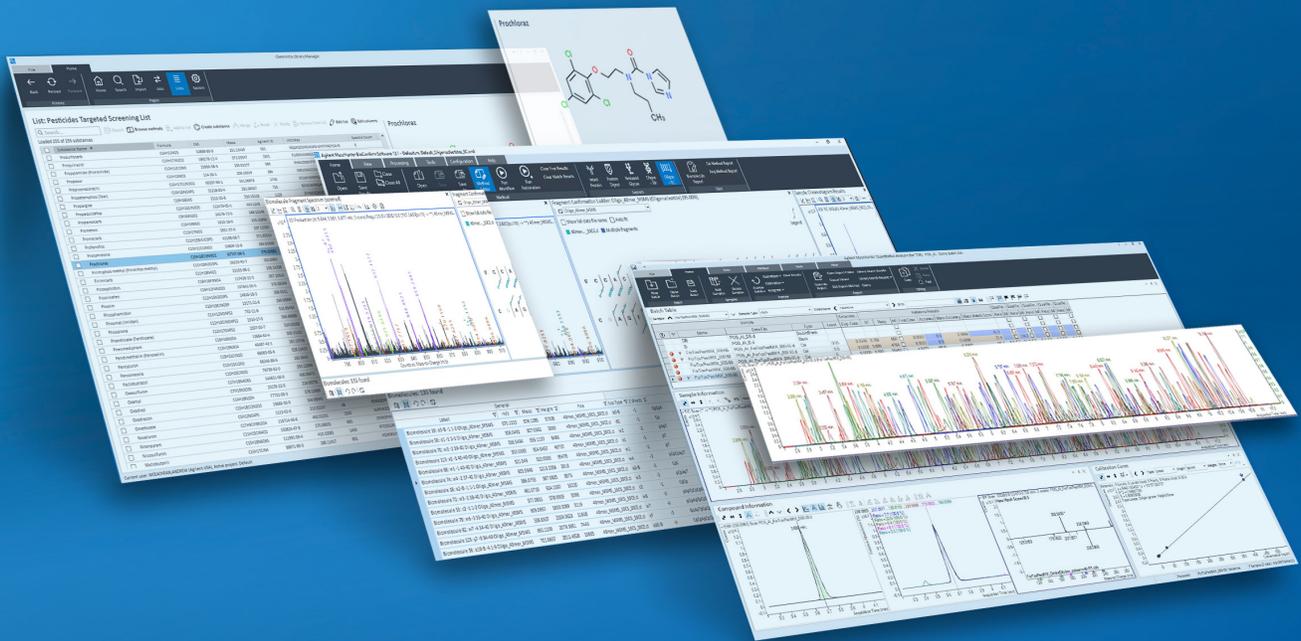


Maximum Throughput, Confident Results

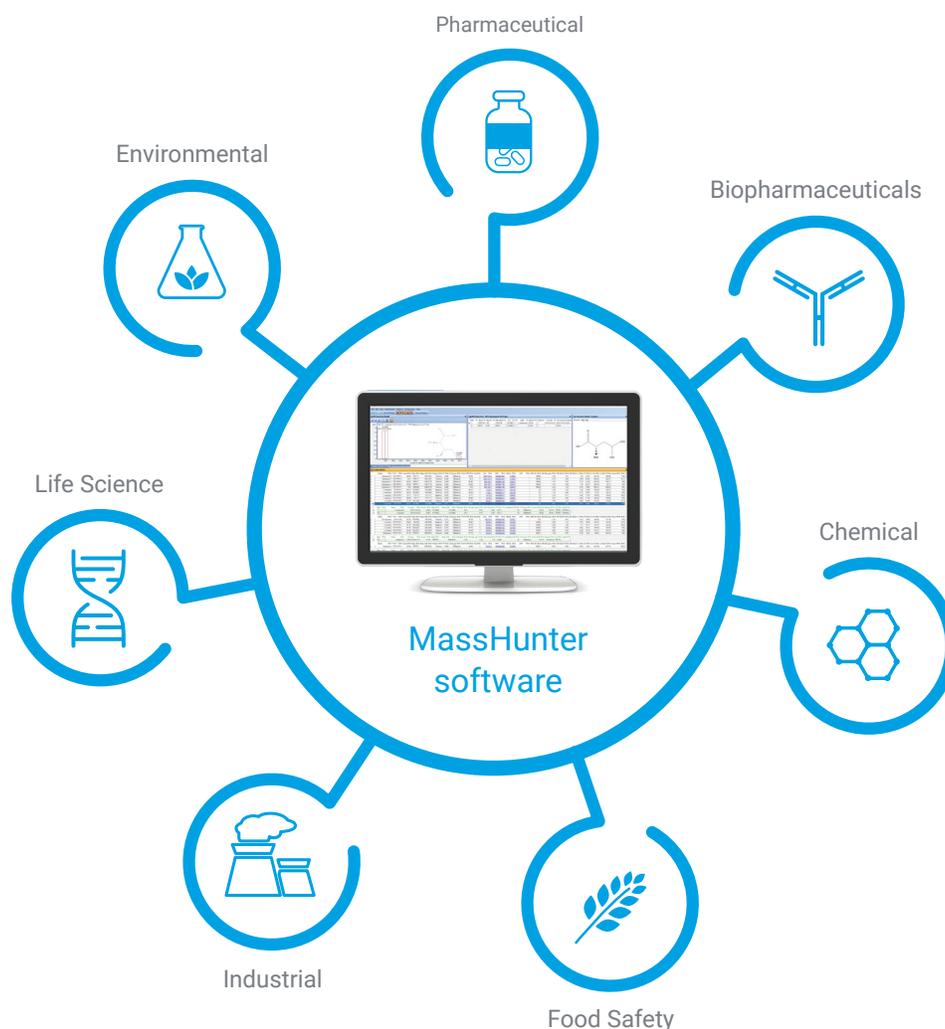
Agilent MassHunter software suite



Make Your Mass Spectrometry Analysis Faster, Easier, and More Productive

Designed to solve your everyday struggles, Agilent MassHunter software provides customizable features and capabilities that support diverse applications in mass spectrometry.

Operators at all levels can use MassHunter software to drive confident results for your laboratories. This intuitive software suite for Agilent mass spectrometers features easy-to-use methods and templates, intelligent instrument and automation control, as well as comprehensive curated databases and libraries.



Maintain Sustainability While Increasing Your Productivity

Lower your environmental impact and optimize your workflows

At Agilent, we believe that efficiency, productivity, and sustainability are interlinked. Together, we can help your lab achieve its sustainability goals—while increasing output, maintaining accuracy, and staying competitive.



Agilent 5977C GC/MSD



Agilent 7010C GC/TQ



Agilent 1290
Infinity II LC system



Agilent 6495D TQ LC/MS



Agilent Revident LC/Q-TOF



Agilent 8890 GC/Q-TOF



Agilent 6230 TOF LC/MS

 ACT. The Environmental Impact Factor Label.		US
Product Name		
Product Location		
SKU 0000		
Environmental Impact Scale Decreasing Environmental Impact		
		
Manufacturing		
Manufacturing Impact Reduction	3.0	
Renewable Energy Use	No	
Responsible Chemical Management	1.0	
Shipping Impact	1.0	
Product Content	10.0	
Packaging Content	5.0	
User Impact		
Energy Consumption (kWh/day)	12.7	
Water Consumption (gallons/day)	N/A	
Product Lifetime	1.0	
End of Life		
Packaging	4.8	
Product	1.0	
Innovation		
Innovative Practices	-1.0	
Environmental Impact Factor: 38.5		
Label Valid Through: June 2024		
 act.mygreenlab.org		

Selected Agilent instruments have been independently audited for the Accountability, Consistency, and Transparency (ACT) label. Similar to food nutrition labels, the ACT label provides a transparent, easy-to-understand measure for several factors related to sustainability.

MassHunter Acquisition Software

Simplify Data Acquisition

Make LC/MS and GC/MS instrument control easier with MassHunter acquisition software

Intuitive

Simple and easy to use, from instrument control to method setup.

Optimized

Autotune ensures reliable operation.

