

# Agilent 6410 Triple Quad LC/MS

# **Concepts Guide**

# The Big Picture



# Notices

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## In This Guide...

The Concepts Guide presents "The Big Picture" behind the operation of the Agilent 6410 Triple Quad LC/MS by helping you understand how the hardware and software work.

### **1** Overview

Learn how the Agilent 6410 Triple Quad helps you do your job.

### 2 Inner Workings – Triple Quadrupole MS versus Single Quadrupole MS

Learn the concepts you need to understand how the Agilent triple quadrupole mass spectrometer works.

### **3** Agilent Triple Quad MS and Sensitivity

Learn how the Agilent triple quadrupole mass spectrometer achieves high sensitivity.

### 4 Agilent MassHunter Workstation Software

Learn concepts behind the design of the Agilent MassHunter Workstation Software, which consists of these components:

- Acquisition
- Quantitative Analysis
- Qualitative Analysis

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Agilent 6410 Triple Quad LC/MS Concepts Guide

# **Overview**

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This chapter provides an overview of the Agilent 6410 Triple Quad LC/MS components and how they help get the job done.



#### 1 Overview

What kind of system do you have?

### What kind of system do you have?

ESI – Electrospray Ionization APCI – Atmospheric Pressure Chemical Ionization APPI - Atmospheric Pressure Photo Ionization HPLC-Chip/MS – High Performance Liquid Chromatography on a Chip MMI - Multimode Ionization You can set up an Agilent 6410 Triple Quad LC/MS in several configurations:

- For normal flow LC/MS with a binary pump, quaternary pump, well-plate sampler (or autosampler or CTC PAL autosampler). The supported ion sources are ESI, APCI, APPI, and MMI.
- For microflow LC/MS with a capillary pump, micro well-plate sampler (or CTC PAL micro-plate autosampler) and ESI, APCI or MMI ion sources
- For nanoflow LC/MS with a nanopump, micro well-plate sampler and HPLC-Chip/MS interface (used in place of standard or dual nanospray source) to increase reliability and boost performance with narrow peak dispersion and lower dead volumes.

Each Agilent combination has advantages for different applications. Each uses the same Agilent MassHunter Workstation Acquisition program and Agilent MassHunter Workstation Quantitative and Qualitative Analysis programs to enable these advantages.

### **Help for applications**

You can use one or more of the Agilent 6410 Triple Quad LC/MS combinations to quantitate trace organic compounds in complex matrices:

- Food safety studies
- Environmental studies
- Drug discovery
- Toxicology
- Forensics

Paired with Agilent's 1200 Series LCs, the 6410 Triple Quad MS delivers sensitive, reproducible analyses of target compounds in complex matrices.

• Femtogram-level limits of detection and quantitation

The *dwell time* is the amount of time allotted for analyzing each ion during a scan.

- Minimized memory effects even at very short dwell times
- Simplified operation with Agilent's newly developed data analysis software

# Help for acquisition

To help you use the Agilent Triple Quad LC/MS for these applications, the software lets you perform the following tasks in a single window with the Agilent MassHunter Workstation Acquisition Software:

### Prepare the instrument

- Start and stop the instruments from the software
- Download settings to the Agilent 1100 LC or 1200 LC and the Agilent Triple Quad mass spectrometer in real time to control the instrument
- Evaluate if the MS parameters are within the limits to produce the specified mass accuracy and resolution with a Checktune report
- Optimize MS parameters automatically (Autotune) or manually through Agilent tuning programs and print an Autotune report
- Monitor the actual conditions of the instrument
- View the real-time plot for chromatograms and instrument parameters (both UV/Vis and MS) and print a real-time plot report
- View the centroided line spectrum of a peak or the mass range profile spectrum of a peak in real time

### Set up acquisition methods

- Enter and save parameter values for all LC modules and the MS to an acquisition method
- Select and label the total ion chromatograms or extracted ion chromatograms that you want to appear in the real-time plot
- Set up time segments for each scan type and analysis where parameters change with the time segment or with the scans within the time segment
- Print an acquisition method report

To learn how to get started with the Agilent Triple Quad LC/MS, see the Quick Start Guide.

To learn more about how to use the Agilent Triple Quad LC/MS with real samples and data, see the Familiarization Guide

To learn how to perform individual tasks with the LC/MS, see the Online Help.

*To learn more about an Agilent 1100LC module or 1200LC module, see the* Agilent 1100LC or 1200LC User's Guide *for the module.* 

### Acquire data

• Enter sample information and pre- or post-analysis programs (scripts) and run single samples interactively

• Enter and automatically run both individual samples and samples organized in a *worklist* (sequence of samples)

- Set up pre- and post-analysis scripts to run between samples in a worklist
- Set up and run a worklist to optimize MS acquisition parameters
- Print a worklist report
- View system events, including start and stop times, run events and errors and print an event log report

### Help for data analysis

#### Agilent MassHunter Workstation Software Quantitative Analysis

Agilent has designed the quantitative analysis software to help quantitate very low amounts of material with the following unique features:

- Imports information directly from the acquisition method
- Provides a curve-fit assistant to test all fits and statistics on curve quality
- Integrates with an automated, parameter-free integrator that uses a novel algorithm, optimized for triple quad data
- Presents a Batch-at-a-Glance results window to help you review and operate on an entire batch of data at once
- Automatically detects outliers
- Provides preconfigured templates for basic reporting and enables the capability to create custom reports in Microsoft Excel

A worklist is a list of individual samples and batches (sequences) that you enter and run automatically with the Agilent MassHunter Workstation Acquisition program.

#### 1 Overview

Help for data analysis

### Agilent MassHunter Workstation Software Qualitative Analysis

For fast method development, this software is used to quickly review the qualitative aspects of the data, such as the optimum precursor to product ion transitions.

Agilent designed the Qualitative Analysis program to present large amounts of data for review in one central location. With the program you can perform these operations for any type of mass spectrometer data that you open:

- Extract chromatograms
- View and extract peak spectra
- Subtract background
- Integrate the chromatogram

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



Agilent 6410 Triple Quad LC/MS Concepts Guide

# 2 Inner Workings – Triple Quadrupole MS versus Single Quadrupole MS

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In this chapter you learn about concepts to help you understand the inner workings of the Agilent Triple Quad LC/MS.

The foundation for understanding the operation of a triple quadrupole mass spectrometer is the operation of a single quadrupole mass spectrometer. Therefore, an explanation of the workings of a single quadrupole mass spectrometer is presented first.



