

Expect more productivity

with the evolution of ultimate performance GC-MS/MS

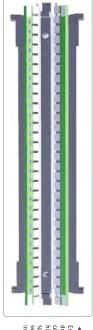




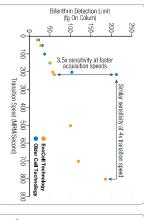


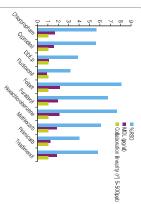
instrument control Fast EvoCell technology and intelligent

with the high efficiency timed-SRM capability of the TSQ 8000 Evo GC-MS/MS, opens the door to new possibilities for high-complexity methods in routine laboratories. This enables higher levels of productivity and efficiency in analytical workflows. Innovative EvoCell collision cell technology, in combination More capacity is offered through the acquisition of more information, more compounds, and more results per unit time in your laboratory.



Enhanced velocity optics driving EvoCell collision cell technology provide high SRM transition speeds, precision, and sensitivity for even the most complex methods





Increased sensitivity at fast transition speeds allows the use of up to 4x transition speeds of standard collision cell technology

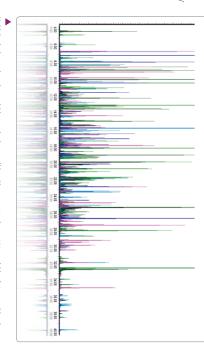
Analysis of pesticides in rice (10 $\mu g/kg)$ with the EvoCell collision cell at 500 μs dwell time, acquiring 800 transitions per second

	m.	EvoCell Collision Cell Technology	
_	Features	Benefits	Impact
	Increased number of transitions per compound	More points of confirmation More resistance to matrix interference	Higher confidence More on-time results
	Increased number of compounds	Higher capacity methods	More method consolidation More efficiency in result production
	Fast GC compatible	Faster run times	Faster turnaround More on-time results
	Wider SRM windows	More resistance to the effects of RT shift caused by matrix	Faster turnaround More on-time results

Truly powerful methods use speed efficiently

particular compound. by automatically optimizing the targeting of a MS system dramatically reduces this complexity increasingly complex. The TSQ 8000 Evo GC-MS/ run, managing acquisition windows becomes As more compounds are added to a single

mass spectrometer is not consuming valuable required to capture the peak, and t-SRM takes times when those compounds are not eluting. resources scanning for compounds at the sensitivity of each individual analyte. The method without compromising the excellent and more compounds can be added to the detection is optimized for maximum sensitivity care of the rest. This ensures that compound Simply enter the retention time and the time



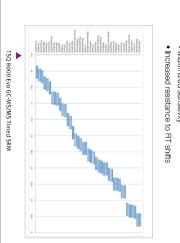
Multiple single quadrupole pesticides methods consolidated into one 40-minute, 350 pesticides (700 transitions) run

Traditional Segmented SRM

- Complicated set-up
- Wasted dwell time
- Reduced sensitivity
- Reduced tolerance to RT shifts

TSQ 8000 Evo GC-MS/MS Timed SRM

- Automated set-up
- Optimized dwell time
- Maximized sensitivity

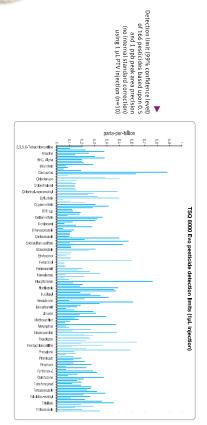


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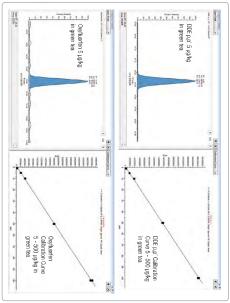
Traditional Segmented SRM

Ultimate sensitivity SRM

GC-MS/MS offered on the market sensitivity selected reaction monitoring (SRM) enabled by the TSQ 8000 Evo GC-MS/MS, which is the highest performing triple quadrupole Laboratories performing analyses in food safety, environmental monitoring, and sports doping can all benefit from high selectivity and ultimate



quadrupole instruments. is a perfect fit for laboratories in above and beyond former triple productivity. Expect performance into real routine efficiency and speed, and sensitivity translate methods. Enhanced selectivity, need of high efficiency pesticide The TSQ 8000 Evo GC-MS/MS

