Agilent

Agilent OpenLab ChemStation
For InfinityLab LC/MSD Series and 6100 Series LC/MS
Quick Start Guide

Where to find more information 2
To start the OpenLab ChemStation program 4

Configuration Diagrams 5

InfinityLab LC/MSD Series with 1100/1200/1260/1290 Series LC 5

OpenLab ChemStation Views 6

Method and Run Control View 7
Data Analysis View 10
Report Layout View 13
Diagnosis View 14
MSD Tune View 15

Basic Operation 16

Start up and shut down the system 16
Tune the MSD 17
Acquire MSD Data 19

Deconvolution 21

To deconvolute a mass spectrum 21
Deconvolution Report 22

Operating Tips 23

Troubleshooting Tips 25

This manual applies to OpenLab ChemStation LTS 01.11 software.

Where to find more information

Online Help Use online Help for in-depth information beyond what is shown in this *Quick Start Guide*. Display online Help in any of the following ways:

- Click the **Help** button in the toolbar.
- Select Help Topics from the **Help** menu.
- Click the **Help** button on most dialog boxes to show task-specific help.

User Guides Besides this *Quick Start Guide*, the following guides are available from the Resource App after installation.

- *Agilent OpenLab ChemStation Familiarization Guide*
- *Agilent InfinityLab LC/MSD Series System Installation Guide*
- *Agilent InfinityLab LC/MSD Series and 6100 Series LC/MS Maintenance Guide*

After installation, open the LC-MSD Resource app to access MSD-specific user guides and training material. For other user guides, refer to the **OpenLab ChemStation User Resources & Learning Platform**.

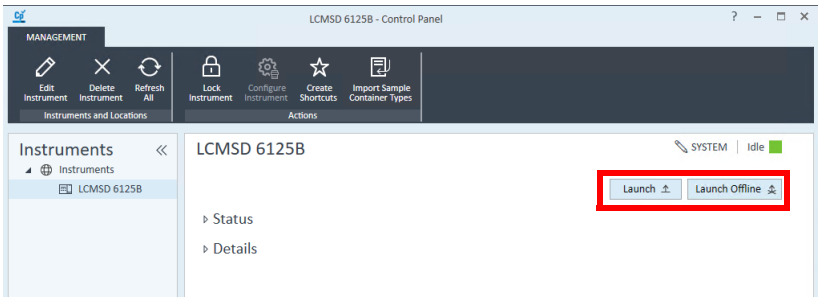
Getting Started Self-paced training modules to learn about the mass selective detector and OpenLab ChemStation software are available from the Resource App.