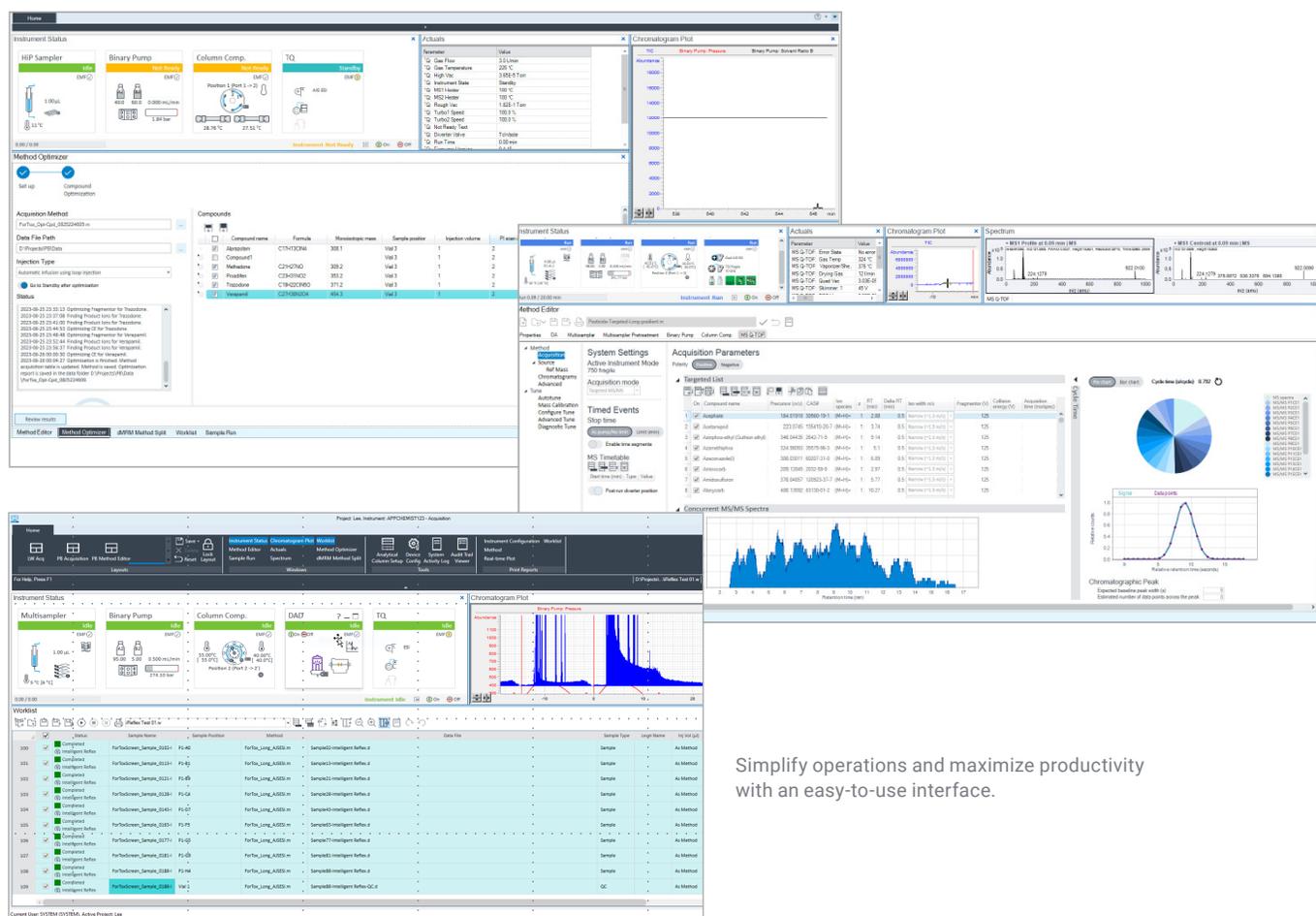
Automated

Intelligent reflex reinjection workflows maximize productivity, confidence, and throughput.

Intelligent

Maintenance feedback ensures highest instrument uptime.

Built-in intelligence. From instrument calibration to compound optimization and method development, MassHunter acquisition software gives you flexibility and delivers accurate answers quickly.



Simplify operations and maximize productivity with an easy-to-use interface.

Streamline Quantitation and Screening Workflows

Generate quantitative answers while preserving the integrity of your data

MassHunter Quantitative Analysis software simplifies compound identification and quantitation across all LC/TQ, LC/TOF, LC/Q-TOF, GC/MSD, GC/TQ, and GC/Q-TOF, from small- to large-target lists.

Features include:

- Built-in workflow templates
- Compliance-ready controls
- Rich data visualization
- Customizable data review panels

Easy data review provides flexibility for different users and workflows.

The screenshot displays the Agilent MassHunter Quantitative Analysis software interface. The main window shows a 'Batch Table' with columns for Name, Sample, Data File, Type, Level, Acq. Date-Time, Exp. Conc., RT, Resp, MI, Okc, Conc, Final Conc, Accuracy, Date, MI, Ratio, and MI. Below the table, there are several panels: 'Compound Information' showing a chromatogram with a peak at 3.12 min; 'Calibration Curve' showing a linear plot; 'Samples' table with columns for Name, File Name, Components, HPLC, and Acquire; 'Components' table with columns for Component, Compound Name, Molec. Weight, Formula, and Acquire; 'Chromatogram' showing a plot of Abundance vs. Acquisition Time (min); 'New Peaks' showing a zoomed-in view of a peak; 'Spectrum' showing a mass spectrum plot; and 'Molecular Structure' showing the chemical structure of the identified compound.

From quantitation to identifying and analyzing unknowns, built-in workflows give you confidence in your results.

Improve Confidence for Any Targeted Screening Assays

Enable method setup for all MS instruments and screening lists of any size

Easily manage large lists of targeted compounds using either ChemVista, Personal Compound Database and Library Manager (PCDL) software, MRM databases, or an acquisition method. All Agilent libraries include a comprehensive list of important compounds for accurate screening and identification through GC/MSD, GC/TQ, GC/Q-TOF, LC/TQ, LC/TOF, and LC/Q-TOF.

Workflows center around compound lists associated with a specific analysis.

Lists serve as organizational tools, allowing a compound to exist in multiple lists with multiple spectra.

Summary	
Substances	1402
Substances with spectra	962
Substances with RTs	180
Substances with method data	1078
Spectra	3681

Automated chemical class tags are assigned from manually curated PCDLs.

Enhanced cheminformatics generate structures and identifiers, eliminating duplicates and supporting downstream workflows.