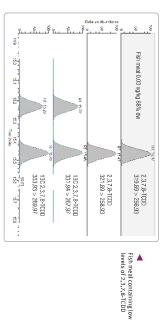


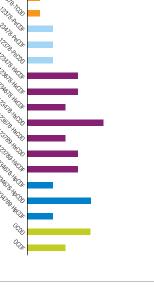
High selectivity of low concentrations in complex tea matrix

Expect **More** Performance

at the lowest levels Ultimate sensitivity for contaminants regulated

GC-MS/MS delivers routinely extremely powerful SRM that the TSQ 8000 Evo criteria in force. Meeting that challenge requires extremely low regulated levels and stringent analytical contaminant analysis. This challenge is due to the challenging of the ultra-trace analytical methods in dibenzodioxins and furans (PCDD/Fs) is one of the most The confirmatory determination of polychlorinated





Concentration [pg/µL]

0.03

0.08-0.09-

Instrument detection limit (IDL) for PCDD/Fs
when satisfying all confirmatory criteria CSL x 5
dilution (n=10) injected using splitless mode

	State the state of	
European Commission GC-MS/MS Confirmatory Performance Criteria	TSQ 8000 Evo GC-MS/MS Capabilities	Compliance Confirmed
Two specific precursor ions with two specific production ions	All recommended methods developed as defined in criteria Fully optimized by AutoSRM	Yes
Tolerance of ion ratios within ± 15%	< 10% measured at EPA 1613 CSL level (n=14)	Yes
Resolution of each quadrupole equal to or better than unit mass resolution	All recommended methods developed Q1 and Q3 at 0.7 Da	Yes
The % RSD of the five (or more) Relative Response Factors (RRFs) for each unlabeled POD/PODF and ballets) for each unlabeled moust not exceed 20%	6 point curve EPA 1613 CSL-CS4 2%	Yes

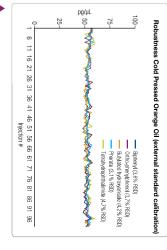
Expect **More**Robustness

Expect **More**Flexibility

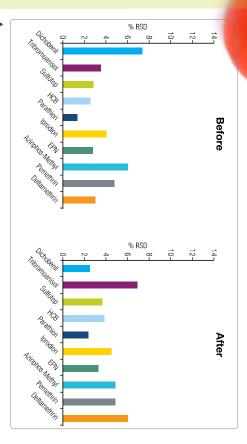
Unstoppable productivity

The Thermo Scientific" ISQ" Series single quadrupole GC-MS and TSQ 8000 triple quadrupole GC-MS/MS have a strong reputation in allowing laboratories to tackle the demands of producing results on time, every time. The TSQ 8000 Evo triple quadrupole GC-MS/MS continues that legacy by enabling routine-grade robustness for ultimate performance applications in some of the most challenging matrices.

The patented Thermo Scientific"
ExtractaBrite" ion source is heated throughout and includes the first Fir egion (RF lens), commonly the first place susceptible to ion burn. This design ensures high matrix tolerance for high throughput applications. Crucially, it is fully removable, even when hot, under vacuum for necessary cleaning or swapping with a spare source. There is never a need to vent to clean your instrument or to heat or clean quadrupoles. Enjoy constant productivity.







%RSD pesticides (10 ppb) before and after 1,400 vegetable matrix injections

Flexibility to best solve your analytical challenge

The TSQ 8000 Evo GC-MS/MS addresses many laboratory mass spectrometry needs through various scan modes along with source and probe options Excellent performance is enabled for single quadrupole-type scan modes which, in turn, expand system versatifty in the modern laboratory.