To start the OpenLab ChemStation program



1 Click the **Control Panel** icon on the Desktop to open the **Control Panel**.

2 Click **Launch**.

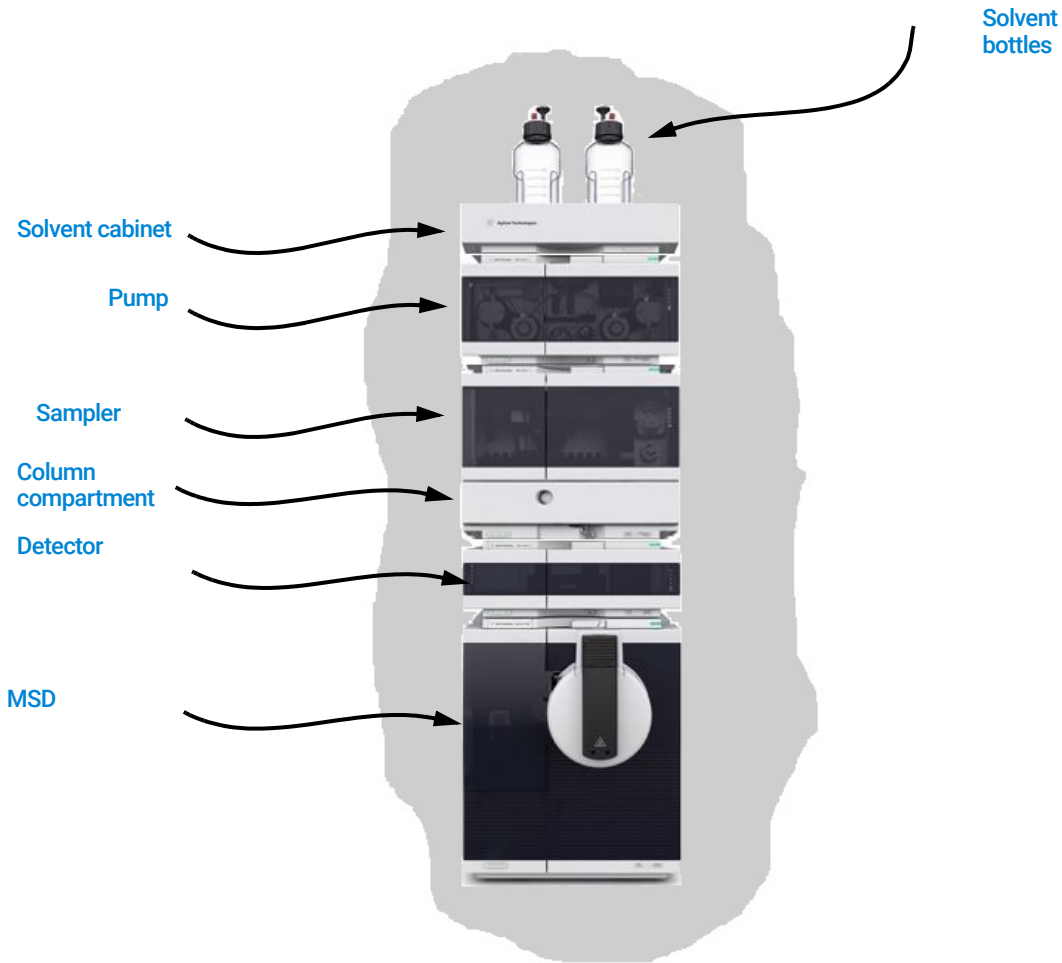


3 Click the ChemStation icon.



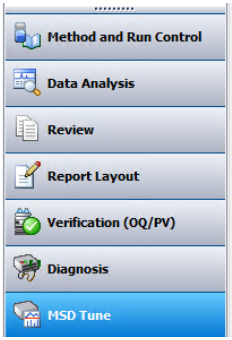
Configuration Diagrams

InfinityLab LC/MSD Series with 1100/1200/1260/1290 Series LC



OpenLab ChemStation Views

The OpenLab ChemStation software consists of several views. Click one of the view buttons in the lower left corner of the screen, or from the **View** menu, to change to a different view.



Method and Run Control View Allows you to set up methods and adjust instrument parameters to inject samples and acquire data one sample at a time or in automated sequences. See the next page for more information.

Data Analysis View Allows you to perform various data evaluation tasks on chromatograms and spectra, such as integration, quantitation, checking peak purity, deconvolution, and report generation. See **page 9** for more information.

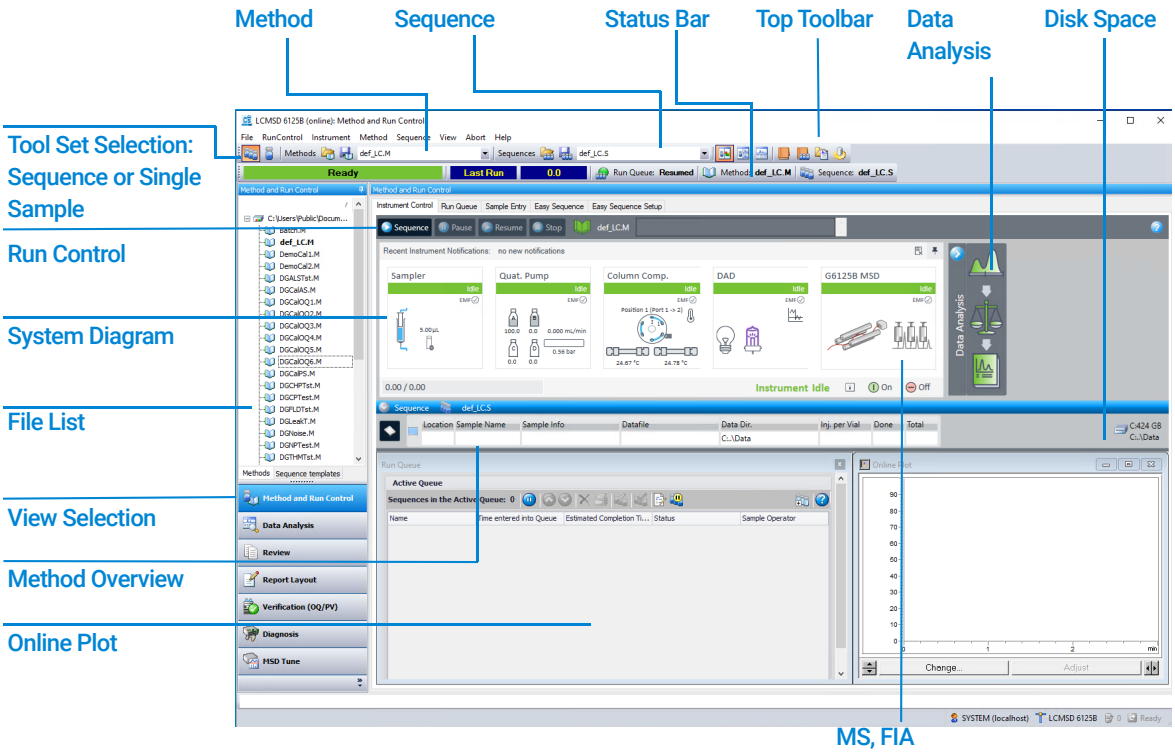
Report Layout View Allows you to design custom report templates to use with the data generated by your OpenLab ChemStation. See **page 12** for more information.

Verification (OQ/PV) This view is no longer supported. Use the Agilent Automated Compliance Engine (ACE) for your verification needs.

Diagnosis View Allows you to run tests to diagnose instrument problems and access information on how to resolve these problems. The early maintenance feedback (EMF) feature can notify you when it is time to perform system maintenance before a problem occurs. See **page 13** for more information.

MSD Tune View In this view, you can calibrate your MSD automatically. You can also set MS parameters manually for specific types of molecules. See **page 14** for more information.

