



# Short Course in Bioinformatics

17<sup>th</sup> - 19<sup>th</sup> March 2025

## AGENDA

### March 17<sup>th</sup> 2025

Time	Duration	Topic	Speaker
10h00	15'	Welcoming session	Pedro Leão
10h15	45'	Introduction to Bioinformatics in Natural Products Discovery	Adriana Rego
11h00	30'	NLP/AI for Functional Annotation of Enzymes: A Case Study with FAALPred	Leandro Pereira
11h30	30'	<b>Coffee-break</b>	
12h00	30'	Introduction to antiSMASH and BiG-SCAPE workflows	Catarina Loureiro
12h30		<b>Lunch break</b>	
14h00	60'+30'	Hands- on session antiSMASH and BiG-SCAPE	Catarina Loureiro
15h30	30'	<b>Coffee-break</b>	
16h00	30' + 60	Introduction to GNPs and MicrobeMASST Hands-on session - MicrobeMASST	Mauricio Caraballo

# March 18<sup>th</sup> 2025

Time	Duration	Topic	Speaker
9h00	60'+30'	Phylogenetic approaches to natural product discovery  (short) Hands-on session NaPDos2	Hans Singh
10h30	30'	<b>Coffee-break</b>	
11h00	30'+45'	Genomic context of target biosynthetic genes  (short) Hands-on session genomic context	Adriana Rego  Adriana Rego and Catarina Loureiro
12h15			
<b>Lunch break</b>			
13h45	45+60'	Introduction to computational protein structure prediction  Hands-on session to computational protein structure prediction, molecular docking	Arménio Barbosa
15h30		<b>Coffee-break</b>	
15h45	30'	EFI-SSN and Sequence-Based Functional Prediction in Enzyme Discovery	Raquel Castelo Branco



# March 19<sup>th</sup> 2025

Time	Duration	Topic	Speaker
10h	15'	Welcoming session - Presentation of the speakers	
10h15	30'	<i>Open-science-driven development of genome mining tools and their application on microbiomes</i>	Catarina Loureiro
10h45	15'	Q&A	
11h00	30'	<b>Coffee-break</b>	
11h30	45'	<i>Advancing Biotechnology with Molecular Modeling: Applications in Drug Discovery, Protein Purification and Self-Assembly</i>	Arménio Barbosa
12h15	15'	Q&A	
12h30		<b>Lunch break</b>	
14h30	45'	<i>Targeted gene cluster mining with GATOR-GC</i>	Marc G. Chevrette
15H15	15'	Q&A	
15h30	30'	<b>Coffee-break</b>	
16h00	45'	<i>A journey for the discovery of microbial molecules, the need of reference datasets, and using small molecules to understand symbiotic relationships in ants' ecosystems</i>	Mauricio Caraballo
16H45	15'	Q&A	
17h00	10'	Final remarks	

