

## **Molecular Discovery North America User's Meeting**

Processing Waters Datasets with MassMetasite: A Post-ASMS –  
Russell Mortishire-Smith, Waters Corporation

Met ID for Peptides – Mark Cancilla, Merck

Leveraging SSID in the Lead Optimization Space: A PI's  
Perspective – Kerry Fillgrove, Merck

Cross Platform Comparison for Soft Spot ID – Kevin Bateman,  
Merck and Ismael Zamora, Molecular Discovery, Ltd.

Towards Efficient Metabolite Identification Workflows – Suma  
Ramagiri, AB SCIEX

GSH Workflow by MassMetaSite/WebMetabase – Ismael Zamora,  
Molecular Discovery, Ltd.