Method and Run Control View



NOTE

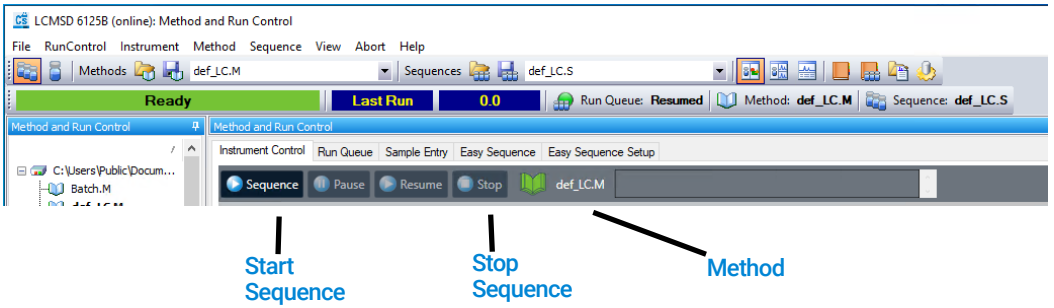
The user interface for Method and Run Control can be configured in two ways during installation. See *Agilent InfinityLab LC/MSD Series System Installation Guide* for information on installation and configuration.

If your instrument is configured with the Classic user interface, then see OpenLab ChemStation online Help and these manuals on the Resource App:

- *Quick Start Guide*
- *Familiarization Guide*
- *Concepts Guide*

Status and Run Bars: Instrument Control Tab

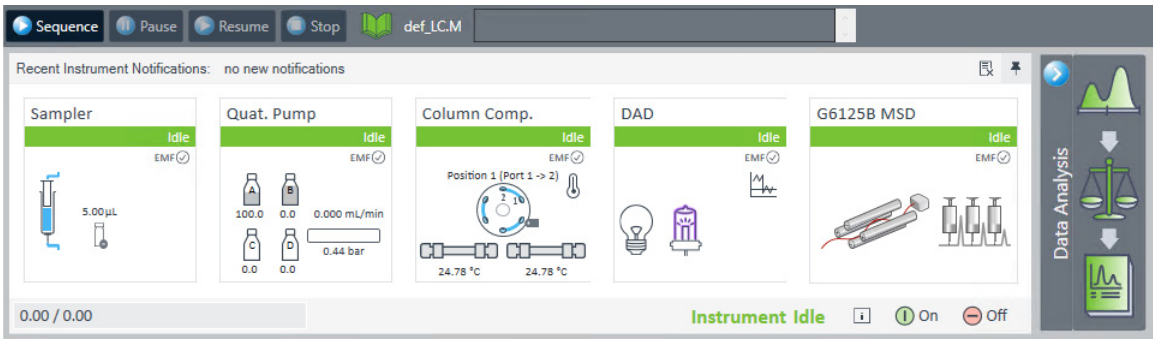
Right-click the Method icon to see the following commands: Run Time Checklist, Method Information, Edit Entire Method, Method Audit Trail, Print Method, and Help.



System Diagram

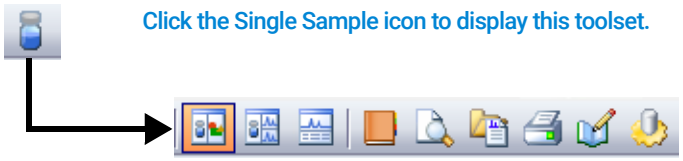
Each icon of the system diagram represents one component or module of your system. Click on an icon if you want to edit the method parameters or go to the online Help for that particular component.

Injector Pump & Solvents Column Detectors



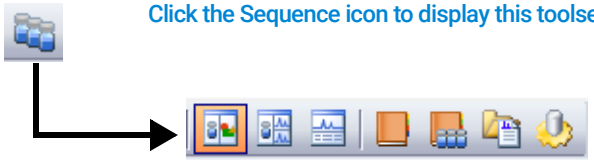
Single Sample Tool Set

This toolset allows you to work on methods and run an analysis for a single sample. Move the cursor over a button to view a description of it.



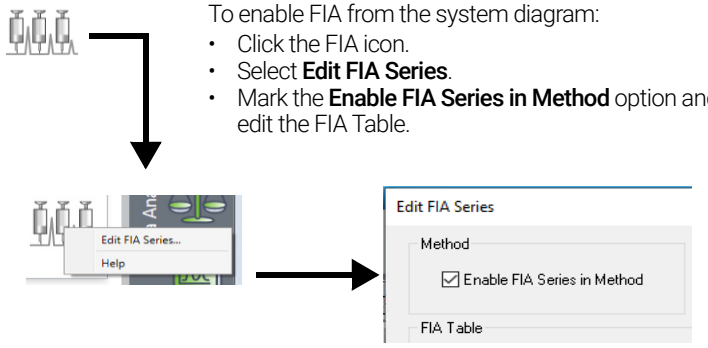
Sequence Tool Set

This toolset allows you to work on sequences and run automated analyses of multiple samples. Move the cursor over a button to view a description of it.