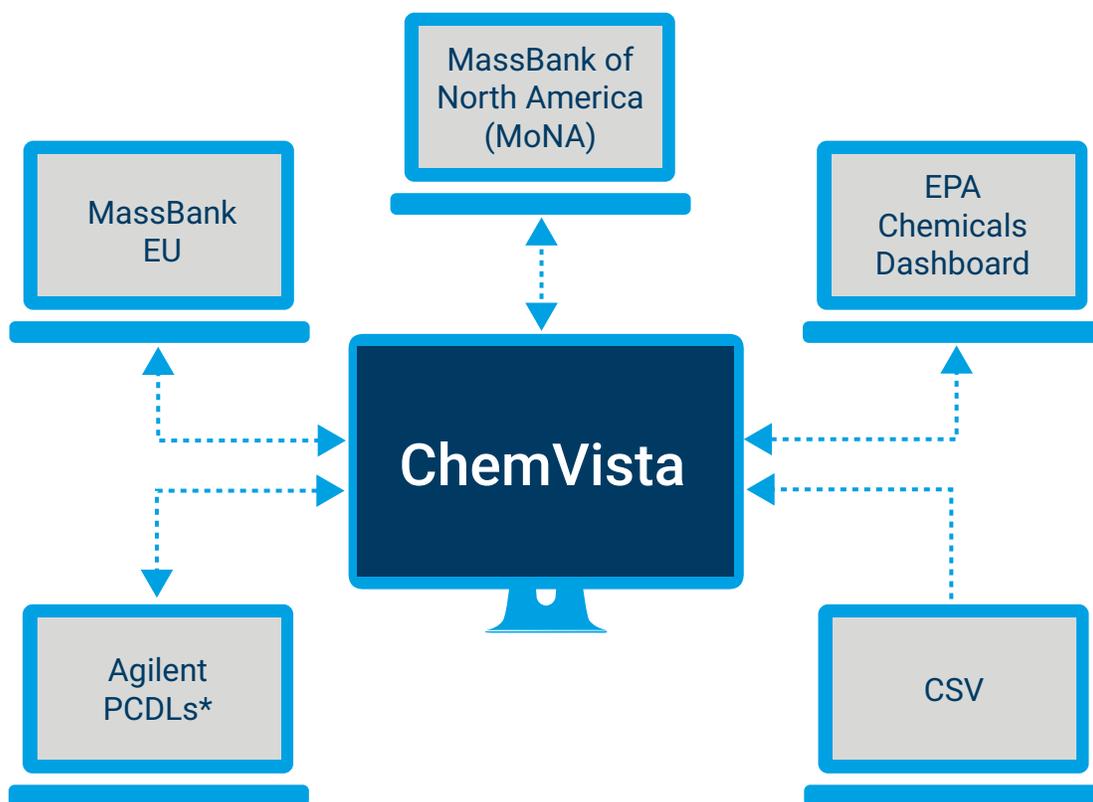
Import/export data with ChemVista in multiple file formats (SDF, MassBank Text, PCDL, CSV). To help you refine the exchanged information, these processes are supported with filters and customizable options.

Agilent ChemVista: An innovative tool for confident identification

Agilent ChemVista is a standalone software application that manages spectral libraries created by LC/Q-TOF and GC/Q-TOF mass spectrometry. It integrates compound details, retention time, and spectral information from multiple sources, allowing you to:

- Access multiple public databases and curated libraries (MassBank [EU and MoNA], EPA Chemicals Dashboard, Agilent PCDLs, and customer CSV).
- Organize, manage, edit, or create spectra.
- Facilitate identification workflows within MassHunter data analysis applications.
- Identify compounds with greater confidence.

In addition, ChemVista includes extensive preloaded library and database content.



