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Bioinformatics analysis of biological sequences: from sequence to structure

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WHAT YOU WILL LEARN

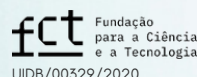
Acquire knowledge regarding bioinformatic tools available to predict nucleic acid and protein three-dimensional structure, as well as autonomy and critical thinking in the use of those tools. Develop skills on the use of bioinformatics software freely available on the Internet and to interpret the biological meaning of the results. At the end of this course, researchers, molecular biology students, or health professionals will be able to predict the three-dimensional structure of nucleic acid and proteins from their sequence and use the acquired knowledge to improve or create new methodologies in molecular biology research and diagnostics.

FEE

1st year PhDs at FCUL	Free
BIODIV PhD students	Free
1st year PhDs BEAG & SC	Free
Other PhDs at cE3c	30,00€
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BIODIV – Biodiversity, Genetics and Evolution (ULisboa & UPorto)
BEAG – Biology and Ecology of Global Changes (ULisboa & UAveiro)
SC – Sustainability Sciences (ULisboa)
CFE – Centre for Functional Ecology

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