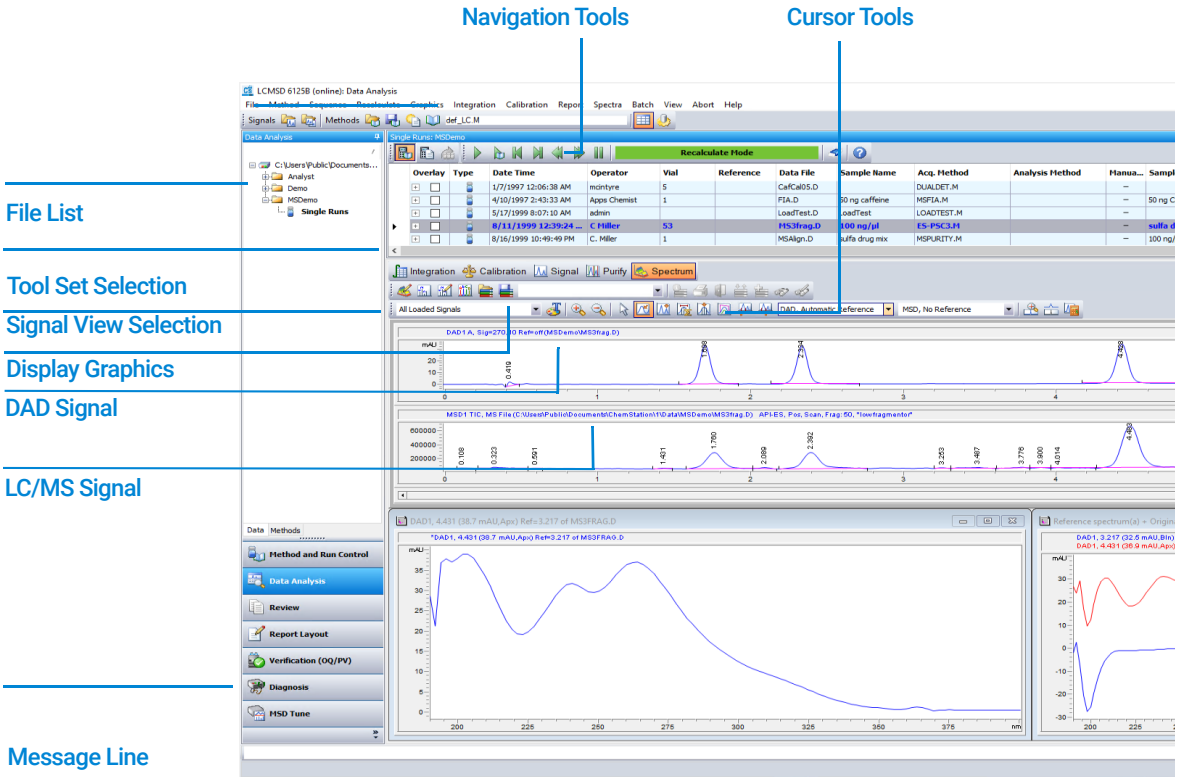


Flow Injection Analysis (FIA)

This toolset allows you to inject multiple samples directly into the detector, which lets you bypass the chromatographic column. The results are sent to a single data file. FIA can be used for method development or for applications that do not require chromatography.



Data Analysis View

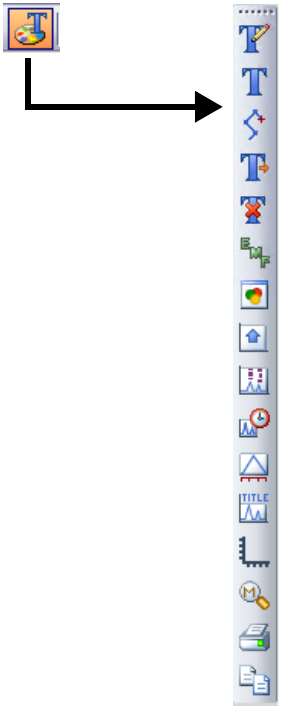


The various tool sets available in Data Analysis View are shown on the following pages. You can view a description of a button by moving the cursor over it.

Common Tool Set This toolset is always present in Data Analysis view.



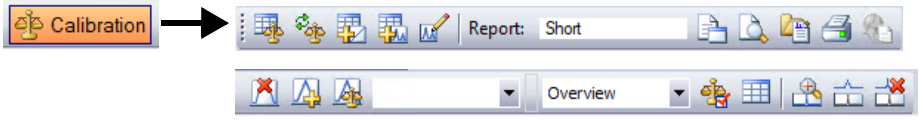
Graphics Tool Set Allows you to manipulate the graphic display.



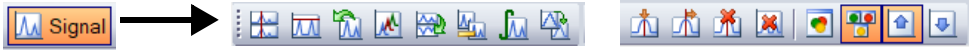
Integration Tool Set Allows you to perform integration and reporting tasks on a chromatogram.



Calibration Tool Set Allows you to perform calibration tasks for quantitation.



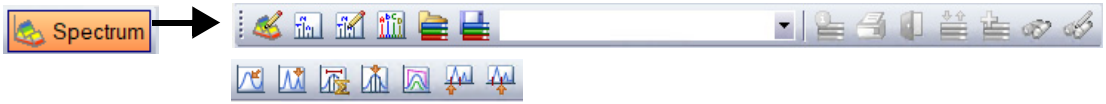
Signal Tool Set Allows you to work graphically with the UV or MS signal.



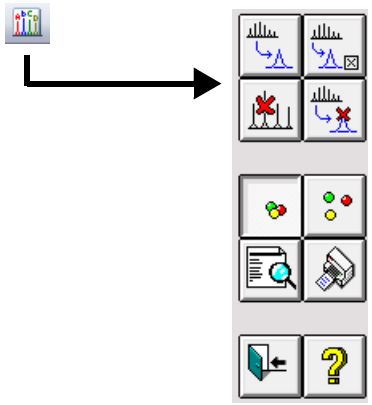
Purify Tool Set Allows you to work graphically with your purification data.