**Curated Personal Compound Databases and Libraries*

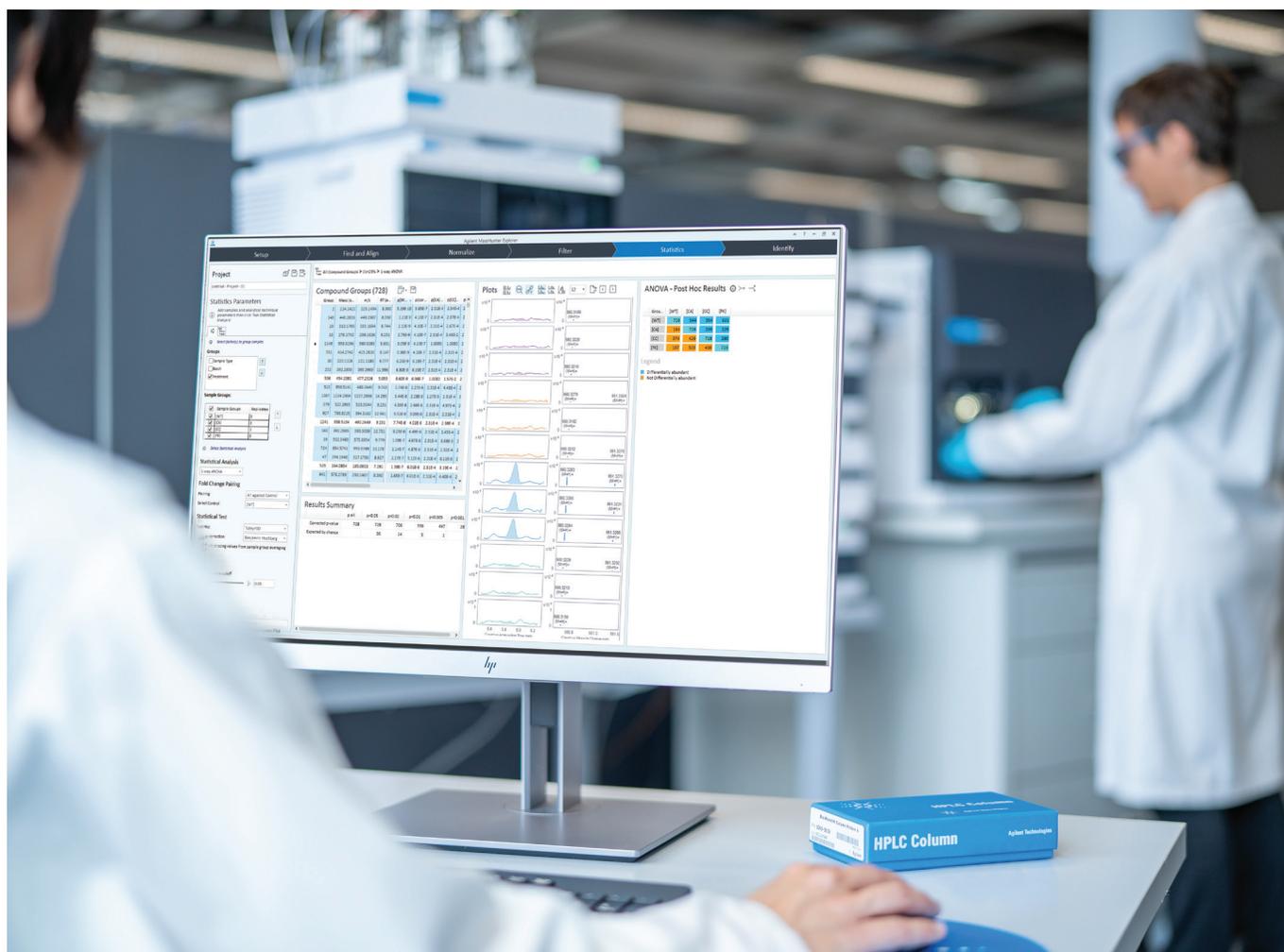
For a closer look at how ChemVista improves the efficiency and productivity of high-resolution mass spectrometry (HRMS) spectral management, [download](#) our technical overview.

Perform Fast, Reliable, and Simple Nontargeted Data Analysis

Make nontargeted data analysis easier, faster, and smarter

MassHunter Explorer features an integrated user interface that seamlessly combines advanced data extraction with normalization, statistical