2 Inner Workings – Triple Quadrupole MS versus Single Quadrupole MS Single quadrupole MS operation

# Single quadrupole MS operation

To better understand the specific hardware features of the Agilent Triple Quadrupole Mass Spectrometer, this section first reviews the fundamental aspects of the single quadrupole mass spectrometer.

### Design for a single quadrupole mass spectrometer

Mass spectrometry is based on the analysis of ions moving through a vacuum.

The ionization of a sample occurs in the ion source that is shown, schematically, on the left. The ions are analyzed by a mass analyzer (mass filter) that controls the motion of the ions as they travel to the detector to be converted into actual signals.



Figure 1 Schematic for single quadrupole mass spectrometer

m/z - mass/charge ratio The quadrupole mass analyzer consists of four parallel rods to which specific DC and RF voltages are applied. These rods filter out all ions except those of one or more particular m/z values as determined by the voltages applied.

The RF is applied to all four rods, but the negative (–) rods are 180 degrees out of phase with the positive (+) rods. The rods are labeled + and – in reference to the DC voltages applied to them.

All ions that comprise the sample are generated at the source. However, when a specific set of voltages is applied, only ions of the corresponding m/z value may pass through the quadrupole to reach the detector. As the voltages are increased to other values, ions with other m/z values are allowed to pass through. A full MS scan is obtained by increasing the DC and RF voltages applied to the four rods over an expanded range of values.

### How a single quadrupole mass spectrometer works

A diagrammatic model can be used to illustrate the concept of how a single quadrupole instrument works. See Figure 2.





In the model,

• All of the ions contained in a sample are formed in the external ionization source and collected in a funnel. The balls of different colors and sizes represent different ions having different m/z values.

### 2 Inner Workings – Triple Quadrupole MS versus Single Quadrupole MS

How a single quadrupole mass spectrometer works

		• The quadrupole mass analyzer is represented by a moving belt that serves to filter the ions as they pass through openings of various sizes. The ions pass from the funnel, through the filter, to the detector.
		• The detector is represented by the collecting funnel below the filtering belt.
		As the belt (the analyzer) moves, or the voltages on the rods are changed, ions with different $m/z$ values are filtered through the mass spectrometer.
		As the analyzer moves from a small $m/z$ value to increasingly larger values, a full MS scan is created.
	SIM – Selected Ion Monitoring	If the belt does not move, the detector continues to monitor the same single $m/z$ value over the entire scan period. This type of analysis is known as SIM. It is the most sensitive operating mode for a single quadrupole mass spectrometer.
		The scan period is selected (fixed) by the user. The user may set the dwell time to scan a specific mass range (e.g. $m/z$ 50 to 1000) <i>or</i> to remain on one selected ion (SIM) <i>or</i> to move to several selected ions during the scan period. The quadrupole mass filter is not scanned in this mode. The required RF/DC voltages are often set to filter a single mass at one time.
		For comparison, see "How a triple quadrupole mass spectrometer works" on page 22.
		Single quadrupole: SIM
		To obtain the best consitivity or quantitation, the single

To obtain the best sensitivity or quantitation, the single quadrupole is operated in SIM mode (Figure 3). The duty cycle is the measure of the instrument's time actually devoted to measuring signals. In SIM mode, the single quadrupole analyzes the signal of a specific m/z ion almost all of the time. This results in nearly 100% acquisition during the duty cycle.

How a single quadrupole mass spectrometer works



**Figure 3** Single quadrupole: SIM

In this example,

- 1 All of the ions (+, -, and neutrals) are formed in the API source.
- 2 Ion optics guide the ions to the quadrupole mass analyzer.

The Agilent Ion Guide is an octopole filter of eight equally spaced rods.

- **3** In the analyzer, only ions of a particular m/z value, represented by blue balls, are allowed to pass through to the detector.
- **4** The detector completes the analysis.

This system has several advantages:

- provides the best sensitivity for quantitation
- increases selectivity
- improves chromatographic specificity
- provides no structural information

#### 2 Inner Workings – Triple Quadrupole MS versus Single Quadrupole MS How a single quadrupole mass spectrometer works

#### Single quadrupole: Full Scan MS

In a full MS scan, the quadrupole serves as a mass filter over time, and a scan is carried out by stepping through increasing DC and RF voltages. This provides filtering through the corresponding m/z values across a mass spectrum. See Figure 4.



Figure 4 Single quadrupole: Full scan MS

The full scan MS mode is less sensitive because the duty cycle for each m/z is considerably less than 100%. The quadrupole mass analyzer scans sequentially, passing each m/z in the selected mass range to the detector.

A full scan MS is still a useful mode of operation because it shows all of the ions that are being formed in the ion source. This is useful for developing SIM acquisitions but also alerts analysts to other compounds co-eluting with compounds of interest.

### What about fragment ions?

Full scans with a single quadrupole instrument can also be used to study fragment ions. See Figure 5.



Figure 5 Fragment ions with single quadrupole MS

The diagram shows that fragment ions, also known as product ions, are formed by fragmenting or breaking apart precursor ions. Precursor ions formed in the ion source travel through the mass analyzer without change, unless extra energy is applied to their motion in a region where fragmentation can occur.

This fragmentation or collisionally induced dissociation (CID) can be carried out in a low pressure region between the ion source and the mass analyzer. The ion source is under atmospheric pressure, while the mass analyzer is at a much lower pressure because it has been evacuated of gas with a vacuum pump.

On the Agilent single quadrupole mass spectrometer, this region is between the capillary exit and the skimmer, where the gas pressure is about 2 Torr, or about three orders of magnitude below atmosphere pressure (760 Torr). Under normal operation, a voltage is applied across this region to keep the

### $\label{eq:constraint} 2 \qquad {\rm Inner \ Workings-Triple \ Quadrupole \ MS \ versus \ Single \ Quadrupole \ MS}$

How a single quadrupole mass spectrometer works

ions passing through to continue on to the mass analyzer. Even if these ions collide with the gas molecules in this region, they usually do not have enough energy to fragment.

#### CID – Collisionally Induced Dissociation

However, as the voltage is increased, the ions have more translational energy. Then, if the ions run into gas molecules, the collisions convert the translational energy into molecular vibrations that can cause the ions to fragment (Figure 6). This is collisionally induced dissociation (CID). Even though this fragmentation does not occur where the ions are formed at atmospheric pressure, it's a tradition to call this type of fragmentation "In-source CID."



