Algorithms for Computational Biology

Zsuzsanna Lipták

Masters in Molecular and Medical Biotechnology a.a. 2015/16, fall term

Organisation

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- course times: Thu 10:30 12:30 (aula L), Fri 13:30 16:30 (aula H)
- language: English, but you can ask questions in English or in Italian
- webpage: coming soon!

will include: current info, slides, materials etc.

 attendance: not obligatory but recommended (not all material will be available on webpage, plus: read what I wrote on the webpage of the course)

Organisation (cont.)

- email: zsuzsanna.liptak@univr.it Please put "Algorithms for Computational Biology" in the subject line
- office: CV 2, 1st floor, room 1.79
- student hours: Wed 10:30-12:30 ???

Organisation (cont.)

- this course is 6 CFU
- exam:

written and oral: admitted to oral only if you pass the written test

 for those who want this course to count for the Masters in Medical Bioinformatics (Dip. Inf.): it can replace 6 CFU of the "Fundamental Algorithms in Comp. Biology" only if you do an additional assignment on a topic from algorithms in computational biology (to be chosen together with lecturer)

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Questions?

Goals of this course

- 1. to learn about some basic problems and algorithms behind common bioinformatics applications (sequence alignment, sequence similarity, phylogenetic reconstruction)
- 2. to get an idea of some basic computational issues involved (problem specification, efficiency, limitations)

Overview

• Part I: Sequence Analysis

- Pairwise sequence alignment
- Detour: Algorithm analysis
- Multiple sequence alignment
- String similarity and distance
- Scoring matrices
- Heuristic database search: FASTA, BLAST

• Part II: Phylogenetics

- Detour: Trees and graphs
- algorithms for distance-based data
- character-based data, Perfect Phylogeny
- Small Parsimony: Fitch's algorithm
- Large Parsimony: heuristics

Books

- Neil C. Jones and Pavel A. Pevzner: An Introduction to Bioinformatics Algorithms (2004).—3 copies in library
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- João Setubal, João Meidanis: Introduction to Computational Molecular Biology (1997).—my favourite, 1 copy in library
- **Dan Gusfield**: Algorithms on Strings, Trees, and Sequences (1997).—the bible of string algorithms, a bit dated now
- Joseph Felsenstein: Inferring Phylogenies (2004).—important book on phylogenetics, very understandably written
- Cormen, Leiserson, Rivest (& Stein): Introduction to Algorithms (different editions, 1990-onwards).—the bible of algorithms, a must have for anyone interested in algorithms (buy second hand, old editions are also fine)