Bioinformatics



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Students' assistance:

Wednesday, from 10.00 to 12.00, room 120

By appointment, agreed in advance via email

Course program

- Main topics
 - Molecular biology and biochemistry
 - Biological database searching and pairwise alignments
 - Substitution patterns
 - Distance-based phylogenetic methods
 - Character–based phylogenetic methods
 - Genomics and gene recognition
 - Prediction of protein and RNA structures
 - Proteomics
 - Machine learning tools for bioinformatics
- Seminars from academic and industry experts

Textbooks

- Dan E. Krane, Michael L. Raymer, Fundamental Concepts of Bioinformatics, Benjamin–Cummings Pub. Co.
- → Arthur M. Lesk, An Introduction to Bioinformatics, Oxford Press, 4th ed.
- Slides, freely downloadable at www.diism.unisi.it/~monica/bioinfo.html

Exam

- Practical exam choose one of the two options:
 - Use of machine learning tools for biological data processing (based on existing software)
 - Team work
 - Discussion of the obtained results during the oral exam
 - Preparation of a small presentation based on the reading of some scientific papers
- Oral exam on the whole contents of the lectures

Collaborations

- Department of Biotechnology, Chemistry and Pharmacy, University of Siena
- Department of Medical Biotechnologies, University of Siena
- Department of Medicine, Surgery and Neuroscience, University of Siena
- Toscana Life Sciences
- Experimental and Clinical Medicine, University of Florence
- Microbiology and Virology Laboratory, Careggi University Hospital of Florence
- Department of Chemistry and Applied Biosciences, ETH Zürich
- Radboud University Medical Center, Nijmegen, The Netherlands
- GSK Vaccines (Siena)
- Philogen/Philochem (Siena/Zürich)
- bioMérieux (Florence)