**Figure 6** Ion fragmentation caused by collision-induced dissociation

A single quadrupole mass spectrometer cannot be used to perform MS/MS because all of the ions formed in the ion source are transferred to the quadrupole whether fragmented or not. At the end when the mass analyzer filters the ions, it is not possible to identify which product ions came from which precursor ions.

A triple quadrupole mass spectrometer can perform MS/MS, with fragmentation within its collision cell as described in the next section.

# **Triple quadrupole MS operation**

Be sure to read the previous section on the concepts behind the operation of a single quadrupole mass spectrometer. Understanding these concepts helps you understand the operation of the triple quadrupole mass spectrometer.

## **Design of the Agilent Triple Quad MS**

The triple quadrupole mass spectrometer consists of an ion source, followed by ion optics that transfer the ions to the first quadrupole positioned to the right of it. A diagram of the spectrometer is shown in Figure 7.



Figure 7 The Agilent Triple Quad MS

The quadrupole consists of four parallel hyperbolic rods through which selected ions are filtered before reaching a collision cell where they are fragmented. The collision cell is typically called the second quadrupole, but in this case, geometrically it is actually a hexapole filled with nitrogen, the same gas used in the ion source. The fragment ions formed in the collision cell are then sent to the third quadrupole for a second filtering stage to enable a user to isolate and examine one precursor and one product ion.

### How a triple quadrupole mass spectrometer works

Quadrupoles provide a user with the capability to perform MS/MS in several ways (see Figure 8).



**Figure 8** Conceptual model of a triple quadrupole mass spectrometer: With SIM, MS1 (Q1) and MS2 (Q3) are set at a single *m/z*, while with Full Scan MS/MS, MS1 (Q1) is set at a single *m/z* and MS2 (Q3) is scanned.

Representing the quadrupole mass analyzers as moving belts, a collision cell can be placed between the belts to fragment the ions. The first belt can be fixed to select which precursor ion travels to the collision cell. Different types of collision cells can be used. The cell can be another quadrupole, a hexapole (six rods like the one used in the Agilent Triple Quad LC/MS), an octopole (eight rods), or even a transverse wave guide.

Whichever geometry is used, a collision gas is required—an inert, non-reactive gas such as nitrogen or argon. Nitrogen is used here. In addition, the voltages applied to the collision cell must be different from those applied to the quadrupoles to enhance the movement of all of the product ions toward the third quadrupole.

In this example, a precursor ion is selected using the first quadrupole and is sent to the collision cell for fragmentation. The fragments are scanned through the third quadrupole resulting in a product-ion scan MS/MS. Since the fragment ions are pieces of the precursor, they represent portions of the overall structure of the precursor molecule. A triple quadrupole instrument can be used in this way to identify a compound's fingerprint.

A full scan MS/MS using a triple quad MS is also not the most sensitive mode for the same reason that a full scan MS using a single quadrupole is not the most sensitive mode of operation possible (first belt remains steady; second belt moves). The most sensitive mode of operation for the triple quad MS instrument is to fix both belts and only monitor a specific precursor ion and a specific product ion. This mode is called *selected reaction monitoring or SRM*.

In normal operation, a triple quad MS instrument involves running multiple SRMs for the same precursor ions. This is called *multiple reaction monitoring or MRM*.

### 2 Inner Workings – Triple Quadrupole MS versus Single Quadrupole MS

How a triple quadrupole mass spectrometer works



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3

# **Agilent Triple Quad MS and Sensitivity**

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This chapter shows how the Agilent triple quadrupole mass spectrometer reduces chemical and electronic noise and how each component contributes to enhanced instrument sensitivity.



# How the Agilent Triple Quad MS improves sensitivity

Triple quadrupole mass spectrometers exhibit multiple sources of noise, including noise from all chemical and cluster backgrounds and electronic noise (Figure 9).

## **Noise reduction**

The problem of noise must be addressed at several stages of the instrumentation from the ion source (1) to the detector (10).



Figure 9 Multiple sources of noise

### How the Agilent Triple Quad instrument minimizes noise

- **1** Agilent's patented orthogonal spray sources maximize ionization while minimizing solvent and matrix noise.
- **2** This combination of a heated counter-current drying gas, dielectric capillary and skimmer enhances desolvation while minimizing chemical noise.
- **3** RF Octopole ion guide provides high efficiency ion capture while optimizing wide mass bandwidth ion transmission.

- 4 L2 RF enhances high mass ion transmission (Agilent patent).
- **5** Quadrupole 1 uses hyperbolic quadrupoles to optimize ion transmission and spectral resolution.
- **6** RF quadrupole segment enhances ion transmission into and out of the collision cell.

7 High pressure collision cell with linear acceleration optimizes MS/MS fragmentation while eliminating crosstalk, even at very low dwell times. A small diameter high frequency hexapole assembly assists with capturing and focusing fragmented ions.

- **8** Quadrupole 2 uses hyperbolic quadrupoles to optimize ion transmission and spectral resolution.
- **9** The off-axis matching dual high energy dynode detector with log amp signal compression permits a high gain with rapid polarity switching, a long life and low noise. The off-axis design allows neutrals to pass without hitting the detector.
- **10** The multiplier has a long life since only electrons impact its surface, never ions.

# **Example of chemical noise reduction**

The Agilent Triple Quad mass spectrometer passes through four transitional steps in translating a signal in the MRM process (Figure 10).

Crosstalk is the interference caused when two signals become partially superimposed on each other. In this case residual product ions can interfere with the product ion spectrum of a subsequent MRM experiment. 3 Agilent Triple Quad MS and Sensitivity

**Example of chemical noise reduction** 



Figure 10 Multiple reaction monitoring (MRM)

**Step 1** The spectrum at the far left represents everything that is being ionized at the ion source. This example shows the ESI spectrum of a phenylurea pesticide. A triple quad mass spectrometer reduces chemical noise for low-level quantitation in a dirty matrix more than a single quadrupole LC/MS does.

**Step 2** This step is accomplished by first selecting the pesticide of interest at m/z 210 from the co-eluting interferences seen in the rest of the spectrum. The second spectrum shows the result after passing through the first quadrupole, or MS1 (Q1).

**Step 3** After MS1 (Q1), fragment ions are generated in the collision cell. The corresponding MS/MS spectrum is shown below the collision cell.

**Step 4** Particular fragment ions can be selected to pass through the MS2 (Q3) quadrupole. These are selected for quantitation and confirmation. For example, the product ion at m/z 158 is more intense than the product ion at m/z 191. Therefore, the MRM transition 210 to 158 would be used for quantitation and the 210 to 191 transition would be used for confirmation, where the m/z 191 ion is considered a qualifier ion.