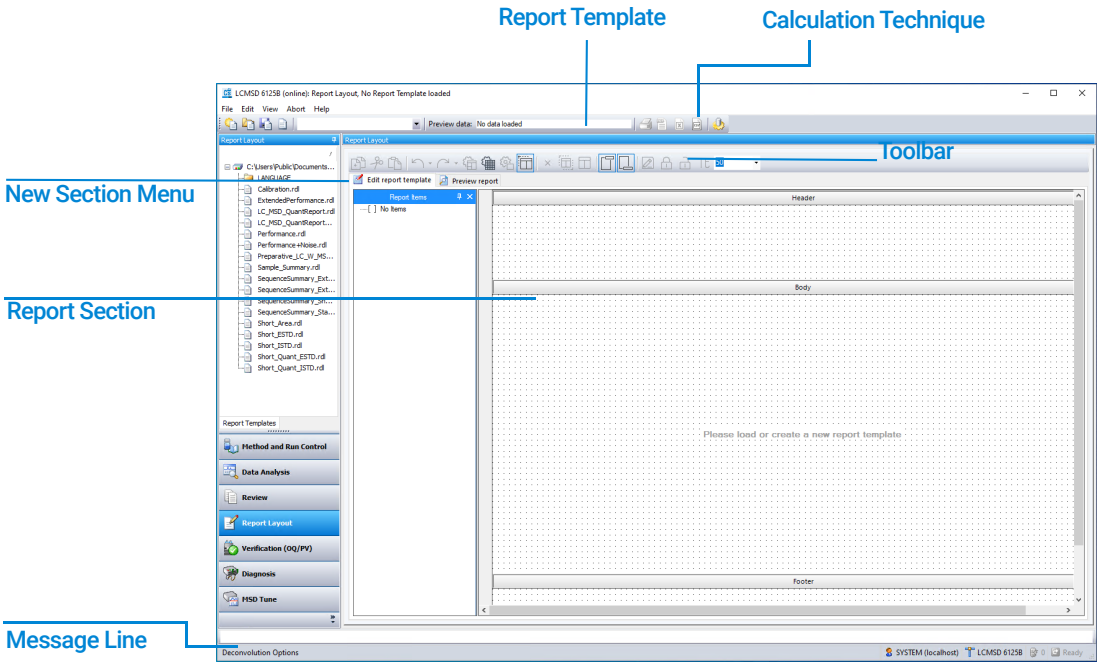
Spectral Tool Set Allows you to perform spectral evaluation tasks.



Deconvolution Tool Set Allows you to transforms mass spectra from multiply-charged ions into a calculated molecular weight.



Report Layout View



Report Layout

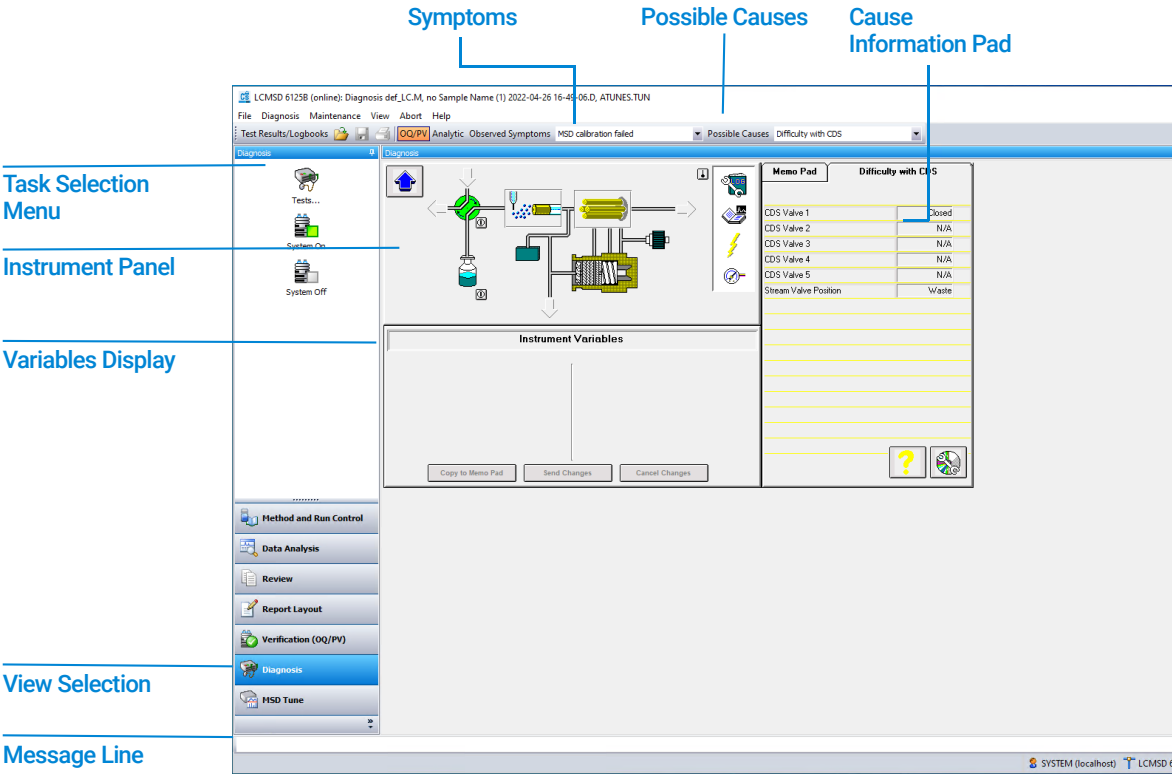
Tool Set This toolset is present in the Report Layout view. Move the cursor over a button to view a description of it.



