

Please mark the training sessions that you would like to do.

Name: _____

- MassMetaSite 3.4.....
 - Simple example to understand the MassMetaSite settings
 - Batch and auto process
 - UNIFI-MassMetaSite connection

- Molecular Discovery Portal.....
 - WebMetabase 4.0:
 - Small Molecule:
 - Protocol definition
 - Experiment and sample list generation
 - Upload a multiple condition experiment
 - Macros
 - Approval process
 - Analysis tools
 - Single molecule Analysis tools:
 - Metabolic pathway and kinetic analysis
 - 32D docking
 - Metadesign
 - Reporting
 - Multiple molecule analysis tools
 - Search
 - Cluster view
 - Comparator
 - SMRT
 - Fragment analysis
 - MacroMolecule
 - Protocol definition
 - Experiment and sample list generation
 - Upload a multiple condition experiment
 - Macros
 - Approval process
 - Helm notation
 - Frequency analysis
 - Site of Cleave predictions

- WebChembase:
 - Chemical degradation experiments
 - Protocol definition
 - Experiment and sample list generation
 - Upload and sample list generation
 - Macros
 - Approval process
 - NMR data for structure elucidation

- Compound library:
 - Data collection
 - Data visualization

- MetaSite 6.0
 - Prediction for:
 - CYP
 - AOX
 - FMO
 - Design tools:
 - 32D
 - MetaDesign

- MassChemSite 3.0.....
 - Chemistry workflows
 - Late stage functionalization using Mass Spec and NMR data
 - Compound degradation studies
 - Reaction Tracking

Thank you very much for your help in organizing hopefully this unique event.