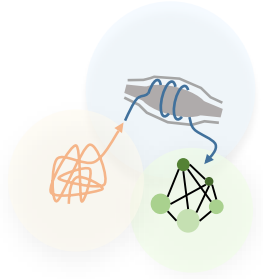

Quantitative Proteomics and Bioinformatics for life scientists

Nov 23 - 25, 2016

CECAD, Joseph-Stelzmann-Str. 26, Vortragsaal EG (Raum 0.36)

23rd

11:30 - 12:30	**Pitter Hüsgen** (Jülich)	Terminomics - System-wide protein termini identification reveals proteolytic protein modification
		- Lunch -
13:30 - 14:30	**Rene Zahedi** (Dortmund)	The power of chromatography for pre-fractionation and mass spectrometry
14:30 - 15:30	**Boris Macek** (Tübingen)	Cancer Proteogenomics
15:30 - 16:30	**Felix Meissner** (Martinsried)	Quantitative Proteomics to Study Immune Cell Communication

24th

10:45 - 11:45	**Jesper Olsen** (Copenhagen, DK)	Phosphoproteomics & Signaling
11:45 - 12:45	**Blagoey Blagoev** (Oddense, DK)	Studying Protein Ubiquitination by Mass Spectrometry
		- Lunch -
13:45 - 14:45	**Jörn Dengjel** (Fribourg, CH)	Degradation of protein translation machinery by amino acid starvation-induced macroautophagy
14:45 - 15:45	**Michael Lund Nielsen** (Copenhagen, DK)	Proteome-wide analysis of arginine mono-methylation reveals widespread occurence in human cells
		- Coffee Break -
16:00 - 17:00	**Sriram Aravamudhan** (Leiden, NL)	PTMScan- antibody enrichment of PTM peptides for MS-based proteomics
17:00 - 18:00	**Jeroen Krijgsveld** (Heidelberg)	Genome meets proteome: a novel strategy to investigate chromatin-bound protein networks
		- Get-together (Bistro Olivieri) -

25th

10:15 - 12:00	**Jürgen Cox** (Martinsried)	MaxQuant & Perseus

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