analysis, visualization, and identification procedures. Together, in one easy-to-use application, these tools help you confidently gain insights into your data quickly, improving your research productivity.

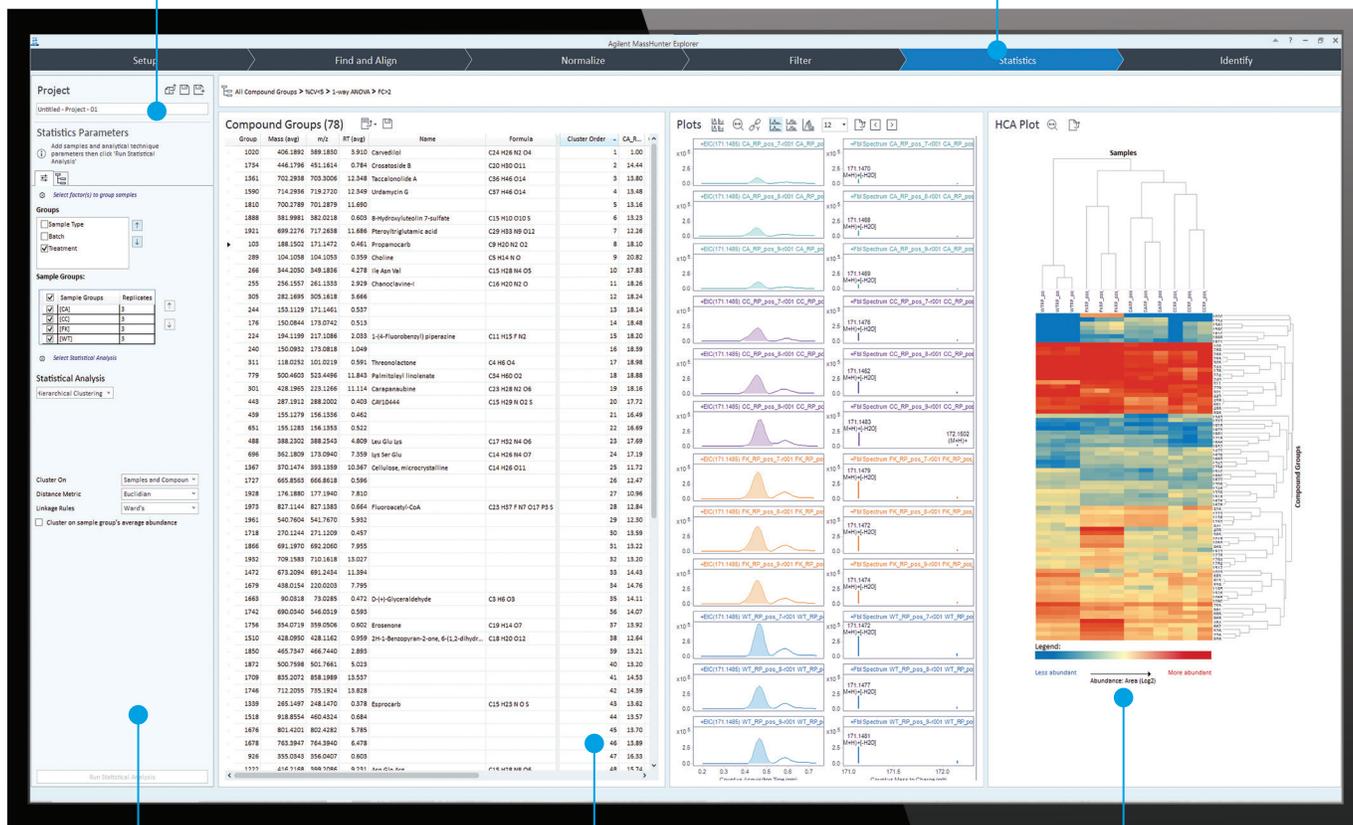
- Methods can be shared across your organization, ensuring consistent results.
- Project-based workflows allow for easy setup and transferability for beginners and experts.
- Different statistical analysis techniques help identify compounds that are significantly different or common between sample groups.



