Computational Mass Spectrometry-Based Proteomics

Quantitative proteomics applied to life sciences and systems biology

23-28 June 2013
Munich, Germany

The Summer School will provide hands-on training in computational analysis of proteomics data generated by high-resolution mass spectrometry. This powerful technology is increasingly applied to systems-wide investigation of different aspects of cell biology. The practical part will enable participants to perform sophisticated quantitative data analysis in their own future research using the publicly available MaxQuant software (Cox and Mann, Nature Biotechnology, Dec 2008).

Organizers
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Registration deadline is 15th February 2013

Further details will be available in December at www.biochem.mpg.de/summerschool
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