The second stage of selectivity using the MS2 (Q3) quadrupole removes much of the chemical background. Typically, the chance of an isobaric interference at the same exact mass as the fragmentation ion is remote.

# Sensitivity and linearity of the Agilent Triple Quad MS

### Sensitivity based on reserpine

The reserpine sensitivity specification for the Agilent Triple Quad MS is

- 500 fg reserpine on-column, with
- a signal-to-noise ratio of 20:1 or greater

Three concurrent MRM transitions are monitored, but only the 609.3 to 195.1 transition is used for the quantitation. Reserpine's precursor ion mass is 609.3.

The other two transitions are used to simulate the acquisition of qualifier ions, which represent real-world applications for confirmation. The instrument can detect femtogram levels of material (Figure 11).

### 3 Agilent Triple Quad MS and Sensitivity

Sensitivity and linearity of the Agilent Triple Quad MS



Figure 11 Reserpine Sensitivity (Specifications: 500 fg with S/N 20:1)

**NOTE** Currently, the sensitivity for reserpine is based on the ESI ion source.

### Linearity

Reserpine can also be analyzed with repeated injections of the sample, showing a linearity of  $r^2 = 0.987$  (Figure 12).



**Figure 12** Demonstrating linearity in a Reserpine dilution series, in mobile phase (Reserpine, 609>195 EID with three transitions

#### **3** Agilent Triple Quad MS and Sensitivity

Sensitivity and linearity of the Agilent Triple Quad MS





Figure 13 Alprazolam results (LP5) with 1 pg–10 ng positive ESI mode

# How each component works to improve sensitivity

This section describes in more detail how each of the components of the Agilent Triple Quad MS contributes to reducing noise (Figure 9).

## LC/MS ion sources

Agilent provides a choice of four ion sources to use with its triple quadrupole mass spectrometer: ESI, APCI, APPI and MMI. You can also use the HPLC-Chip.

This section describes how the different ion sources affect sensitivity.

### **ESI** ion source design

The patented orthogonal source reduces the introduction of unwanted sample components that interfere with analysis. The advanced nebulizer design produces a uniform droplet size, which ensures maximum sensitivity. Since the source is at ground, the source is safe to use and has the advantage of reducing solvent cluster background (Figure 14).



Figure 14 Orthogonal introduction and electrospray ionization

### 3 Agilent Triple Quad MS and Sensitivity

LC/MS ion sources

The capillary is glass, dielectric "cold" capillary that enhances desolvation and improves the focusing of high mass ions while minimizing chemical noise. Some desolvation occurs in the capillary. This appears to work better with glass capillaries rather than stainless steel capillaries and reduces cluster noise. Ion focusing is also improved.

#### **Atmospheric Pressure Chemical Ionization (APCI)**

Atmospheric Pressure Chemical Ionization (APCI) is a popular complement to electrospray. Because APCI does not generate multiply charged ions, and operates at higher temperatures, it is commonly used to analyze smaller, thermally stable polar and non-polar compounds. Agilent's APCI source is sensitive, yet extremely robust thanks to orthogonal spray and counterflow drying gas. Like the ESI source, it can generate both positive and negative ions, and ion polarity can be switched on a spectrum-to-spectrum basis.

#### Agilent multimode source

The most versatile ion source for the Agilent Triple Quad MS is Agilent's G1978A Multimode Source shown diagrammatically in Figure 15.





Neutral analytes and ESI charged analytes pass through the divided chamber entering either the APCI Zone or adjacent zone.

**NOTE** Analytes are distributed spatially between the two sections.

ESI and APCI are essentially incompatible processes because each needs its own conditions for aerosol drying and electrical fields. However, it is possible to form ions simultaneously from ESI and APCI if the two ionization regions are separated in space.

The HPLC effluent is nebulized using the same sprayer that is used for a dedicated ESI source. The droplets are emitted into the "ESI zone," where a high voltage electrode charges the droplets and induces ion formation. The ions formed in this region pass through the source and enter the capillary. Residual droplets are dried using two infrared lamps (not shown) that emit at the absorption frequency of water. The vapor and analyte(s) enter the APCI zone where they are ionized. Ions are then drawn into the capillary the same way as they would be with dedicated ESI and APCI sources.

**Expected** When the Multimode source is operated as an ESI or APCI source only, no loss in sensitivity is found for the compounds studied. For many compounds run in mixed mode, an equal signal response (compared to single mode operation) or sensitivity gain can be achieved.

However, studies also show that when operating the Multimode source in ESI and APCI simultaneously, there can be a loss of sensitivity up to a factor of two for some compounds.

Therefore, weighing the benefits of running analyses in both modes simultaneously versus a potential loss in sensitivity is important. For most applications, a loss in sensitivity of less than 2 is negligible. 3 Agilent Triple Quad MS and Sensitivity LC/MS ion sources

LC/IVIS ION SOURCES

#### **APPI (Atmospheric Pressure Photo Ionization)**

For analysis of compounds that ionize poorly by ESI and APCI, the atmospheric pressure photoionization (APPI) source provides a useful alternative. It combines Agilent's proven orthogonal spray nebulization and counterflow drying gas with innovative photoionization from Syagen Technology. The long-lasting krypton lamp emits photons at energy levels high enough to ionize many large classes of compounds, but low enough to minimize the ionization of air and common HPLC solvents. Relatively low ionization energy means the APPI source causes minimal fragmentation and generates mostly molecular ions and protonated molecules.

APPI may ionize compounds that do not ionize well by ESI or APCI, such as Polyaromatic Hydrocarbons (PAHs). Also, APPI may provide better overall sensitivity for some compounds than either ESI or APCI. Some examples of these compounds are Tetrahydrocannabinol (THC), Benzoic acid, and fat soluble vitamins. APPI may provide better sensitivity at lower flow rates than APCI. Reproducibility data indicates APPI is robust and highly reproducible.


## **HPLC-Chip**

Traditional nanospray mass spectrometry has proven its potential as a cost-effective, sensitive and reproducible technique for the identification of peptides at femtomol to atomol levels. However, connecting nano capillaries, columns and valves frequently is a tedious procedure and requires user skills and routine. When handled incorrectly, nano flow connections are prone to leakage which are difficult to detect and result in poor system performance and extended downtime of the complete system. Quartz nanospray needles are prone to blockages and require frequent replacement.

With the invention of HPLC-Chip technology, Agilent is significantly reducing the need for user interaction and making nanospray a rugged state-of-the-art technology.

