

Agilent
7200 Accurate-Mass
Quadrupole
Time-of-Flight (Q-TOF)
GC/MS System

**Quick Start** 



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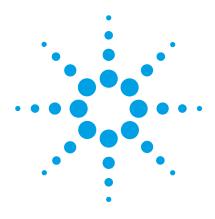
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This book provides a brief summary of your Agilent 7200 Accurate-Mass Quadrupole Time-of-Flight (Q-TOF) GC/MS system and each document written about it.





## **Overview**

#### **Hardware**

The **Agilent 7200 Accurate-Mass Q-TOF GC/MS** is a quadrupole time-of-flight mass spectrometer for use with the Agilent 7890 Gas Chromatograph. The 7200 Q-TOF features:

- Three turbomolecular vacuum pumps
- A rotary vane foreline pump
- A quick change independently heated removable ion source (RIS)
- A proprietary internal reference mass (IRM) delivery system for "on the fly" mass axis correction
- A high temperature heated hyperbolic quadrupole mass filter
- A single hexapole collision cell blanketed with nitrogen gas
- A constant temperature flight tube with dual-stage ion mirror
- Fast electronics, allowing fast sampling rates
- An analog-to-digital detector
- An independently heated GC/MS interface
- New ion optics for focusing the ions prior to the flight tube and detector

**Documentation** for the Agilent 7200 Accurate-Mass Q-TOF GC/MS system includes the Concepts Guide, Operation Manual, and the Troubleshooting and Maintenance Manual. Each of these are described later in this document.

## **Software**

Your Agilent 7200 Accurate-Mass Q-TOF GC/MS system comes configured with the powerful **MassHunter Workstation** software which includes:

• Data Acquisition

Unknowns Analysis

Qualitative Analysis

Library Editor

Quantitative Analysis

• Report Designer Add-in

Online Help, software demonstration videos, and familiarization guides are provided for each of these, and are described later in this document.

## **Online Help**



#### Create a new batch 🗐



A Batch is a group of related data files that are processed using the same data analysis method. It is saved as a .bin file.

The following describes how to create a batch and add samples to it so they may be analyzed by the MassHunter program.

To create a new batch:

- 1. Select File > New Batch, and navigate to, or create the directory where you want to store the batch and the results of your analysis (the batch directory).
- 2. In the File name field, type the name you wish to assign to this batch.
- 3. Press Open. The MassHunter program will:
  - a. Create a folder named QuantResults

#### Software demonstration videos

Your online **Help** system includes beginner to advanced software demonstration videos.

- The **Getting Started** videos are designed to give you the information you need to start using your system quickly and efficiently.
- The **Advanced** videos explain some of MassHunter's more intricate, but perhaps less frequently used functions.

## **Step-by-step instructions**

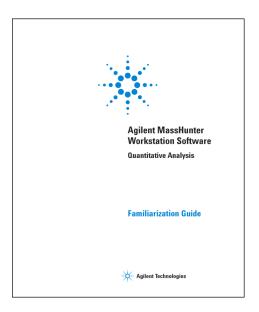
Most Help topics are written as detailed step-by-step instructions which walk you through routine daily tasks. Also included are links to other relevant material and video demonstrations.

# Camera icon 🗐



Any topic with the camera icon has a video linked to it. Watch these videos to see a demonstration of using the described feature in your MassHunter software.

## **Familiarization Guides**

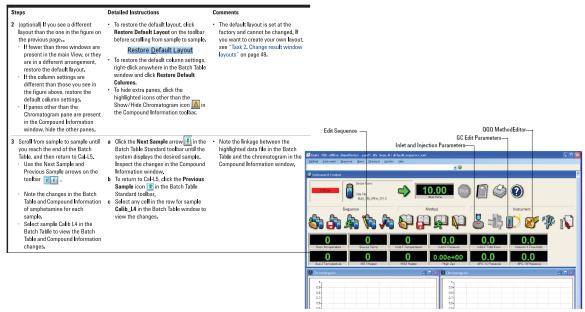


There are several familiarization guides available for your system, including one for:

- Data Acquisition
- Quantitative Analysis
- · Qualitative Analysis
- Unknowns Analysis
- Reporting

These guides provide detailed step-by-step instructions on how to prepare your system for data acquisition, how to create acquisition methods, qualitative methods, and quantitative methods, how to use the Unknowns Analysis software, and how to use the MassHunter reporting features.