		Q2 (EvoCell)		
Selected ion monitoring (SIM)	Transmission only	Transmission only	Selected m/z	Target quantitation
Full scan	Transmission only	Transmission only	Scanning	Unknown ID, library search
Selected reaction monitoring (SRM)	Selected m/z	CID	Selected m/z	Target quantitation
Product ion scan	Selected m/z	CID	Scanning	SRM method development and compound elucidation, build and search libraries
Precursor ion scan	Scanning	CID	Selected m/z	Searching for unknown compounds giving characteristic fragments
Constant neutral loss	Scanning	CID	Scanning (- neutral loss offset)	Searching for unknown compounds with common structures, e.g. acids
SRW/Full Scan	Alternating selecting m/z and transmission only	Alternating CID/ Transmission only	Alternating selecting m/z and scanning	Target quantitation and unknown ID by library search

lon sources and probes are removable without requiring a break in the vacuum, and they enable you to perform MS and MS/MS experiments on solids in real time while effortlessly switching between electron ionization (EI) and chemical ionization (CI).

Direct Insertion Probe (DIP)
Slower volatilization with heated capillary
tube for solid samples. The DIP can be
utilized for rapid analysis of solids or trace
components in solid matrices such as
forensic samples, tissue, etc.

Direct Exposure Probe (DEP)
Rapid heating filament for liquids or solutions.
The DEP is ideal for rapid molecular weight
confirmation of solids dissolved or suspended
in a suitable solvent.

Chemical Ionization

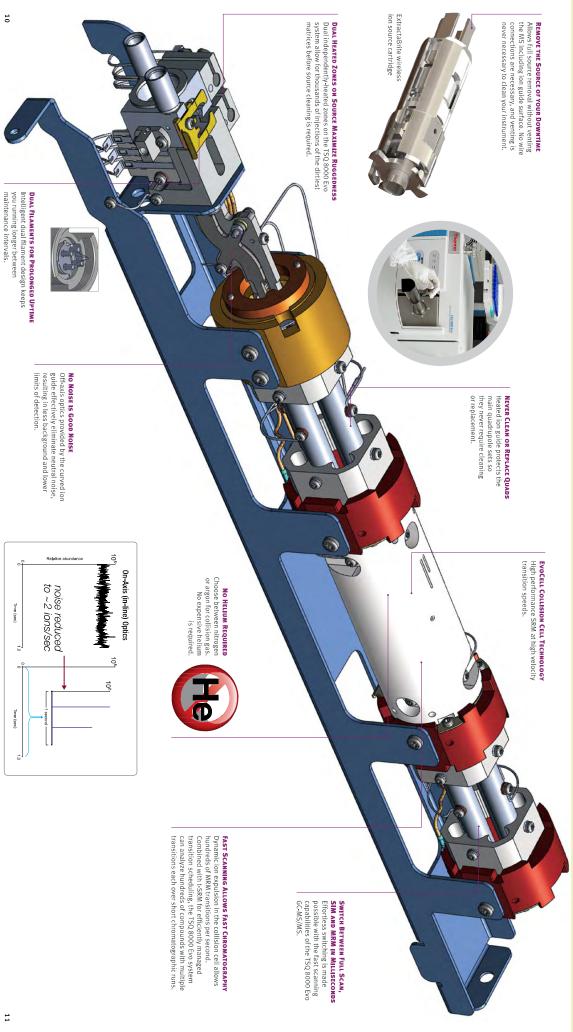


Chemical ionization is a softer form of ionization than electron ionization (El). It offers a greater opportunity to generate more abundant high mass and molecular ions and a higher degree of selectivity or sensitivity in the source ionization process, especially with negative chemical ionization (NCI), favoring electronegative compounds, such as halogenated species. Chemical ionization is ideally used for structure elucidation also in MS/MS mode.

brilliant **Design** for brilliant Results

Industry-Leading Detector Linearity

the ultimate quantitative instrument. Combined with the low detection limits attainable by SRM, this detector makes the mass spectrometer The Thermo Scientific DynaMax detection system standard on the TSQ 8000 Evo system offers industry-best linearity.