Quickly achieve accurate results with an integrated, guided, and intuitive workflow

Project-based workflows allow for easy setup and transferability for beginners and experts.

Streamlined, integrated procedures simplify navigation.



Multiple statistical algorithms provide the right tools for the given data.

Preprocessing during acquisition speeds up data processing.

Chromatographic, spectral, and statistical data are pictured for quick review.

Take a Giant Step Forward in Protein and Oligonucleotide Characterization

Enable routine characterization through easy-to-use workflows

Agilent MassHunter BioConfirm software is ideal for intact protein analysis, oligonucleotide characterization, peptide mapping, and routine glycan profiling.

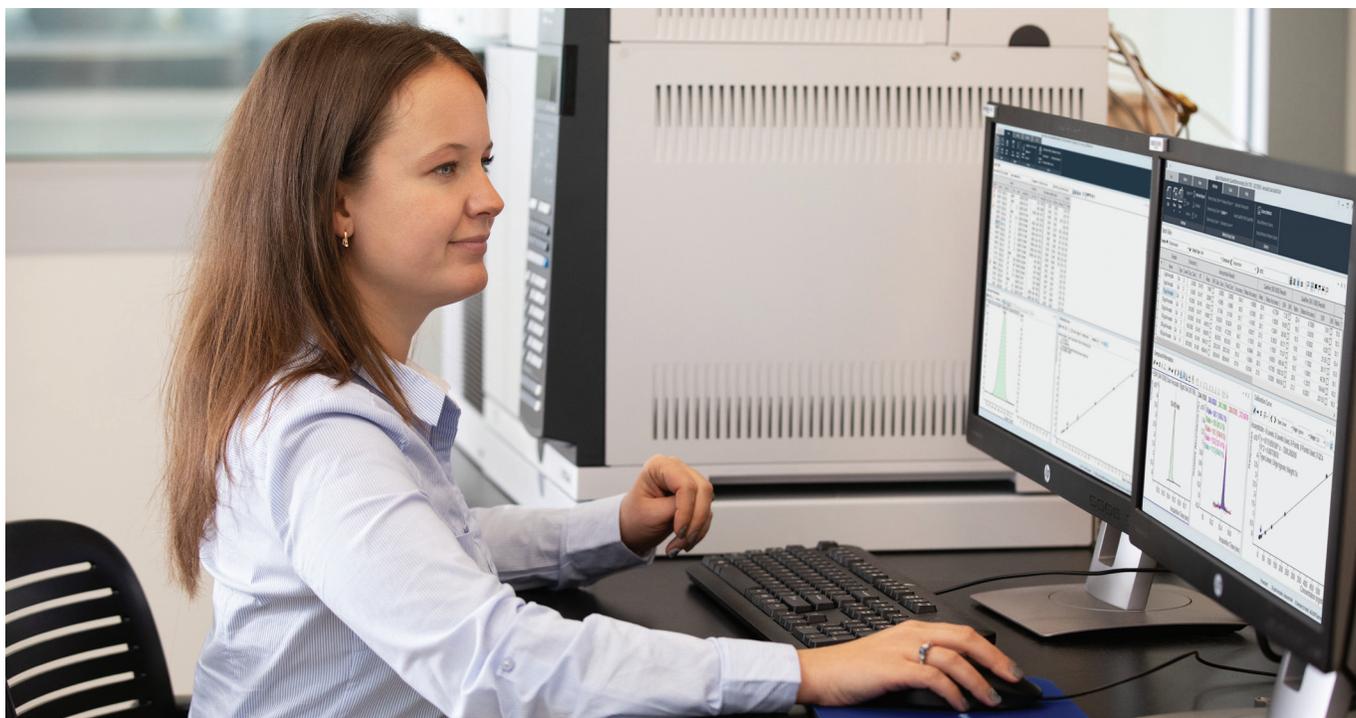
- Confirm biomolecule sequences with confidence.
- Rapidly deconvolute intact biomolecule data and confirm peak composition and potential modifications.
- Analyze in a secure, auditable manner using technical controls such as roles, permissions, and audit trail.