The familiarization guides are provided as PDF files, accessible from online Help.

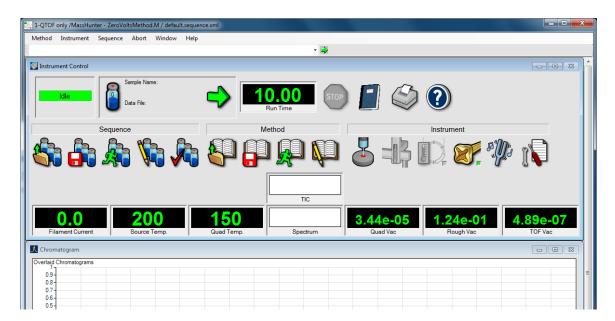


# **Data Acquisition Software**

From one screen in the **MassHunter Data Acquisition** software you can:

- Start and stop the instrument
- Download instrument settings in real time
- · Load, edit, create, or run a method
- Load, edit, create, check, or run a sequence
- Evaluate the MS parameters
- Autotune the MS parameters
- Monitor the real-time conditions of the instrument parameters
- View the instrument logs

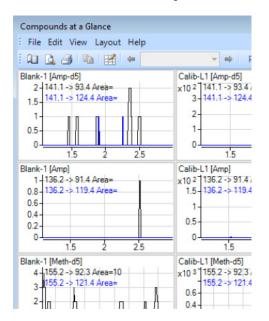
- View and print the real-time plot for chromatograms and instrument spectra
- Use the interactive video to help you quickly change the ion source
- Enter acquisition method parameters for the GC and the Q-TOF
- Select and label the total ion chromatograms (TIC) or extracted ion chromatograms (EIC) that you want to appear in the real-time plot
- Press **F1** to see context sensitive online Help



Online Help (F1) for the MassHunter Workstation Data Acquisition software includes:

Step-by-step detailed instructions Software demonstration videos Data Acquisition Familiarization Guide

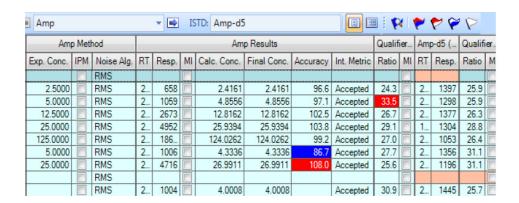
## **Quantitative Analysis Software**



The **MassHunter Quantitative Analysis** software will quantify very low amounts of sample.

From one screen in the MassHunter Quantitative Analysis software you can:

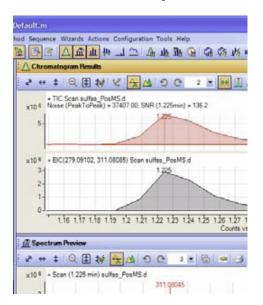
- Import information directly from the acquisition method
- Use the curve-fit assistant to test all fits and statistics on the calibration curve quality
- Integrate signals with an automated, parameter-free integrator that uses a novel algorithm, optimized for Q-TOF data
- Use the Compounds-at-a-glance feature to review all the chromatograms in your batch at once, side-by-side.
- Quickly scan the batch table and see your quality metrics such as high outliers (in red) or low outliers (in blue)
- Press **F1** to see context sensitive online Help



Online Help (F1) for the MassHunter Workstation Quantitative Analysis software includes:

Step-by-step detailed instructions Software demonstration videos Quantitative Analysis Familiarization Guide

# **Qualitative Analysis Software**

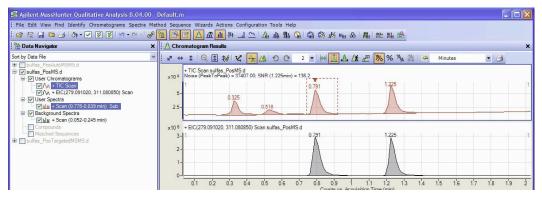


The MassHunter Qualitative Analysis software uses advanced data mining and processing tools, including integrated deconvolution algorithms, to rapidly and accurately find all detectable compounds in your samples and easily confirm targets or identify unknowns.