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Expect More Simplicity

Easy MS/MS

The advantages of selected reaction monitoring (SRM) and multiple reaction monitoring (MRM) analysis often come with challenges, especially with respect to adoption of mer technology in the laboratory. These challenges are due to the method set-up and optimization and management of complex analytical methods. Laboratories that strive to realize the benefits of MS/MS must overcome some barriers in the tuning, set-up and optimization to reduce the time required to achieve a smooth operating routine.

The TSQ 8000 Evo GC-MS/MS was built to guarantee an easy start-up. Whether you are managing retention times, starting a completely new analysis, transferring a method from a single quadrupole system, or porting a known MRM method from another instrument, the TSQ 8000 Evo system ensures the fastest route to routine high performance results through its integrated software tools.

AutoSRM: A purpose-built software for automated SRM method creation and optimization from full scan to the complete analytical method setup

SRM Optimization

- TSQ 8000 Evo GC-MS/MS Instrument Method: Offers true timed-SRM operation, allowing for high sensitivity and ease of use for the most complex SRM methods
- Thermo ScientificTM TraceFinderTM Software: State of the art, multi-platform user-friendly chromatography analysis software

om the Star

For new compounds, choose to begin the AutoSRM function, ask for a precursor ion study (Step 1), and place a vial with the compound mix in the autosampler. The AutoSRM function reports a full scan chromatogram of the injection, and you simply identify your peaks and choose the most appropriate precursors that AutoSRM presents. In the case of coelluting compounds, AutoSRM automatically performs one injection per compound.

Method Sync

From an Existing SIM Method

If you are moving from a single quadrupole GC-MS method using SM analysis, or you already know which ions you want to select for your precursor ions, you can start with Step 2. The AutoSRM function performs product ion scans for each selected precursor and presents the results in a single window. Simply select (or ask AutoSRM to select) the best product ion for that compound.

From a Previous MS/MS Method

If you have already performed previous AutoSRM steps or are simply moving from another MS/MS system, then Step 3 can take you to the final fully optimized SRM transitions for your compounds. Fully automated collision energy optimization is performed to ensure that transitions are as sensitive as they can be. These optimized SRM transitions are then exported into a finished MRM method ready for analyzing real samples in routine.

Your Starting Point

Unknown SRM transitions

Start from the very beginningStart with existing SIM method

Known SRM transitions

From an existing method

From SRM transition list

Method

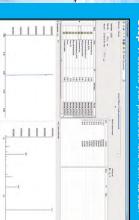
Optimized SRM transitions

From Compound Data Store
From existing TSQ 8000 Evo
GC-MS/MS method

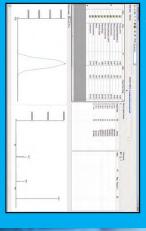
GC-MS/MS method

AutoSRM Workflow

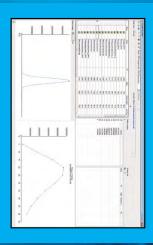
p 1: Selects your precursor ions from full scar



Step 2: Selects your product ions from product ion scans



Step 3: Optimizes the collision energy for selected transitions



Be More Productive

Fast, flexible, easy-to-use mass spectrometry software

The challen

The constant pressure to increase productivity challenges laboratory

safar at every level, from the technician to management. Streamlined, workflow-driven software solutions are needed to address these challenges. At the same time, the growing list of compounds that laboratories must quantify and screen, combined with increasingly complex regulatory requirements, necessitate flexible method development and powerful reporting tools.

TraceFinder software is the solution

TraceFinder software makes routine quantitation simple, fast and productive. It offers a complete solution that can be used to develop methods, acquire and process data, and generate standard and customized reports with the full portfolio of Thermo Scientific mass spectrometers.

Powerful method development

TraceFinder software provides powerful tools for method developers, including:

- Compound databases (CDB) for rapid selected reaction monitoring (SRM) method development
- User-defined flags for nearly every parameter
- Automated retention time and ion ratio adjustment

| Composition |

Compound Database contains hundreds of compounds and parameters for SRM work flows, allowing rapid method development

Simple, full-featured data review

TraceFinder software offers simple, yet exceptionally feature-rich data review capabilities. The intuitive, workflow-driven arrangement of information simplifies the technician's role in reviewing data. The technician can quickly and easily observe the data processing status, view compound-related flegs to locate compounds which need attention, and take appropriate steps to address any problems.