Tips

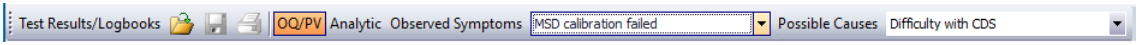
- Edit or create a report layout as described in the online Help.
- To test your report layout, select a calculation technique from the drop-down list box to define how the results are calculated.
- Load a data file. The results are loaded into the report template using the selected calculation.
- Select **File / Add to Report Styles** to add your completed report template to the list of available report styles. Now you can use your customized report within a method.

Diagnosis View

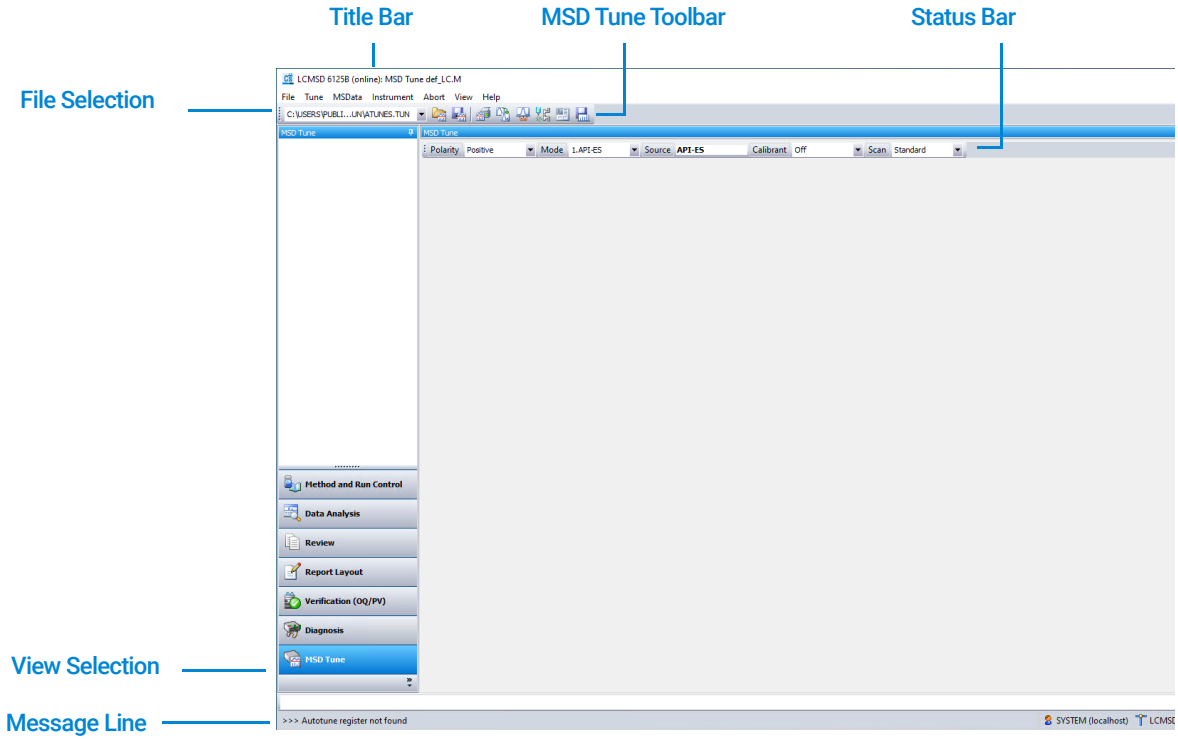


This view allows you to carry out diagnostic and maintenance activities on your system.

Diagnosis Toolbar The Diagnosis toolbar is displayed when you select Show Top Toolbar from the View menu. Move the cursor over button to view a description of it.



MSD Tune View



MSD Tune Toolset

This toolset is present in MSD Tune view.

	Load an MSD tune file.		Calibrate the mass axis.
	Save the current tune file.		Change spray chamber parameters.
	Generate a profile and scan report.		Edit MSD parameters for manual tuning.
	Autotune the MSD and print a tune report.		Acquire multiple scans and send results to data file.

Basic Operation

Start up and shut down the system

Task	Instruction	Comment
To start up the system	1 From the Method and Run Control view, click the ⏻ button on the system status bar. When you move mouse over the button, if system is currently idle, it turns green. 2 Set up the LC conditions (pump, column heater, and detector). 3 Set up the MSD spray chamber conditions as described in the MSD Spray Chamber dialog box topic in the online Help. Allow 15 minutes for the system to warm up.	This procedure assumes that the system is under vacuum and the LC and MSD are properly connected. It also assumes that the liquid flow path is properly set up.
To put the system in Standby Mode	1 Flush the system with pure mobile phase to ensure that the flow path is clear of buffers. This prevents the nebulizer from plugging. A plugged nebulizer can cause a high back pressure, which can damage the LC flow cell. It is good practice to flush the flow path (including the column) for 5 to 10 minutes with a mobile phase without buffers such as 50:50 water/acetonitrile. 2 Click the ⏻ button on the system status bar. All modules are set to standby mode (indicated by the gray color in the system diagram).	Put the system in standby mode overnight or whenever you are not analyzing samples for an extended time. The standby state for the MSD leaves the nebulizer and drying gas on at low flow. Standby for the MSD is: • 20 psi for the nebulizer, • 3 L/min for drying gas, • 300 °C for drying gas temperature, • 325 °C for the APCI vaporizer, if present • the MS stream selection valve is set LC to Waste. For Agilent Jet Stream source: • Sheath temp 100°C • Sheath gas flow 2.2 L/minute

Tune the MSD

When the MSD is used as a detector for the LC, a mass spectrum is associated with each data point in the LC chromatogram. To obtain high quality, accurate mass spectra, optimize the MSD to:

- Maximize sensitivity
- Maintain acceptable resolution
- Ensure accurate mass assignment

Tuning is the process of adjusting MSD parameters to achieve these goals. After the MSD parameters have been optimized, they must be saved in a tune file (.tun). This tune file is then specified in the method that is used to acquire data for your samples.