Intuitive windows give you increased access to information for sequence confirmation and target-plus-impurity workflows for oligonucleotides and proteins.

The screenshot displays the Agilent MassHunter BioConfirm Software 12.1 interface with several key windows open:

- Biomolecule Fragment Spectrum (Zoomed):** Shows a mass spectrum with peaks labeled with m/z values (e.g., 752.5, 810.5, 831.5, 852.5, 873.5, 894.5, 915.5, 936.5, 957.5, 978.5, 999.5, 1020.5, 1041.5, 1062.5, 1083.5, 1104.5, 1125.5, 1146.5, 1167.5, 1188.5, 1209.5, 1230.5, 1251.5, 1272.5, 1293.5, 1314.5, 1335.5, 1356.5, 1377.5, 1398.5, 1419.5, 1440.5, 1461.5, 1482.5, 1503.5, 1524.5, 1545.5, 1566.5, 1587.5, 1608.5, 1629.5, 1650.5, 1671.5, 1692.5, 1713.5, 1734.5, 1755.5, 1776.5, 1797.5, 1818.5, 1839.5, 1860.5, 1881.5, 1902.5, 1923.5, 1944.5, 1965.5, 1986.5, 2007.5, 2028.5, 2049.5, 2070.5, 2091.5, 2112.5, 2133.5, 2154.5, 2175.5, 2196.5, 2217.5, 2238.5, 2259.5, 2280.5, 2301.5, 2322.5, 2343.5, 2364.5, 2385.5, 2406.5, 2427.5, 2448.5, 2469.5, 2490.5, 2511.5, 2532.5, 2553.5, 2574.5, 2595.5, 2616.5, 2637.5, 2658.5, 2679.5, 2700.5, 2721.5, 2742.5, 2763.5, 2784.5, 2805.5, 2826.5, 2847.5, 2868.5, 2889.5, 2910.5, 2931.5, 2952.5, 2973.5, 2994.5, 3015.5, 3036.5, 3057.5, 3078.5, 3099.5, 3120.5, 3141.5, 3162.5, 3183.5, 3204.5, 3225.5, 3246.5, 3267.5, 3288.5, 3309.5, 3330.5, 3351.5, 3372.5, 3393.5, 3414.5, 3435.5, 3456.5, 3477.5, 3498.5, 3519.5, 3540.5, 3561.5, 3582.5, 3603.5, 3624.5, 3645.5, 3666.5, 3687.5, 3708.5, 3729.5, 3750.5, 3771.5, 3792.5, 3813.5, 3834.5, 3855.5, 3876.5, 3897.5, 3918.5, 3939.5, 3960.5, 3981.5, 4002.5, 4023.5, 4044.5, 4065.5, 4086.5, 4107.5, 4128.5, 4149.5, 4170.5, 4191.5, 4212.5, 4233.5, 4254.5, 4275.5, 4296.5, 4317.5, 4338.5, 4359.5, 4380.5, 4401.5, 4422.5, 4443.5, 4464.5, 4485.5, 4506.5, 4527.5, 4548.5, 4569.5, 4590.5, 4611.5, 4632.5, 4653.5, 4674.5, 4695.5, 4716.5, 4737.5, 4758.5, 4779.5, 4800.5, 4821.5, 4842.5, 4863.5, 4884.5, 4905.5, 4926.5, 4947.5, 4968.5, 4989.5, 5010.5, 5031.5, 5052.5, 5073.5, 5094.5, 5115.5, 5136.5, 5157.5, 5178.5, 5199.5, 5220.5, 5241.5, 5262.5, 5283.5, 5304.5, 5325.5, 5346.5, 5367.5, 5388.5, 5409.5, 5430.5, 5451.5, 5472.5, 5493.5, 5514.5, 5535.5, 5556.5, 5577.5, 5598.5, 5619.5, 5640.5, 5661.5, 5682.5, 5703.5, 5724.5, 5745.5, 5766.5, 5787.5, 5808.5, 5829.5, 5850.5, 5871.5, 5892.5, 5913.5, 5934.5, 5955.5, 5976.5, 5997.5, 6018.5, 6039.5, 6060.5, 6081.5, 6102.5, 6123.5, 6144.5, 6165.5, 6186.5, 6207.5, 6228.5, 6249.5, 6270.5, 6291.5, 6312.5, 6333.5, 6354.5, 6375.5, 6396.5, 6417.5, 6438.5, 6459.5, 6480.5, 6501.5, 6522.5, 6543.5, 6564.5, 6585.5, 6606.5, 6627.5, 6648.5, 6669.5, 6690.5, 6711.5, 6732.5, 6753.5, 6774.5, 6795.5, 6816.5, 6837.5, 6858.5, 6879.5, 6900.5, 6921.5, 