

Program

Wednesday, December 21st 2022

8:30 Registration

9:30 Opening Session

Pedro Patacho, Councilor for Education and Science, Municipality of Oeiras

Cláudio M. Soares, Dean of the ITQB-NOVA

Manuel N. Melo, 3D-BioInfo-PT-Community

9:45 Carlos S. H. Shiraishi

“Virtual screening of a library of natural compounds against COX-2 protein”

10:00 Gabriel F. Martins

“Polyphenols as Aggregation Inhibitors: Application to α -Synuclein and Related Peptides”

10:15 Israa Aljnadi

“In Silico Studies and Chemical Synthesis of Ligands Targeting c-MYC G-Quadruplex”

10:30 Marta S. P. Batista

“Structural characterization of the permeation of water and glycerol derivatives through PfAQP for the development of new antimalarial therapies”

10:45 Rita I. Teixeira

“The impact of SARS-CoV-2 Omicron variant on the interaction with human ACE2”

11:00 Coffee Break and Poster Session I

11:45 Wallfuture

11:55 Keynote Lecturer – Carla Sousa

“MD simulations applied to Infection research: improving studies of drug permeation across biomembranes”

12:55 Tomás F. Silva

“Tackling the realism of transmembrane peptides simulations with a pH-Gradient/CpHMD approach”

13:05 Pedro C. Rosado

"Exploring PBP2a protein to fight β -lactam resistance in MRSA"

13:25 Rodrigo Barriga

"Simulating substrate binding sites in the *S. aureus* Type II NADH Dehydrogenase"

13:40 Tatiana F. Vieira

"What can computational methods do to help find new molecules with antimicrobial properties"

13:45 Lunch

15:50 Armindo Salvador

Presentation of the Portuguese Biophysical Society (SPBf)

SPBf Session

16:00 Pedro Beltrão

"Studying protein interactions and domain evolution using AlphaFold"

16:45 Coffee Break and Poster Session II

17:30 Bruno Calçada

"Development of machine learning models to predict thyroid peroxidase inhibition"

17:45 João G. N. Sequeira

"Extending the stochastic titration CpHMD to the CHARMM36m and AMBER14SB force fields"

18:00 Urszula Orzeł

"Oligomerization of metabotropic glutamate receptors"

18:15 Andreia Fortuna

"Tackling halogen anisotropy in biomolecular simulations: assessment of force field parameters using hydration free energies"

18:30 Bárbara Bahls Bruni

"Design and virtual screening of Indoloisoquinoline derivatives as c-MYC G4 binders"

18:45 Closing Session/Remark