The cornerstone of chip technology is the HPLC-Chip, a 3-dimensional structure made of sandwiched polyimide films. Grooves of specific dimensions are laser-ablated into a layer of polyimide film. The lamination of a top and bottom layer then forms channels of trapezoidal or triangular shape inside the chip which can either be used as capillaries or, if filled with packing material, as nanocolumns. One end of the chip tapers off into a polyimide nanospray emitter.



By ablating small holes into the top and bottom layers, the chip can be used as a self-connecting and sealing microvalve. The unique rotor-in-rotor microvalve in the Chip Cube moves two

#### 3 Agilent Triple Quad MS and Sensitivity

LC/MS ion sources

concentric valve rotors from one side towards the chip while a 6-port stainless steel stator face closes in from the other side, forming a pressure tight connection without any user interaction.

The very flexible HPLC-Chip concept can combine multiple columns, the valve plumbing and the nanospray emitter on one chip. A rugged chip holder protects the chip and retracts the fragile tip whenever the HPLC-Chip is removed from the Chip Cube.

# **Front-end ion optics**

Several innovations enhance sensitivity across a broad mass range. Improved ion optics is one of those innovations and provides a 10X advantage. The key components are shown in Figure 16.





Lens 2 RF (transmission of higher masses)

Cuadrupole

Quadrupole and hyperbolic post-filter

Figure 16 Front-end optics

#### 3 Agilent Triple Quad MS and Sensitivity Collision cell

Skimmer	Agilent uses a small diameter skimmer orifice with very short distances from the capillary to the skimmer to the octopole. Consequently, more of the ions exiting the capillary are captured by the skimmer, thereby keeping the ion beam very tightly focused.
Octopole 1	Higher multipoles provide better ion transmission over a wider mass range. The depth of the potential well is steeper for the higher multipoles (at like voltages), especially those close to the rods resulting in the loss of fewer ions. This explains the predominant use of octopoles as ion guides in mass spectrometers where the main objective is to transmit rather than filter the ions.
Lens 2 RF	The phase of lens 2 is matched to that of the subsequent quadrupole, MS1 (Q1), resulting in a significantly increased sensitivity. This is an Agilent patented technology.
Quad mass filters	The quadrupoles consist of hyperbolic rods that optimize ion transmission and spectral resolution. There tends to be more ion loss with circular rods.
Pre-filter	The end section of the MS1 (Q1) quadrupole also consists of short hyperbolic rods, but their RF voltages are only high enough to guide ions into the collision cell. A similar set of rods on the exit side of the collision cell are part of the MS2 (Q3) quadrupole. These short RF-only rods act as pre- and post-filters to the collision cell to ensure optimum ion transmission.

# **Collision cell**

## What is the collision cell?

The collision cell is another innovation. The collision cell is a high pressure hexapole assembly with its linear acceleration adjusted to optimize MS/MS fragmentation while eliminating crosstalk even at very low dwell times (Figure 17).



Figure 17 Collision cell technology produces higher sensitivity and faster responses without memory or cross-talk effects

The components that contribute to this higher sensitivity and faster response are

- Small diameter hexapole collision cell
- High frequency hexapole collision cell
- Linear axial acceleration
- High pressure collision cell
- High speed digital electronics

The collision cell contains nitrogen, the same gas that is used in the ion source. The small diameter of the hexapole assembly assists in capturing fragmented ions.

#### 3 Agilent Triple Quad MS and Sensitivity Collision cell

#### Why a hexapole?

The geometry of a hexapole provides advantages in two domains: ion focusing and ion transmission (Figure 18).

- The first advantage is in ion focusing where a quadrupole is better than a hexapole, which is better than an octopole, that is, quadrupole > hexapole > octopole.
- The second advantage involves ion transmission across a wide mass range, or m/z bandwidth. In this case, the octopole is better than the hexapole, which is better than the quadrupole.

The hexapole is chosen because, overall, it is the best for both ion focusing and ion transmission.



## Mass Range Transmission (octo > hex > quad)

Figure 18 Broad mass range transmission and improved transmission efficiency using a hexapole

#### Collision cell design

The collision cell hexapole consists of six resistively coated rods used to generate a potential difference across the length of the collision cell (Figure 19).



Figure 19 Collision cell design

A potential difference is always present. This ensures that the precursor ions coming from MS1 (Q1), or fragment ions generated in the collision cell, are transmitted and not allowed to drift around at random.

Sweeping out the ions in this manner avoids the issue of crosstalk where residual product ions from a previous MRM experiment can interfere with the product ion spectrum of a subsequent MRM experiment (see Figure 20). A collision energy voltage is applied over the accelerating linear voltage to generate fragments or product ions.

## Length of time for collision cell flushing

The low degree of crosstalk can be demonstrated by examining how long it takes to evacuate ions from the collision cell (Figure 20). 3 **Agilent Triple Quad MS and Sensitivity** 



Figure 20 Collision cell clearing profile (500 pg Alprazolam, 20 ms dwell time)

The figure shows that the higher the mass the longer it takes to evacuate the collision cell. For example, m/z 922 takes about 600 µsec to evacuate the collision cell using the linear potential, while m/z 118 only takes 350 µsec. This also demonstrates the low degree of crosstalk since the Y axis is logarithmic, showing complete clearance of the cell. This means that an inter-scan delay of 5 msec will be more than adequate to flush the collision cell of all ions.

# Detector

The High Energy Dynode detector assembly is unique to Agilent (Figure 21).





The two dynodes are orthogonal to the ion beam and neutrals. This orientation reduces the possibility of neutral molecules impacting the detector while at the same time attracting the ions with high voltages. The dynodes serve to convert the ions to electrons before they impact the multiplier. The off-axis design also allows neutrals to pass through without hitting the detector.

The multiplier has a long lifetime since only electrons are allowed to impact it. Ions never impact its surface. 3 Agilent Triple Quad MS and Sensitivity Pumping system

# **Pumping system**

A single three-stage turbo pump is used for the entire system. This is achieved by partitioning the turbo to create the multiple vacuum stages. It removes the need for two turbo pumps. This pump is backed by a single mechanical (rough) pump (Figure 22).