6942.5, 6963.5, 6984.5, 7005.5, 7026.5, 7047.5, 7068.5, 7089.5, 7110.5, 7131.5, 7152.5, 7173.5, 7194.5, 7215.5, 7236.5, 7257.5, 7278.5, 7299.5, 7320.5, 7341.5, 7362.5, 7383.5, 7404.5, 7425.5, 7446.5, 7467.5, 7488.5, 7509.5, 7530.5, 7551.5, 7572.5, 7593.5, 7614.5, 7635.5, 7656.5, 7677.5, 7698.5, 7719.5, 7740.5, 7761.5, 7782.5, 7803.5, 7824.5, 7845.5, 7866.5, 7887.5, 7908.5, 7929.5, 7950.5, 7971.5, 7992.5, 8013.5, 8034.5, 8055.5, 8076.5, 8097.5, 8118.5, 8139.5, 8160.5, 8181.5, 8202.5, 8223.5, 8244.5, 8265.5, 8286.5, 8307.5, 8328.5, 8349.5, 8370.5, 8391.5, 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15573.5, 15594.5, 15615.5, 15636.5, 15657.5, 15678.5, 15699.5, 15720.5, 15741.5, 15762.5, 15783.5, 15804.5, 15825.5, 15846.5, 15867.5, 15888.5, 15909.5, 15930.5, 15951.5, 15972.5, 15993.5, 16014.5, 16035.5, 16056.5, 16077.5, 16098.5, 16119.5, 16140.5, 16161.5, 16182.5, 16203.5, 16224.5, 16245.5, 16266.5, 16287.5, 16308.5, 16329.5, 16350.5, 16371.5, 16392.5, 16413.5, 16434.5, 16455.5, 16476.5, 16497.5, 16518.5, 16539.5, 16560.5, 16581.5, 16602.5, 16623.5, 16644.5, 16665.5, 16686.5, 16707.5, 16728.5, 16749.5, 16770.5, 16791.5, 16812.5, 16833.5, 16854.5, 16875.5, 16896.5, 16917.5, 16938.5, 16959.5, 16980.5, 17001.5, 17022.5, 17043.5, 17064.5, 17085.5, 17106.5, 17127.5, 17148.5, 17169.5, 17190.5, 17211.5, 17232.5, 17253.5, 17274.5, 17295.5, 17316.5, 17337.5, 17358.5, 17379.5, 17400.5, 17421.5, 17442.5, 17463.5, 17484.5, 17505.5, 17526.5, 17547.5, 17568.5, 17589.5, 17610.5, 17631.5, 17652.5, 17673.5, 17694.5, 17715.5, 17736.5, 17757.5, 17778.5, 17799.5, 17820.5, 17841.5, 17862.5, 17883.5, 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20235.5, 20256.5, 20277.5, 20298.5, 20319.5, 20340.5, 20361.5, 20382.5, 20403.5, 20424.5, 20445.5, 20466.5, 20487.5, 20508.5, 20529.5, 20550.5, 20571.5, 20592.5, 20613.5, 20634.5, 20655.5, 20676.5, 20697.5, 20718.5, 20739.5, 20760.5, 20781.5, 20802.5, 20823.5, 20844.5, 20865.5, 20886.5, 20907.5, 20928.5, 20949.5, 20970.5, 20991.5, 21012.5, 21033.5, 21054.5, 21075.5, 21096.5, 21117.5, 21138.5, 21159.5, 21180.5, 21201.5, 21222.5, 21243.5, 21264.5, 21285.5, 21306.5, 21327.5, 21348.5, 21369.5, 21390.5, 21411.5, 21432.5, 21453.5, 21474.5, 21495.5, 21516.5, 21537.5, 21558.5, 21579.5, 21600.5, 21621.5, 21642.5, 21663.5, 21684.5, 21705.5, 21726.5, 21747.5, 21768.5, 21789.5, 21810.5, 21831.5, 21852.5, 21873.5, 21894.5, 21915.5, 21936.5, 21957.5, 21978.5, 21999.5, 22020.5, 22041.5, 22062.5, 22083.5, 22104.5, 22125.5, 22146.5, 22167.5, 22188.5, 22209.5, 22230.5, 22251.5, 22272.5, 22293.5, 22314.5, 22335.5, 22356.5, 22377.5, 22398.5, 22419.5, 22440.5, 22461.5, 22482.5, 22503.5, 22524.5, 22545.5, 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