

Code: 1328 Computational Biology**Degree:** 2nd cycle - Functional Biology**Curricular Year:** 1st**Credits:** 6 ECTS**Semester Course:** 2nd**Compulsory****Language:** Portuguese/English**Responsible:** Marta Guerreiro Duarte Mesquita de Oliveira**Other lecture(s):** Maria Leonor Mota Morais Cecílio and Maria Manuela Costa Neves Figueiredo**Web Site:** <http://www.isa.utl.pt/home/node/3769>**1. Contact hours:****Lecture/Practicals 70 Others 14 Total 84****2. Objectives:**

The aim is to link biology and computer science knowledge in order to analyze and model biological data.

3. Programme:

Algorithms design
Evolutionary algorithms
Graphs
DNA sequencing: Fragments assembly
Sequence alignment
The Motif finding problem
Probabilistic Models
Markov Chains
Statistical Inference
Microarrays
Clustering and Biclustering
The Bioconductor software R

4. Bibliography:**Main Bibliography**

- An Introduction to Bioinformatics Algorithms, N. C. Jones and P. Pevzner, MIT Press, 2004.
- Computational Genome Analysis. An Introduction, Richard Deonier, S Tavaré, and Michael S. Waterman, Springer Verlag, 2005.
- Statistical Methods in Bioinformatics. An introduction (2001). W. Ewens and G. Grant. Statistics for Biology and Health
- Evolutionary Computation in Bioinformatics, Fogel, G.; Corne, D.W. (Eds), Morgan Kaufmann Publishers, Elsevier Science, 2003.

Other Bibliography

- Bioinformatics and Computational Biology Solutions using R and Bioconductor, Gentleman, R.; Carey, V.; Huber, W.; Irizarry, R.; Dudoit, S. (Eds), Springer 2005.

5. Assessment:

- 2 midterm exams and 3 homeworks
- Or
- 1 final exam.

6. Estimated Workload:

168	Hours
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7. Last Update:

14/7/2010