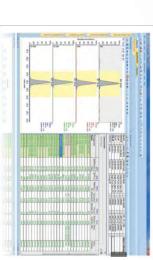
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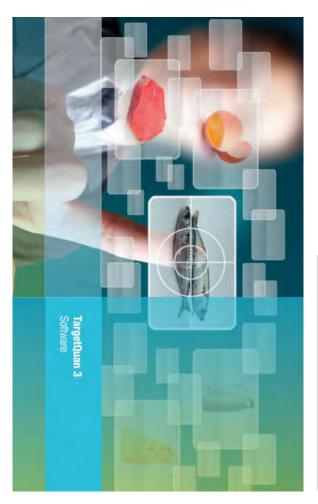
Powerful, automated reporting

The new TraceFinder software report designer offers enhanced laboratory productivity with customizable, automated reporting with numerous standard report templates.

Leading the way in regulatory POPs quantitation

Thermo Scientific TargetQuan software is uniquely tailored for a workflow-orientated POPs quantitation, like PCDD/Fs, PCBs, PBBEs and others. This software package is designed specifically to serve laboratories charged with performing routine quantitation of POPs in a regulated environment. With TargetQuan 3 software, comprehensive processing of MS, MS/MS and HRMS data is possible on a single software platform, compliant with the US EPA 1613 requirements.





Experience More innovation

Example of plug-in module installation by user

Tackle future challenges through GC modularity

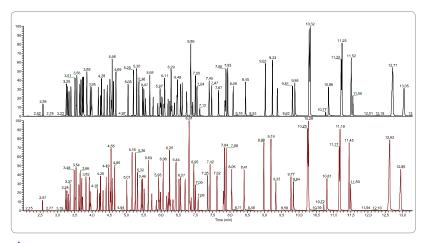
Enable Quick Customization with GC Modularity

Tailor the Thermo Scientific™ TRACE™ 1300 Series GC to your needs with its proprietary user-exchangeable Instant Connect injector and detector modules. Swapping modules is easily done by the removal and replacement of three screws, accessible from the top of the GC. The entire process takes less than five minutes without requiring specialized service assistance. This enables budget-conscious laboratories to purchase a base-configured GC today, then easily expand their capabilities to accommodate new application and throughput demands.



GC and GC-MS	Estimated Cylinder Lifetime		
Operating Conditions*	Conventional Operation	With the Instant Connect Helium Saver Module	
Around the Clock Analysis 24 hours a day, 7 days a week, 365 days a year	5 months	3.5 years	
Daily Operations Leaving the GC idle at the end of workdays and on weekends	15 months	14.6 years	

^{*} Conditions: Operating a TRACE 1300 Series GC with helium at 4 mL/min (sccm) using a typical helium cylinder of 48 L volume at 2250 psig.



Using the Instant Connect Helium Saver Module, results from US EPA Method 8270C (semi-volatiles) remain unchanged

www.thermoscientific.com/tsq8000evo

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Ensure Uninterrupted Analyses and Save Budget by Conserving Helium

The helium shortage continues to inflict price pressure and supply uncertainty on laboratories that require these gas supplies to perform their research and analyses.

Realizing the importance of finding a safe, cost-effective solution, the proprietary Instant Connect Helium Saver Module has been introduced to drastically reduce helium carrier gas consumption and extend helium cylinder lifetime up to 14 years per instrument, without any GC or GC-MS method modifications. Helium is continuously saved, both while the GC is in operation and while it is idle. Previously acquired retention times remain unchanged, and no method revalidation is required.

Backflush and Large Volume Capabilities

The capabilities of the Instant Connect injector modules are further enhanced by the available concurrent backflush options. These solutions enable the user to reverse the flow inside the injector, eliminating heavy or "undesired" compounds concurrently during the analysis run, protecting the column and detector while cutting down non-productive times, thus increasing throughput.

Instant Connect-SSL concurrent backflush. The T-piece for column connections and its carrier gas control are integrated within the module





A Thermo Fisher Scientific Brand