This software presents large amounts of data for review in one central location. From one screen you can:

- Extract chromatograms
- · View and extract peak spectra
- · Subtract background
- Integrate the chromatogram
- Find compounds and generate formulas
- Press F1 to see context sensitive online Help

You can also set up methods to automatically do the tasks in the above list, as well as others, when you open the data files.



Online Help (F1) for the MassHunter Workstation Qualitative Analysis software includes:

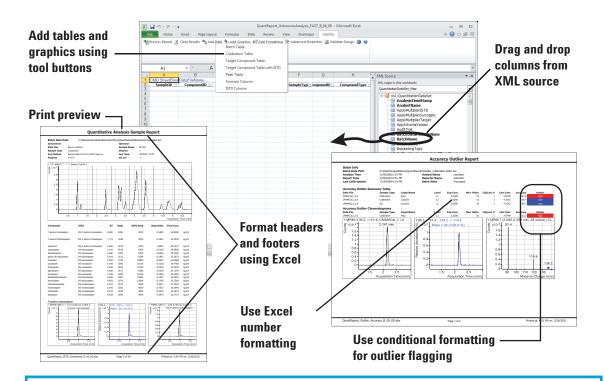
Step-by-step detailed instructions Qualitative Analysis Familiarization Guide

## Report Designer Add-in

MassHunter stores all results in XML format and uses Microsoft Excel and the **Report Designer Add-in** for reporting. Many application-specific templates and customized reports, including custom calculations for your specific application needs, are available in the familiar Excel environment.

Using these preconfigured templates, you can easily create reports in the **Data Acquisition** software, the **Qualitative Analysis** software, and the **Quantitative Analysis** software.

You can also customize many of the application-specific templates, including the custom calculations, with the **Report Designer Add-in**.



Online Help (F1) for the MassHunter Workstation Report Designer Add-in includes:

Step-by-step detailed instructions

Software demonstration and training videos and quizzes with instant feedback Reporting Familiarization Guide

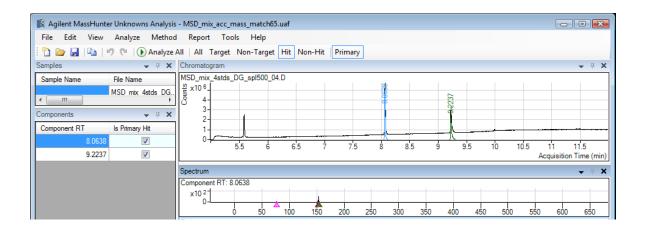
The Reporting User Information DVD also includes extensive software demonstrations and training videos.

## **Unknown Analysis Software**

The **MassHunter Unknowns Analysis** software identifies compounds that may be present in a sample batch beyond those that have been identified in the MassHunter Quantitative Analysis software.

From one screen in the MassHunter Unknowns Analysis you can:

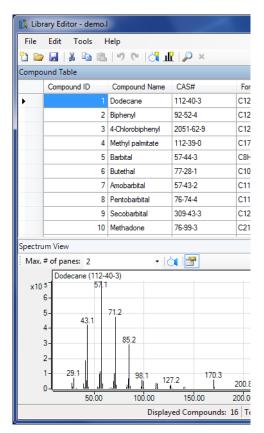
- Import data directly from the MassHunter Quantitative Analysis software.
- Use optimized deconvolution and library match algorithms to identify target and non-target hits
- View chromatograms, component lists, spectrums, ion peaks, and molecular structures side-by-side.
- Easily review library match and peak shape metrics of unknown components in table and spectrum formats.



