectory analysis (MatLab, CHARMM and VMD)		Methods paper: dynamics in explicit solvent	FRS/JSF

Tues Nov 21	Student presentations		Putting it together, first draft: (Introduction, Methods, Results, Discussion, thus far)	students
Tues Nov 28	Student presentations			students
Tues Dec 5			Final paper due (including free energy calculations); discussion to include comparison to other student results and to literature)	

*Reading assignments for labs:

- Amino acid web sites: <http://wbiomed.curtin.edu.au/teach/biochem/tutorials/AAs/AA.html>; <http://www.rothamsted.ac.uk/notebook/courses/guide/aa.htm>
- Richardson review, 1981: Anatomy and Taxonomy of Protein Structure, J. S. Richardson (1981, as pp. 167-339 in volume 34 of Advances in Protein Chemistry published by Academic Press); electronic version: <http://kinemage.biochem.duke.edu/~jsr/index.html>
- Richardson 1994: Richardson JS. Introduction: protein motifs. FASEB J. 1994 Dec;8(15):1237-9.
- VMD and PDB web pages: <http://www.ks.uiuc.edu/Research/vmd>; www.rcsb.org;
- Eglin xtal and NMR papers:
 - Bode, W., Papamokos, E., Musil, D. The high-resolution X-ray crystal structure of the complex formed between subtilisin Carlsberg and eglin c, an elastase inhibitor from the leech *Hirudo medicinalis*. Structural analysis, subtilisin structure and interface geometry. Eur.J.Biochem. v166 pp.673-692, 1987.
 - Hyberts, S.G., Goldberg, M.S., Havel, T.F., Wagner, G. The solution structure of eglin c based on measurements of many NOEs and coupling constants and its comparison with X-ray structures. Protein Sci. v1 pp.736-751, 1992.
- Eglin research paper: Reich L, Weikl TR. Substructural cooperativity and parallel versus sequential events during protein unfolding. Proteins. 2006 Jun 1;63(4):1052-8.
- DEAC cluster web pages: http://www.deac.wfu.edu/index.php?title=WFU_DEAC_Cluster
- NAMD web pages: <http://www.ks.uiuc.edu/Research/namd/>
- MD introduction paper: Hansson T, Oostenbrink C, van Gunsteren WF. Molecular dynamics simulations. Current Opinion In Structural Biology 12 (2): 190-196 APR 2002
- Free energy calculation paper: Kollman PA, Massova I, Reyes C, et al. Calculating structures and free energies of complex molecules: Combining molecular mechanics and continuum models Accounts of Chemical Research 33 (12): 889-897 Dec 2000