

Computing for Life Sciences (CS 59000)

Course Content: The accumulation of biological data, such as genome sequences, protein structures and sequences, metabolic pathways, has opened up a new way of research in biology – bioinformatics. In this course we will learn programming, bioinformatics databases, tools, and algorithms behind these tools. Covered topics will include methods for protein sequence comparison, protein structure comparison, protein structure prediction/modeling, protein docking prediction, protein function prediction, and protein network analysis.

This is a core course for CLS specialization of graduate degree.

See <http://www.gradschool.purdue.edu/CLS/index.cfm>

Grading: Grades in the course will be based on midterm & final exam (40%), homework assignments (30%), and projects (30%). The course will be concluded by presentations of group projects. Semester grades will be awarded based on the following minima of performance: 90% = A, 80% = B, 70% = C and 60% = D; < 60% = F. These thresholds will not be raised but may be lowered at the instructor's discretion.

Projects: Considering that this course is cross-listed between different departments, students will choose either of programming-type or mini-review-type (+ using existing software) project by the consent of the instructor.

Instructor: Daisuke Kihara (dkihara@purdue.edu), Robert Skeel (skeel@cs.purdue.edu) Professor Skeel will teach in the first 3 weeks (Aug 22nd – Sep. 9th) on basics of unix commands and Python programming. Professor Kihara will teach for the rest of the semester on bioinformatics methods.

Time & Place: MWF, 11:30am - 12:20pm, LWSN 1106

TextBook: We don't use a specific textbook. But here are good books to refer to:

- Learning Python, 2nd Edition, by Mark Lutz, and David Ascher, O'Reilly, December 2003, ISBN: 0-596-00281-5,
- Biological Sequence Analysis, Durbin, Eddy, Krogh, Mitchison, ISBN 0-521-62971-3
- Protein Bioinformatics, by Ingvar Eidhammer, Inge Jonassen & William R. Taylor, Wiley, ISBN: 0-470-84839-1

Fall 2011: Provisional Schedule

Week	Lec. #	Date	Topic
1	1	Aug. 22	Introduction of the course, Unix file system (Skeel)
	2	Aug. 24	Unix commands
	3	Aug. 26	Computing and scripting in Python
2	4	Aug. 29	Strings and files
	5	Aug. 31	Regular expressions Homework #1 (programming) out
	6	Sep. 2	Control structures and variables
3	-	Sep. 5	(no class, Labor day)
	7	Sep. 7	containers
	8	Sep. 9	Functions Homework #1 (programming) in
4	9	Sep. 12	Numbers and graphics (Skeel)
	10	Sep. 14	Sequence alignment (Kihara)
	11	Sep. 16	Sequence alignment Homework #2 (sequence alignment) out
5	12	Sep. 19	Amino acid similarity matrices
	13	Sep. 21	Multiple sequence alignment
	14	Sep. 23	Sequence motifs Homework #2 turn in
6	15	Sep. 26	Profile motifs (Cont.) Homework #3 (motif) out
	16	Sep. 28	Database search, Statistical Analysis
	17	Sep. 30	Database search, Statistical Analysis (cont)
7	18	Oct. 3	several databases tools for motif, domain search Homework #3 in
	19	Oct. 9	Hidden Markov models
	20	Oct. 7	HMM (cont.)
8	-	Oct. 10	(no class, October break)
	21	Oct. 12	Protein Structure Homework #4 (HMM) out
	22	Oct. 14	Protein Structure (cont.)
9	23	Oct. 17	Protein structure comparison
	24	Oct. 19	neural network: Protein secondary structure prediction

	25	Oct. 21	Protein Secondary Str. prediction cont.
10	26	Oct. 24	Template based structure modeling Homework #4 in
	27	Oct. 26	Template based str. modeling (cont.)
	28	Oct. 28	Geometrical properties of protein structures Homework #5 (protein structure) out
11	29	Oct. 31	Mid term exam
	30	Nov. 2	Ab initio protein prediction
	31	Nov. 4	Protein-protein interaction network
12	32	Nov. 7	Protein-protein complex: Geometric hashing
	33	Nov. 9	Side-chain optimization Homework # 5 in
	34	Nov. 11	Function prediction from structure
13	35	Nov. 14	Project out
	36	Nov. 16	Comparison of chemical molecules (drugs)
	37	Nov. 18	Ligand (drug) protein interaction
14	38	Nov. 21	Comparative genomics
	-	Nov. 23	(no class, university holiday)
	-	Nov. 25	(no class, university holiday)
15	39	Nov. 28	Systems biology (gene expression, pathways etc.)
	40	Dec. 30	Systems biology
	41	Dec. 2	Project presentation
16	42	Dec. 5	Project presentation
	43	Dec. 7	Advanced research Topics
	44	Dec. 9	(Course evaluation)
13		Dec. 12	Final Exam Week
		Dec. 14	
		Dec. 16	