

Monday 29th January

Time	Speaker	Topic
10:30 - 10:40	Brendan MacLean	Opening Remarks
10:40 - 11:40	Sandi Spencer	Lecture: Moving from discovery to targeted Proteomics
11:40 - 12:40	Brendan MacLean	Lecture: Introduction to Skyline
12:40 - 1:30	Lunch	
1:30 - 2:45	Brendan MacLean	Hands-on: Targeted method editing in Skyline
2:45 - 4:00	Brendan MacLean	Hands-on: Targeted method refinement in Skyline
4:00 - 4:30	Break	
4:30 - 5:30	Sandi Spencer	Lecture: Building prior knowledge for SRM assays

Tuesday 30th January

Time	Speaker	Topic
8:30 - 9:45	Brendan MacLean	Hands-on: Data analysis in Skyline
9:45 - 10:45	Sandi Spencer	Lecture: Protein quantification
10:45 - 11:15	Break	
11:15 - 12:15	Brendan MacLean	Hands-on: Calibrated quantification with Skyline
12:15 - 1:15	Brendan MacLean	Hands-on: Group comparison statistics and reports in Skyline
1:15 - 2:00	Lunch	
2:00 - 2:30	Brendan MacLean	Lecture: Overview and keys to success in processing DDA data with Skyline
2:30 - 4:00	Brendan MacLean	Hands-on: Label-free quantitative analysis of DDA data with Skyline
4:00 - 4:30	Break	
4:30 - 6:00	Sandi Spencer	Hands-on: Parallel reaction monitoring (PRM) with Skyline

Wednesday 31st January

Time	Speaker	Topic
9:00 - 10:00	Brian Searle	Lecture: Data independent acquisition
10:00 - 10:30		Break
10:30 - 12:30		Tutorial: DIA data analysis with Skyline (ETH tutorial)
12:30 - 1:30		Lunch
1:30 - 3:30	Brian Searle	Hands-on: Data independent acquisition data analysis
3:30 - 4:00		Break
4:00 - 5:00	Ruedi Aebersold	Keynote: Targeting proteomics at high throughput