Figure 22 Pumping system



Agilent 6410 Triple Quad LC/MS Concepts Guide

4

# Agilent MassHunter Workstation Software

Overview 48 Agilent MassHunter Workstation Software: Acquisition 51 Tuning 52 Acquisition 53 Agilent MassHunter Workstation Software: Quantitative Analysis 55 Quantitative methods 62 Parameter-free integrator 63 Batch-at-a-Glance: Results 65 Compound calibration 67 Compound confirmation 66 Reporting 71 Agilent MassHunter Workstation Software: Qualitative Analysis 73 Overview 73 Capabilities 75

Learn the concepts to help you understand the design and operation of the Agilent MassHunter Workstation Software. This software comprises three parts:

- Acquisition
- Quantitative Analysis
- Qualitative Analysis



# **Overview**

The Agilent MassHunter Workstation Software consists of three major components:

- Acquisition (Figure 23)
  - All LC and MS parameters are immediately visible.
  - Real-time plots show the instrument at work.
  - Running multiple samples is easily handled through a worklist—a spreadsheet-like interface.



Figure 23 Acquisition component

• Quantitative Analysis (Figure 24)

The Quantitative Analysis program provides Batch-at-a-Glance views that let you see quantitation results immediately for the entire batch of data that you choose to open. In addition, Batch-at-a-Glance detects and identifies outliers. This capability enables you to zero in on samples and compounds that may need reanalysis.



Figure 24 Quantitative Analysis component

#### • Qualitative Analysis (Figure 25)

The Qualitative Analysis program allows you to operate on many data files at once from a single screen. Typically, qualitative analysis is used to determine the precursor and product ions associated with a particular compound. This MRM transition information can be used to create the acquisition method and, later, the quantitation method.



Figure 25 Qualitative Analysis component

# **Agilent MassHunter Workstation Software: Acquisition**

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When you first open the Agilent MassHunter Workstation Data Software, you see the following window.

Figure 26 Agilent MassHunter Workstation Acquisition window

With this window you can perform these operations:

- Control and monitor instrument settings
- Tune the instrument
- Set up acquisition parameters for LC and MS
- Monitor the chromatogram and mass spectra as samples are analyzed
- · Set up worklists for sequences of samples

# Tuning

## Autotune and Checktune

A Checktune can be used to determine if the tuning mix ion masses are properly assigned and if the response or sensitivity of these ions is within expectations. This check tune will only take a few minutes. An Autotune taking 18-25 minutes for each polarity can be used if a more extensive tune is recommended (Figure 27).

Everything is automatic since the tuning mix is delivered by the calibrant delivery system (CDS), which is switched on automatically during the tune.

## Manual Tune

A manual tune of user-defined ion masses with six corresponding profile masses is available.



Figure 27 Autotune in progress

A Lens 2 DC voltage ramp with RF Off is shown at the top right. An automated version of this is also available for the Tuning Mix with Autotune.

## **Tune Report**

Tune reports are also available in Microsoft Excel. See "Reporting" on page 71.

# Acquisition

Many Agilent LC modules and the triple quad MS can be controlled and monitored (Instrument Status) from the same screen used for entering acquisition settings and setting up sequence lists (Figure 23).

## 4 Agilent MassHunter Workstation Software Acquisition

The Data Display also can show the MS and UV results in real time.

Because of the large amount of information available, any of these windows can be closed for easier viewing, if necessary.



Figure 28 Instrument status and real-time plot

# Agilent MassHunter Workstation Software: Quantitative Analysis

See the Agilent Triple Quad LC/MS Quick Start Guide for a quick look at how you can use these functions to quantitate data and evaluate results Quantitative Analysis includes seven capabilities that help you integrate, quantitate and review your data more easily and powerfully:

## Batch-at-a-Glance: Batch Table Setup

- New batch Creates a Batch Table in which you can operate on samples and compounds from a single view
- Analyze Recreates calibration curve and requantitates all samples using the method that is currently open
- Quantitate Applies the existing calibration curve to current batch, sample, or compound

The granularity of applying quantitation allows you to quickly manipulate a particular signal.

• Integrate – Integrates signals to the current batch, sample or compound

# **Method Editor**

- MRM Setup Presents a quantitation method in simple stepwise fashion
- Create method from acquired MRM data Creates a quantitation method automatically from the acquisition method after requiring only the assignment of ISTD relationship and concentrations
- Group by time segment Organizes methods by compounds in ordered time segments
- Validate Ensures that a quantitation method meets rigorous criteria
- Isotopic dilution Supports adjustments from (Rx, Ry) Colby constant calculations
- Browse Displays the acquisition method including both the MS and LC parameters

4

Agilent MassHunter Workstation Software: Quantitative Analysis

## Calibration

- CurveFit assistant Calculates all combinations of curves; picks disabled points; and presents results with an equation that is sortable by confidence band and custom filterable by R<sup>2</sup>, standard error and max % residual
- Dilution assistant Calculates and creates calibration levels based on a default or specified serial dilution scheme
- Copy Cal levels Copies calibration levels from one compound to other compounds
- Disable Cal points Disables calibration points based on level, or individual compounds in tables, or interactively through graphs
- Curve fits Supports curves by
  - Type: Linear, Quadratic, First order ln, Second order ln, Average of Response Factors
  - Origin: Ignore, Include, Force, Blank Offset
  - Weight: None, 1/x,  $1/x^2$ , 1/y,  $1/y^2$ , Log,  $1/SD^2$
- Replace curve Creates calibration curves from existing calibration samples
- Average replicates Averages in new replicates into existing calibration curves by compounds
- Import levels Imports calibration levels and concentrations from a file
- Scale graphs Provides graphs with the capability to be auto-scalable by X, Y, X-log, and Y-log; and intelligent zooming to fit specified levels

#### Integrator

- MS-MS integrator Provides a parameter-free integrator at all levels of signals that reduces manual integration efforts
- Integrator metrics Generates metrics that characterize the signal's integration to accept, inspect or reject the integration

These metrics include:

AreaProblem

**Agilent MassHunter Workstation Software: Quantitative Analysis** 

- BaselineProblem
- BedOfSpikes
- HeightProblem
- InterferenceProblem
- MergeProblem
- NoProblem
- SeriousProblem
- Sliced
- SymmetryProblem
- WidthProblem
- Signal-to-noise Calculates signal-to-noise for peaks providing the following choices:
  - ASTM (default)
  - RMS
  - Peak-to-peak
  - Peak-to-peak with drift
- Graphics Shows superior interaction with the graphing of a compound and the display of its peak's
  - Baseline
  - Baseline points
  - Annotations
  - Normalized qualifiers
  - · Uncertainty bands with intelligent autozoom

#### **Batch-at-a-Glance: Results**

- Navigation Moves (previous, next, direct) between samples, compounds, and time segments
- Compound views Switches between the details of the current compound or the summaries of multiple compounds
- Batch table views Enables flat-table layouts or the capability to drill-down to vertically- or horizontally-nested tables for details