Frequent tuning, whether automated or manual, is not required. Once tuned, the MSD is stable. Tuning is needed no more often than monthly, weekly at most.

Wait 4 hours after pumpdown before tuning or operating your MSD. The analyzer takes about 11 hours to reach thermal equilibrium. Tune files that are created or data that is acquired before the MSD is at thermal equilibrium can have incorrect mass assignments and other inaccuracies.

Task	Instructions	Comments
To use Autotune	**1** From the MSD Tune view, select Autotune from the **Tune** menu or click the Autotune toolbar button. **2** Review the tune report that is printed automatically when tuning is completed.	Use autotune for automated adjustment of the MSD performance.

Task	Instructions	Comments
To use Check Tune	1 Select Check Tune from the **Tune** menu. 2 Review the Check Tune report. Adjustments are suggested for values that are outside of acceptable ranges.	Use Check Tune to determine quickly whether the MSD is correctly tuned. It performs a single profile scan of the tune masses and compares the peak widths and mass axes with target values.
To use Manual Tune	1 Select Manual Tune from the **Tune** menu. 2 Set the desired mass resolution (adjusting width gain and width offset). 3 Calibrate the mass axis (adjusting mass gain and mass offset). 4 Optimize the ion transmission through the source ion optics (fragmentor, skimmer(s), lens 1, lens 2, and octopole peak). 5 Adjust the signal strength (setting iris and adjusting the multiplier gain). Fragmentor and gain are method parameters. The fragmentor affects ion transmission and fragmentation. For more information, see the online Help.	Use Manual Tune when you want to: • Achieve maximum sensitivity by sacrificing some resolution • Tune specifically for the very low end (<150 amu) of the mass range • Tune with a compound other than the standard calibrants

Acquire MSD Data

Modes of Acquisition

There are three modes of acquiring data:

- Running a method for a single sample
- Running a sequence for multiple samples
- Running an FIA series

Note the following about acquiring data:

- All three acquisition modes require an appropriate method.
- Samples can be injected either manually or with an ALS.
- Always start a run from the software.
- An FIA method cannot be used in a sequence but multiple FIA methods can be run sequentially.

To edit a method and start a run

Once you know the acquisition mode that you want to use, you must set up an appropriate method. Methods are set up in the Method and Run Control view.

Instructions	Comments
1 Activate the single sample toolset by clicking the button on the toolbar.	
2 Load a method by selecting Method / Load Method or by clicking the button on the toolbar and selecting a method from the list.	
3 Select Edit Entire Method from the **Method** menu or right-click the icon on the Run Control Bar.	
4 Select the method sections you want to edit (select all sections to become familiar with the method parameters that are available) and click OK.	A series of dialog boxes is displayed, which allows you to set up your method and instrument parameters.
5 Add any method comments you want to appear on your reports. **6** Set up the instrument parameters, such as pump, injector, DAD or VWD, column thermostat, MS signals, and MS spray chamber. **7** Set up the Data Analysis parameters, such as signal details, integration events, report parameters, instrument curves, calibration curves, calibration table, and ion parameters. **8** Complete the Run Time Checklist.	Click the Help button on any of the method dialog boxes for descriptive information on the items available in each.
9 Save the method using a different name. Select Method / Save Method As or click the button on the toolbar.	Once you are familiar with the options that are available, use the system diagram menus for quick access to particular method parameters.
10 When you are ready to begin a run, click the Start button.	

Deconvolution

Deconvolution is an optional process (additional license required) that transforms mass spectra from multiply-charged ions into a calculated molecular weight.

To deconvolute a mass spectrum

Instructions	Comments
1 In the Data Analysis view, load an electrospray MS data file. The TIC is displayed.	
2 Generate the MS spectrum that you want to deconvolute. (Select the Spectral task tool set and use the spectrum selection tools to pick the spectrum of interest).	These operations are performed on data acquired in full scan mode.
3 Click the Enter Deconvolution Tool button to set up the deconvolution display area. 	
4 When the spectrum is displayed, click the Label Ions button to locate masses to be used in deconvolution. 	
5 Examine the ions that were found. Very noisy data or data with unresolved regions often need special settings to find ions.	You can use the Ion Label Options button to optimize the set of ions found. 
6 Click the Edit Deconvolution Parameters button to change any of the parameters. 	
7 Click the Run Deconvolution button to begin the deconvolution process. 	Once the deconvolution is done, the components are displayed in the upper right window. The charge states are displayed in the lower right window.

Instructions	Comments
8 Select specific components from the Component list to look at individual components or groups of components.	Optionally, click the Preview Deconvolution Report button  or click the Print Report button. 
9 Optionally, select components from the Component List box and then click the Delete Components button.  **10** Repeat the process.	This action removes the peaks for the selected components from the original spectrum. You can then continue with step 4 (previous), using this new spectrum.

