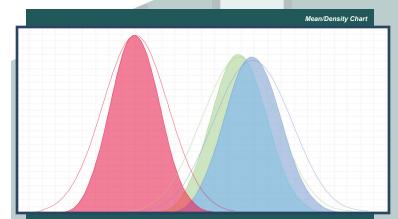


A Full Suite of Tools for Reporter Ion and SILAC Quantitation

Quantify proteins using iTRAQ, TMT, SILAC, or precursor intensity values



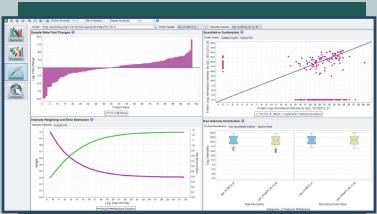
Ensure quantitative confidence regardless of experiment type with parametric and non-parametric statistical analysis

Determine significance and test for validity with assumption testing



Analyze various experimental designs, and normalize across samples using common or pooled references

Compare independent groups, repeated measures, or time-course experiments



Visualize protein-level changes across samples using fold changes, intensity values, or quantitative ratios

Define experiment-specific parameters using the experimental design wizard

Collaborate with colleagues or customers using sharable Scaffold files, Excel reports, and exportable images





