

Scaffold Software Meets Small Molecule LC-MS Analysis

Identify and quantify small molecules using MS1 or MS/MS data

Import CSV files containing sample metadata to automatically configure your experimental design and organize samples



Track isotopically labeled compounds using metabolic flux experiments

Load raw data* from major vendor instruments, including Thermo, Sciex, and Agilent

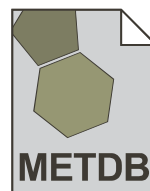
Search multiple spectral libraries including NIST, METLIN, HMDB, and LIPID MAPS

Quantify, determine statistical significance, and visualize results

Create personal spectral libraries using spectra from your instrument

Validate potential identifications using ID scores, fragmentation patterns, isotopic distributions, and chromatographic shapes

Share results with colleagues or customers using the free Scaffold Elements Viewer



*Windows is required to process vendor-formatted raw files