Agilent MassHunter Workstation Software: Quantitative Analysis

- Window layout Reorganizes the screen to its defaults, or saves or loads custom-window layouts
- Float pane Floats any pane onto another monitor to enable dual-monitor presentations
- Export Table Exports Batch-at-a-Glance tables directly to Excel files
- Export Graphics Exports any graphic to a customized size in multiple formats
- Copy/Paste Copies or pastes any graphic directly into Microsoft Office applications such as Word, PowerPoint, Excel, etc.
- Print/Preview Prints or previews screen content in WYSIWYG format (what-you-see-is-what-you-get)
- AutoReview Displays each sample automatically and interactively allowing you to stop at any time for closer inspection
- Filter Displays any combination of sample types
  - Sample
  - Blank
  - Cal calibration
  - QC quality control
  - ConCal continuing calibration
  - DblBlk double blank
  - Matrix
  - MatDup matrix duplicate
  - MatBlk matrix blank
  - TunChk tune check
  - RspChk response check
- Sort Sorts any column that appears in a table
- Columns Enables you to add, remove, reorder, save, load, restore, or reset columns
- Resolution Enables viewing screens at different resolutions:
  - Low (1024x768)

Agilent MassHunter Workstation Software: Quantitative Analysis

- Medium (1280x1024) (This resolution is recommended.)
- High (1600x1200)

#### **Outlier Detection**

- Manage Sets up and selects specific outliers that can be detected and individually controlled
- Highlight Highlights outlier values (high-red, low-blue) in the results table
- Filters Lets you display selected types of filters
- Outliers Supports specific types of data for outlier detection
  - Calibration: Accuracy, Average Response Factor, Calibration Range, CurveFit R2, Relative Response Factor, Response Factor
  - QC: QC, QC Relative Standard Deviation
  - Peak Result: Sample Amount, Integration Metric, Limit of Detection, Limit of Quantitation, Signal To Noise Ratio, Relative Retention Time, Retention Time
  - Qualifier: Qualifier Integration Metrics, Qualifier Ratio
  - ISTD: ISTD Response
  - Blank: Blank Concentration
  - Matrix: Matrix Spike, Matrix Spike Percent Difference, Matrix Spike Percent Recovery
  - Surrogate: Surrogate, Surrogate Percent Recovery
  - Response Check: Response Check
- Quantitation message Warns you of samples that encountered serious problems during quantitation

#### Report

- Generate Generates graphics and report results for importing and formatting for Excel XML
- Custom Lets you customize the Excel template

**Batch-at-a-Glance – Batch Table Setup** 

#### Update

- Update/Average RT Updates or averages compound's retention times
- Update Qualifier Ratios Updates qualifier ratios based on compound's current sample
- Update Mass Assignments Updates mass assignments based on compounds current sample

## Qualitative

- Sample Information lets you display the chromatogram and extracted spectra for the current sample
- Chromatogram/Spectrum Provides significant features that can be used to explore spectra for different types of signals

# Batch-at-a-Glance – Batch Table Setup

All quantitative methods and results are based on operating on batches of acquired data. You first set up a Batch Table, adding samples from multiple data files. This table is called Batch-at-a-Glance because you can perform all operations on the data from this View.

Batch-at-a-Glance – Batch Table Setup

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Figure 29 Batch Table

4 Agilent MassHunter Workstation Software Quantitative methods

# Quantitative methods

The Quantitation Method Editor lets you create a new quantitation method from an MRM acquisition method (Figure 30).

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	-	Cocaine	1	304.1 -> 182.0	MRM	Target	304.1	182.0
Globals Setup		Cocaine-d3	1	307.1 -> 185.0	MRM	ISTD	307.1	185.0
Save / Exit		MDMA	1	194.2 -> 163.2	MRM	Target	194.2	163.2
		MDMA-d5	1	199.2 -> 164.3	MRM	ISID	199.2	164.3
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Figure 30 Quantitative view – Method Editor

A file selected from the Batch Table is used as a reference for developing the method settings. These settings are then used to generate the calibration curve and quantitate the standards, QCs and samples. The MRM transitions are already available in pull-down lists and are assigned the corresponding names of compounds.

# **Parameter-free integrator**

## What is the parameter-free integrator?

Agilent has developed a new peak integrator algorithm that works especially well for triple quadrupole MS data. The parameter-free integrator presents these advantages:

- Handles low-level noisy data by setting a peak's starting and ending points statistically
- Adjusts the threshold automatically
- Eliminates the need for manually re-integrating peaks for low-level MRM signals
- Identifies those peaks that appear reliable and those that should be discarded

## **Example of integration results**

Figure 31 shows data at two extremes.



Figure 31 Parameter-free integrator – Data at two extremes

**Parameter-free integrator** 

The lower chromatographic peak could be easily integrated since it is a nice Gaussian-shaped peak, but it would be difficult to define the baseline of the upper peak. In fact, many integrator algorithms might interpret these results as multiple peaks.

However, Agilent's new algorithm had no trouble defining the baseline and recognized this as a single peak. In fact, the new integrator algorithm would integrate this as a single peak even if the baseline were rising, instead of being flat, as shown.

# **Batch-at-a-Glance: Results**

The integration results obtained from the analysis of amphetamine (Amp) are shown in Figure 32. This is a flat view of the Batch Table, Compound Information and Calibration Curve.

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• The Batch Table shows the integration results from applying the quantitation method to each data file. Colored highlights correspond to results that are lower (blue) or higher (red) than expected.

4 Agilent MassHunter Workstation Software Compound confirmation

- The Compound Information view at the lower left displays the integrated chromatographic peaks.
- The Calibration Curve is shown at the lower right.

# **Compound confirmation**

The format shown in Figure 33 can be of value to certified drug-testing laboratories. It shows two sets of plots that can be obtained from a THC analysis.



Overlap of quantifier and qualifier ions

#### Amphetamine qualifier ion -Normalized





Two product ions must be acquired for confirmation: a quantifier ion and a qualifier ion. Typically, the quantifier ion that is used for quantitation is the most abundant of the two product ions.

To be able to confirm the presence of Amphetamine, the qualifier ion peak area must be at least a certain percentage of the quantifier ion, a number that is set in the quantitation method. In this example, 26.5% is used with a window of  $\pm 20\%$ . This means that the area of the qualifier ion must be in the

range of 21.2% to 31.8% of the quantifier ion for the analyte Amp. The qualifier for the ISTD, or Amp - d5, also has a specific range that it must be in.

