

Workshop at Partner University: Phylogeny & Evolution

Where: University of Mainz/Germany, partner of University of Zagreb

Situated at the River Rhine, Mainz has a rich history and currently about 220.000 citizens; it is close to Frankfurt/Main airport.

Who: Scheduled for six participants (B.Sc., M.Sc. & PhD students; scientists; Biology or related)

When: 15. to 19. April 2024 (20 hours in total)

<u>Criteria for student participants</u>: Depending on the number of applicants study achievements will be considerred.

Financial support by DAAD: 234 € for B.Sc. and M.Sc. students, 324 € for PhD students, 434 € for scientists

<u>Training</u>: Computational analyses of phylogeny and sequence evolution on proteins and coding sequences

Examples will be on mammals, especially primates, but the methods can be applied to any group of organisms. We will additionally carry out phylogenetic exercises based on anatomical data collected from skull models.

Language: English – but don't worry: Everyone will be able to follow.

Benefit: Certificate of participation and ECTSs for students

...besides training in computational biology, meeting local participants and getting to know Mainz

Info meeting: around Christmas '23 or New Year '24; location to be determined

Applicants contact: Prof. Dr. Maria Spoljar (maria.spoljar@biol.pmf.hr)

or Prof. Dr. Holger Herlyn (herlyn@uni-mainz.de); application deadline 20th of Nov. '23

Yours,









Computational

bioloav

Ssc 2 0.6830

- Bta 3 0.7601

- Cia 5 0.2665

- Ppy 7 1.7502

