

Swiss Proteomics Society

2006 Interactive Training Initiative



Applied Proteomics – Tips & Tricks

on-site applied training course with international debriefing workshop

Debriefing workshop – December 4 2006

Program

9 ⁰⁰ -9 ¹⁵	Welcome introduction	
9 ¹⁵ -10 ⁴⁵	Session: Separation techniques	Chairman: Jean-Charles Sanchez
9 ¹⁵ -9 ³⁵	Hans Voshol et al. – Subcellular proteomics	
9 ³⁵ -9 ⁵⁵	Catherine Zimmermann et al. – Shotgun Isoelectric Focusing peptides analysis	
9 ⁵⁵ -10 ¹⁵	Olivier Barré et al. – Two-dimensional liquid separation with the Proteomelab PF2D	
10 ¹⁵ -10 ⁴⁵	Open discussion	
10 ⁴⁵ -11 ¹⁵	Coffee break	
11 ¹⁵ -13 ¹⁵	Session: Mass spectrometry	Chairman: Reto Stöcklin
11 ¹⁵ -11 ³⁵	Michael Afolter et al. – Multidimensional LC-MS/MS analysis and gas phase fractionation	
11 ³⁵ -11 ⁵⁵	Philippe Favreau et al. – De novo ms/ms sequencing	
11 ⁵⁵ -12 ¹⁵	Manfredo Quadroni et al. – Analysing protein phosphorylation sites by mass spectrometry	
12 ¹⁵ -12 ³⁵	Bernd Bodenmiller et al. – Selective isolation and quantitative ms of phosphopeptides	
12 ³⁵ -13 ¹⁵	Open discussion	
13 ¹⁵ -14 ⁴⁵	Lunch	
14 ⁴⁵ -16 ¹⁵	Session: Bioinformatics	Chairman: Hans Voshol
14 ⁴⁵ -15 ⁰⁵	Marie-Claude Blatter et al. – The protein sequence databases and the UniProtKB/Swiss-Prot	
15 ⁰⁵ -15 ²⁵	Christine Hoogland et al. – 2-DE analysis and publishing	
15 ²⁵ -15 ⁴⁵	Patricia Palagi et al. – Mass spectra analysis	
15 ⁴⁵ -16 ¹⁵	Open discussion	
16 ¹⁵ -17 ⁰⁰	Closing remarks and coffee	