From the figure on the left, whether or not the qualifier ion falls within the accepted window is not easily determined because the size of the qualifier peak is normalized by a factor of 1/0.265. In the figure on the right, the acceptance window is centered at 26.5% of the quantifier ion peak and the qualifier ion is drawn not normalized, or on the same scale as the quantifier. If the ion is not within the required acceptance window, then it is shaded blue, but still transparent so as not to hide the quantifier ion. This makes it easier to confirm the presence of compounds visually.

# **Compound calibration**

Agilent MassHunter Workstation Quantitative Analysis Software contains several tools to help calibrate and quantitate compounds.

- CurveFit Assistant
- Cursor Pointer for Data Point Information
- Data Point Zooming

# **CurveFit Assistant**

The CurveFit Assistant provides an analytical view of evaluating the possible curve fits (Figure 34).

**Compound calibration** 



#### Figure 34 CurveFit Assistant

Note that the black line drawn through the data points uses Quadratic as the Fit, 1/x as the Weight, and Include as the Origin as shown at top. Many other combinations of the curve settings are listed below the calibration curve with the selected one highlighted in blue. The highlighted settings are also plotted in blue in the curve window.

You can find the best curve fit, for example, one that corresponds to the highest  $R^2$  value, by ordering all of the possible results from the best to the worse  $R^2$  values and then deciding how many data points to consider as being outliers.

For example, the first set of parameters in the list corresponds to a Linear Fit, Ignore Origin and Equal Weight. The corresponding  $R^2$  value is 0.9998001477, which is very good. The corresponding curve can be plotted by simply clicking this entry in the table.

Using these settings, data can be requantitated. Eliminating outliers is common as a standard operating procedure (SOP) in some laboratories.

## Data point information

Overlapping data points are not unusual in a calibration curve, especially with triple quad MS data, where %RSD values are quite low (Figure 35). To help distinguish the data points from one another, the mouse cursor can be moved over the data points to obtain more information about them.



Figure 35 Amp results: Calibration data point information

This figure shows two examples of this type of information. The first example shows that the data points overlap and you are advised to zoom in to see them separately. The second example shows information on the data point itself.

**Compound calibration** 

# **Data point zooming**

You can zoom in on overlapping data points to see individual data points not visible in the visual presentation.

# Reporting

Reports from the Agilent Triple Quad LC/MS are in Microsoft Excel format to increase the flexibility for design and compatibility with other software programs (Figure 36). A set of example reports is also available. These sample reports can be modified to suit your own purposes. For example, you may want to insert the company's logo into the report.

In addition to customizing any report layout included with the Agilent MassHunter Workstation Quantitative Analysis program, you can create a report by moving additional fields from a list of available fields to a desired location in the redesigned report.

Reporting

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24		•			CMAHCaLLS.d		CHA	M_17	P1-C17	5.00	100 L4	Calibration	APCIautotune.m	
					CMAMQC_L2.d		CHA	M_09	P1-C9	5.00	200 L2	QC .	APCIautobune.m	
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Figure 36 Reporting

- You can add macros in Excel
- You can add tables and graphics using tool buttons
- You can drag and drop additional columns from XML results
- You can format font and numbers in Excel
- You can access Print Preview in Excel
# **Agilent MassHunter Workstation Software: Qualitative Analysis**

# **Overview**

Qualitative Analysis includes many capabilities to help you perform these operations:

- Review your data
- Extract chromatograms
- View spectra and extract peak spectra
- Subtract the background
- Integrate the chromatogram either interactively or automatically

With this software you can easily handle viewing lots of data because most operations are accessible from one screen. You can set up a method while viewing data, then apply the method to the data and see the results change.

4

#### 4 Agilent MassHunter Workstation Software

**Overview** 



Figure 37 Qualitative Analysis Main View

# Capabilities

Here are some of the capabilities:

# Set up methods to automatically perform actions when files are opened

You can choose from a list of file actions to set up for the software to apply when a data file(s) is opened.

Method Editor: Assign Actions on Opening a Data File	
🖸 🚺 🕟 Assign File Open Actions 🔹 🕑 🏢	
Available actions	
Extract Defined Extract Peak Spectra Integrate And Extract Peak Spectra Integrate Chromatograms Extract Peak Spectra Print Qualitative Analysis Report	
Actions to be run	
Extract Defined Chromatograms	
	X

## Easily compare data from many different types of files

Not only can you compare chromatograms and spectra from scan versus SIM versus MRM runs, but you can also compare data from Triple Quad runs versus Q-TOF runs. In addition, you can compare MS data, UV data and instrument curves from one data file.

## Appearance of only relevant fields for data type

The software recognizes file data and presents choices based on the data available.

4 Agilent MassHunter Workstation Software Capabilities

#### **Data Navigator**

Total ion chromatograms, base-peak chromatograms, extracted ion chromatograms (EIC) and spectra can be organized in lists according to data type (chromatograms or spectra) or according to data file.



You can highlight only those chromatograms and spectra you want to view, then mark the appropriate check box, and you can view only these in Data Navigator and the display.

All data in windows are linked so that you know which chromatogram, spectrum, mass or results list belongs to which data file or chromatogram or spectrum. For example, if you click a mass in a spectrum, you will see the peak in the chromatogram the spectrum came from and the data type and data file for that spectrum displayed in the Data Navigator.

#### Actions on more than one chromatogram or spectrum

You can perform any operation on more than one piece of data. Highlight the chromatograms or spectra that you intend to operate on, and the action works on all those items highlighted.

#### **Data filters**

The software presents many choices for filtering data before extracting or integrating.

## **Flexible window layout**

You can look at all the data you want to look at and customize the window layout for doing so. You can also save window layouts.

#### Optimum use of display real estate

You can select how many chromatographic or spectral plots you want to see at one time. The display will contract to show you more and expand to show less.

## Program saves all changes

Any change made to a method or a result can be saved. When you exit and reopen the files, you can reload the results that you created. You can also save methods and results to separate files. 4 Agilent MassHunter Workstation Software Capabilities

#### **Useful for method development**

Agilent MassHunter Workstation Qualitative Analysis Software is most useful with the Agilent Triple Quad LC/MS during method development. With this software you can:

- View the optimum transitions for quantitation
- View the results of optimizing voltages with a worklist of samples to select the best parameter set

## www.agilent.com

# In This Book

The *Concepts Guide* presents "The Big Picture" behind the Agilent 6410 Triple Quad LC/MS to help you to understand how the hardware and software work.

This guide includes concepts for:

- Overview
- Inner Workings
- MS and Sensitivity
- MS Workstation Software

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