Documentation for the MassHunter Workstation Unknowns Analysis software includes:

Step-by-step detailed instructions in online Help Unknowns Analysis Familiarization Guide in online Help Software demonstration videos on the Installation DVD





The **MassHunter Library Editor** software customizes libraries used to identify compounds and components in the MassHunter Qualitative and Quantitative Data Analysis software.

From one screen in the MassHunter Library Editor you can:

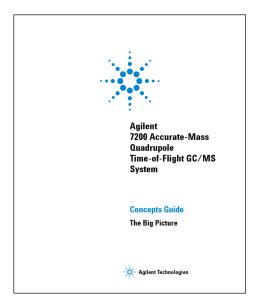
- · Add compounds to a library
- Copy spectral data from your data analysis software to a library
- Add a structure to your custom library
- · Import JCAMP files into a custom library
- Create a reference pattern library

Documentation for the MassHunter WorkStation Library Editor software includes:

Step-by-step detailed instructions in online Help

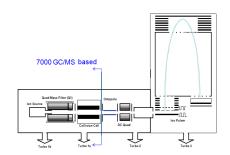
Software demonstration videos on the Installation DVD

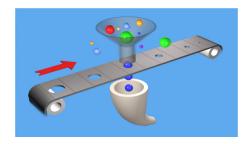
# **Concepts Guide**



To learn how your Agilent 7200 Accurate-Mass Q-TOF GC/MS system works in conjunction with the MassHunter software, take a look at the **Concepts Guide**. This document includes:

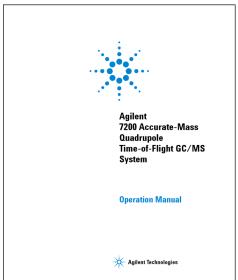
- · Conceptual models
- Illustrations
- Screen captures
- An overview of how the 7200 Q-TOF works
- Concepts behind the design of the Agilent MassHunter Workstation software
- Explanations on how the 7200 Q-TOF achieves high sensitivity
- Plus a lot more





The Concepts Guide is included on the GC and GC/MS Hardware User Information & Utilities DVD.

## **Operation Manual**



Your Agilent 7200 Accurate-Mass Q-TOF GC/MS system is extremely powerful and flexible.

Refer to the **Operation Manual** for extensive information on operating the instrument and performing routine maintenance on it. Included are **step-by-step instructions and videos** on installing GC columns, operating in electron impact (EI) mode, general maintenance procedures, and more.



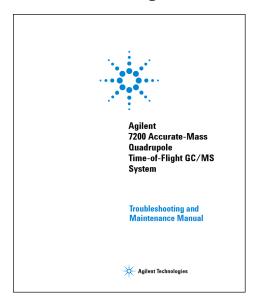
Any topic with the camera icon is linked to a video. Watch these videos to learn more.



The **Operation Manual is** included in the online Help.

A copy is also on the **GC and GC/MS Hardware User Information & Utilities** DVD.

## **Troubleshooting and Maintenance Manual**



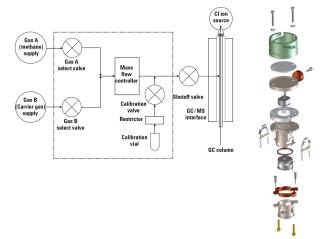
The Agilent 7200 Accurate-Mass Q-TOF GC/MS system consists of a 7890 Gas Chromatograph (GC) and a 7200 Accurate-Mass Q-TOF MS.

The **Troubleshooting and Maintenance Manual** for your system includes step-by-step instructions along with videos, IPBs, and annotated high-resolution photos to help you identify the cause of problems in your system. Also included are detailed procedures for completing common maintenance tasks, sections on the components of the GC/MS vacuum system and analyzer system, and a listing of parts.



Any topic with the camera icon is linked to a video. Watch these videos to learn more.





A copy of this manual is included on the GC and GC/MS Hardware User Information & Utilities DVD.

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