

Proteomics Methods Forum

Day	Time	Opening Session	Chair: Gary Black
Thursday	13:00	Achim Treumann	The results of the PMF2011 survey
June 16 th	13:15	Duncan Smith	Maximising qualitative penetrance and quantitative quality in label free proteomics
	13:45	LUNCH	
		Statistics	Chair: Sarah Hart
	14:30	Laurent Gatto and Kathryn Lilley	Statistics and experimental design
	15:00	Howsun Jow	A Bayesian approach to statistical analysis of isobaric relative MS quantitation
	15:15	David A. Cairns	Learning by looking at clinical proteomics data; simple statistical analysis can be enlightening
	15:45	Marek Gierlinski	Time-resolved proteomics
	16:00	COFFEE	
		PTM	Chair: Duncan Smith
	17:00	Jerry Thomas	Global phosphoproteomics of synaptosomes
	17:15	Mark Skehel	Phosphopeptide analysis
	17:45	Karen Lowdon	Methylation of the Androgen Receptor
	18:00	Ian Fearnley	Proteomic investigation of methylation in mitochondria
	18:30	DRINKS RECEPTION	
Friday		Data, Software, Metabolomics	Chair: Nick Morrice
June 17 th	09:00	Karl Burgess	Metabolomics and proteomics, a bitter perspective on both
	09:30	Pedro Cutillas	LCMS based metabolomics
	09:45	David Knight	Can the analysis of GCMS metabolomics data inspire proteomics?
	10:15	Roman Fischer	How to spend 6 hours with 5 ug of cell lysate and an Orbitrap...
	10:30	COFFEE	
		Quantitation	Chair: Kathryn Lilley
	11:00	Karin Barnouin	Relative Label-Free Quantitation of Post-Translational Modifications
	11:15	Saw Yen Ow	Squeezing that extra bit more from your iTRAQ ratios: Less underestimations to celebrate our 7th anniversary
	11:45	Nick Morrice	SILAC analysis on 5600 Triple Quad and Velos Orbitrap
	12:15	Amy Claydon	Absolute quantification using QconCAT
	12:45	CLOSE MEETING	
	13:00	LUNCH	

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