CSE 427 Introduction to Computational Biology: Generic Syllabus Department of Computer Science & Engineering

This generic syllabus describes CSE427 as approved and initially taught in 2006-7. Subsequent offerings will differ in detail, while retaining the same focus on the mathematical and computational principles underlying this interdisciplinary field.

Topics:

- 1. Basics of Molecular Biology
- 2. Pairwise sequence alignment
- 3. The algorithms and statistics of BLAST
- 4. Amino acid substitution matrices
- 5. Multiple sequence alignment
 - Definition, NP-completeness, progressive alignment, whole-genome multiple alignment
 - Comparative sequence analysis and the UCSC Human Genome Browser
- 6. Inference of phylogenetic trees
- 7. Markov chains and hidden Markov models
- 8. Challenges in Computational Molecular Biology

Reading list:

- 1. <u>Basics of Molecular Biology</u> (class handout)
- 2. Molecular Biology for Computer Scientists, a primer by Lawrence Hunter
- 3. Pairwise sequence alignment: Lectures 3-5 from <u>CSE 527</u>
- 4. Altschul, Gish, Miller, Myers & Lipman. <u>"Basic local alignment search tool"</u>. Journal of Molecular Biology 215:403-410 (1990).
- 5. BLAST statistics: <u>Stephen Altschul's primer</u>.
- 6. Multiple Sequence Alignment: Sections 6.1-6.4 from <u>CSE 527</u>
- Blanchette, Kent, Riemer, Elnitski, Smit, Roskin, Baertsch, Rosenbloom, Clawson, Green, Haussler, and Miller "Aligning Multiple Genomic Sequences With the Threaded Blockset Aligner". Genome Res. 14: 708 – 715 (2004).
- 8. Section on phylogenetic trees in Lecture 4 from Michael Brudno's lecture notes

Programming projects (for more details, see:

http://www.cs.washington.edu/education/courses/cse490c/07wi/homework.html)

- Week 1: Prokaryotic gene prediction
- Week 3: Protein similarity search via dynamic programming
- Week 5: BLAST similarity search versus a sequence database
- Week 7: Sequence conservation via multiple sequence alignment
- Week 9: Hidden Markov models

Grading criteria: 18% per project plus 10% based on participation in class discussion.