



13 - 17 January 2020 in Nyborg (DK)

Kick-off with introductory and **stimulating keynotes**

Half day **educational session**

Followed by hands-on **hackathon** sessions based on the **community interest**

## Hackathon projects

Batch XIC and spectra extraction in ThermoRawFileParser

Vladimir Gorshkov, Niels Hulstaert, Yasset Perez-Riverol

How can we best use Cytoscape for proteomics data analysis?

Nadezhda T. Doncheva, Marc Legeay

Online spectrum identification validation by comparison to predicted/experimental spectra

Mathias Wilhelm, Tobias Schmidt, Patroklos Smaras

Formation of spectral libraries by representative spectra

Lukas Käll

Mapping proteins to functions: method and benchmark development

Bart Mesuere, Pieter Verschaffelt, Henning Schiebenhoefer

Simulating a quantified phosphoproteome for software benchmarking and algorithm development

Marie Locard-Paulet, Veit Schwämmle, Vasileios Tsiamis

## Keynote speakers

Eric Deutsch

Lydie Lane

Ole N. Jensen

Alexander Peltzer

Andy Jones

Olga Vitek

## Registration deadlines:

Early bird: November 28<sup>th</sup>, 2019

General: December 23<sup>th</sup>, 2019

## And more

Intro to computational proteomics

Poster session

Demos of software and tools

Social event

...

## More info and registration:

[eubic-ms.org/2020-dev-meeting](http://eubic-ms.org/2020-dev-meeting)

## Get social!

@EuBIC\_ms #EuBIC2020

