



MS/MS Proteomics Validation at a Glance

Scaffold 3 introduces several new features:

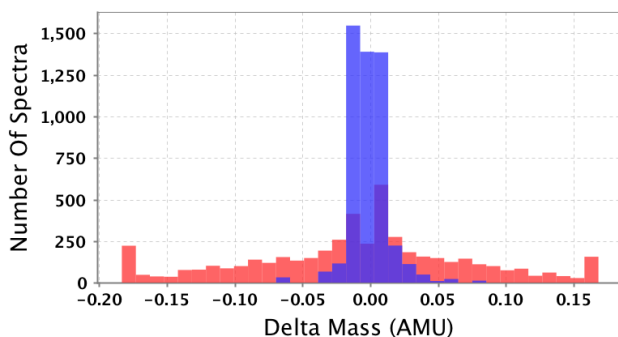
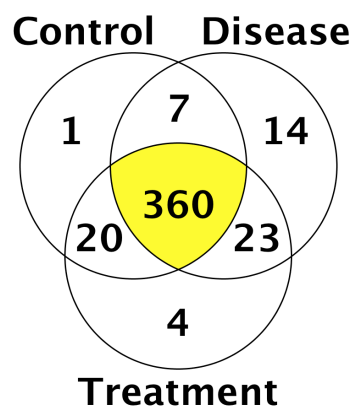


SF3 Format Improves Speed and Stability

Scaffold's new SF3 file format means faster data access and unlimited file size scalability. You'll load more data in less time and always be ready for the next generation of instrumentation.

Powerful Filtering Capabilities

Filter your data like never before. Use Venn diagrams and advanced filtering capabilities to quickly navigate your data set. New search functionality includes spectrum name, peptide sequence and even sequence motif.



New High Mass Accuracy Validation

Sophisticated statistical techniques improve your identification rate with high mass accuracy data. Drift and accuracy are automatically tuned for each sample. Now you can fully realize your advanced instrumentation!