Deconvolution Report

The Deconvolution report contains summary information about each component selected, along with detailed information on how each peak in a component contributes to the molecular weight of the component. In the first part of the report, the components are ranked by percent relative abundance, which is useful for estimating the percent of impurities.

The actual molecular weight can differ from the computed molecular weight even if the data fit a Gaussian curve perfectly, due to other errors such as errors in mass axis assignment or unresolved chemical impurities.

The Deconvolution software is optional and is part of the OpenLab ChemStation or M8363AA ChemStation LC/MS Deconvolution Bioanalysis Software package.

Operating Tips

- Back up your data and methods regularly to avoid loss of data in case the files are accidentally overwritten, deleted, or if a hardware problem develops with your disk drive.
- Put the system in standby mode overnight or whenever you are analyzing samples for an extended time.
- Verify that the tune file you are using is appropriate for your samples.
- Save the Tune reports as a PDF file or print the report, as part of your maintenance log.
- Regular system maintenance can reduce problems. Keep a maintenance record.
- Use the Maintenance Logbook and EMF features in Diagnosis view to help you track when maintenance is needed and to keep an online maintenance record.
- Flush the sample path and clean the spray chamber, capillary tip, and spray shield daily or at the end of each shift. Check the foreline pump fluid level every week.
- The spray chamber vent hose must be connected to a lab vent that is used only for the source. The vent hose must be separate from the vent hose for the foreline pump. Otherwise, pump exhaust can migrate into the spray chamber vent producing chemical noise.
- Samples must be filtered. If no chromatography is used, they must be salt and detergent-free.
- If a UV detector is available, use it in series with the MSD. Try to minimize chromatographic peak broadening by using low dispersion tubing.

- To minimize chromatographic band broadening, be sure that all tubing connections are free of dead volume. Use zero dead-volume (ZDV) fittings when possible.
- Use the following table as a guide to using SIM, condensed scan, and full scan acquisition modes.

Task	Mode
Acquire electrospray data for samples containing large, multiply-charged analytes.	Full Scan
Analyze a mixture with unknown components (small molecules).	Scan
Analyze a mixture with known components in unknown amounts (quantitate).	Scan or SIM
Identify the presence of a few known compounds at low levels within a mixture.	SIM
Use fast chromatography on 1.0-MHz MSD systems (LC/MSD and LC/MSD XT) without missing peaks.	Fast Scan

Troubleshooting Tips

No peaks

? Verify that there is spray from the nebulizer.

? Verify that the capillary voltage is set correctly.

? In APCI mode, verify that the corona needle is installed.

In APCI mode, verify that the Corona Current setting is 2 µA or higher, and that the Corona Voltage is between 1500 and 5000 V.

? Verify that the MS system is tuned correctly.

? Verify that MS vacuum pressures are within normal ranges.

? Check the drying gas flow and temperature.

? Verify that the fragmentor is set correctly.

? Check if any error message indicates a problem with electronics.

? Verify that sufficient sample is present in vial.

Poor mass accuracy

? Recalibrate the mass axis.

? Verify that the ions used for tuning span the mass range of the sample ions and that they show strong, stable signals.

Low signal

? Check the solution chemistry. Verify that the solvent you are using is appropriate for your sample. Mixed samples can exhibit signal suppression of one or more components.

? Verify that the sample is fresh and has been stored correctly.

? Verify that the MS system is tuned correctly.

? Check the nebulizer condition.

? Clean the capillary entrance.

? Check the capillary for damage and contamination.

? Verify that the spray chamber parameters are set correctly, such as drying gas heater and nebulizer pressure.

Unstable signal

? Verify that the drying gas flow and temperature are correct for the solvent flow you are using.

? Verify that the solvent is thoroughly degassed. Do not use ultrasonic degassing with protein samples.

? Verify that the LC backpressure is steady; steady backpressure indicates a steady solvent flow.

High spectral noise

? Use appropriate mass filter values.

? Check the spray shape. Nebulizer can be damaged or incorrectly set.

? Verify that drying gas flow and temperature are correct for the solvent flow you are using.

? Verify that the solvent is thoroughly degassed. Do not use ultrasonic degassing with protein samples.

? Verify that the LC backpressure is steady; steady backpressure indicates a steady solvent flow.

? If you are using water as part of the mobile phase, verify that it is deionized (>18MΩ).

Droplets, not spray, exiting the nebulizer

? Verify that the nebulizing gas pressure is set high enough for the LC flow being used.

? Check the position of the needle in the nebulizer.

? Stop the solvent flow and remove the nebulizer assembly. Use a magnifying glass to examine the end of the nebulizer for damage.

No flow

? Verify that the LC pump is on and there is sufficient solvent in the correct bottle.

? Verify that the solvent supply lines are free of air bubbles. Purge the appropriate pump channels, if needed.

? Check for LC error messages.

? Check for blockages. Repair or replace any blocked components.

? Check for leaks.

? Verify that the MS stream selector valve is set to LC to MSD.

Undesired fragmentation

? Fragmentor is set too high.

? Ionization is causing fragmentation (APCI vs. Electrospray).

? APCI temperature is too high.

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In This Book

This book contains information to get you started with the OpenLab ChemStation software for the InfinityLab LC/MSD Series and 6100 Series LC/MS.

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