

— EuBIC 2020 Developer's Meeting – Save the date & call for project proposals —

We are pleased to announce that the upcoming EuBIC 2020 developer's meeting will be held in Nyborg, Denmark, on **January 13th to 17th 2020**.

The EuBIC 2020 developer's meeting will bring together scientists active in the field of bioinformatics and computational proteomics. It provides a highly interactive platform to develop projects that are relevant to the proteomics community.

Date: January 13-17, 2020

Venue: Sinatur Hotel Storebælt, Østerøvej 121, 5800 Nyborg, Denmark

Website: <https://www.proteomics-academy.org/eubic-dev-meeting-2020>

Key dates

September 18, 2019: Project proposal submission deadline

October 1, 2019: Announcement of selected projects

October 1, 2019: Conference registration opens

January 13-17, 2020: EuBIC 2020 developer's meeting in Nyborg

Call for project proposals

Do you want other researchers working on your project and set up future collaborations?

The call for topics for the hackathon sessions during the EuBIC 2020 developer's meeting is now open!

During the EuBIC 2020 developer's meeting the participants will engage in hackathon sessions where they will develop bioinformatics tools and resources. The participants will break up into small groups to work together on these topics in an open, constructive, and productive atmosphere.

You can now submit project proposals for inclusion into the EuBIC 2020 developer's meeting program. Submit your project proposal before 18 September through the following website: <https://github.com/eubic/EuBIC2020>

5-8 hackathon projects will be selected by a community vote arranged by the EuBIC program committee and announced on the conference webpage.

Topics

Hackathon projects can cover all topics in computational mass spectrometry and proteomics bioinformatics, including, but not limited to:

- Tools
 - Extension of existing bioinformatics tools under the guidance of their original authors
 - Development of novel bioinformatics tools to solve open problems
- Benchmarking
 - Compilation of gold-truth datasets
 - Collection of reference algorithmic implementations

- Specification of benchmarking criteria
- Workflows
 - Definition of formal protocols
 - Development of tutorials for bioinformatics software usage
- Data analysis
 - (Re)analysis of publicly available proteomics datasets
 - Functional data analysis and integration

More information

- Website: <https://github.com/eubic/EuBIC2020>
- Social media hashtag: #EuBIC20
- Conference email: eubic2020@rub.de

Sponsorship inquiries are welcome at eubic2020@rub.de

We are grateful to the European Proteomics Association (EuPA) for supporting EuBIC and to the Novo Nordisk Foundation for their generous support of the conference!

We look forward to welcoming you in Nyborg!

The organizing committee.

Veit Schwämmle
Marie Locard-Paulet
Wout Bittremieux
David Bouyssié
Viktoria Dorfer
Dominik Kopczynski
Alessio Soggiu
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