

Toxicology

Fit for purpose.
Increase productivity.
Consistently accurate results.



Tox Explorer Collection

**An all-in-one LC-MS/MS
solution for toxicology**

Navigate with ease

An all-in-one LC-MS/MS solution for toxicology

Detect an ever-increasing number of substances at ever-decreasing concentrations. The Thermo Scientific™ Tox Explorer™ Collection helps the world's toxicology community in forensics, clinical research, employee drug testing, and sports anti-doping navigate with ease to solve complex analytical challenges and increase productivity. Now you have a choice of either Thermo Scientific™ Orbitrap HRAM or Thermo Scientific™ TSQ triple quadrupole mass spectrometer platforms. Choose the best option based on your needs.



Keep detection capability one step ahead with high-resolution accurate-mass (HRAM) or triple quadrupole mass spectrometry options of the Tox Explorer Collection.



Streamline your methods using sample preparation guidelines, high performance HPLC columns, intuitive software capabilities, and a comprehensive mass spectral library or compound database.



Get excellent dynamic range and highly accurate mass measurements. HRAM option includes over 1,800 compounds of interest in the spectral library.



Increase productivity with SRM acquisition speeds enabling faster polarity switching. Triple quadrupole MS option includes over 1,200 compounds in the compound database.

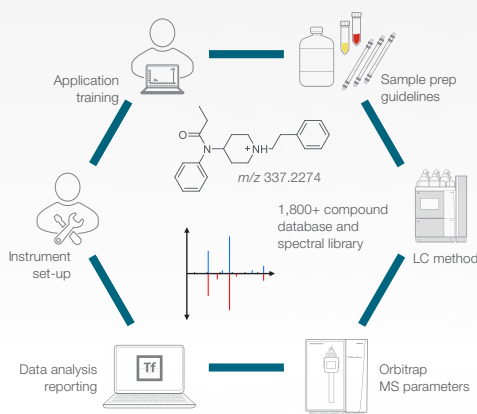
Comprehensive LC-MS solutions

Choose from two LC-MS platforms

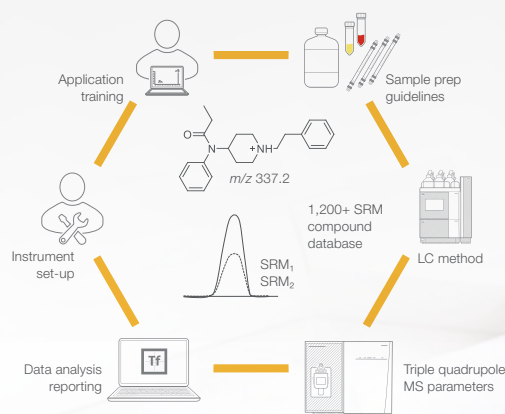
Optimized for quick start-up and ease of use, the Tox Explorer Collection is designed to improve efficiency and reduce cost throughout the preparation, separation, detection and analysis of drug compounds in multiple biological matrices such blood, plasma, and

urine with unprecedented speed, ease, and confidence, regardless of method complexity or user experience. It is equipped with a standardized LC-MS/MS method, UHPLC column, experimentally obtained database with retention times, and LC-MS/MS instrument set-up and

application training enabling smooth start-up and implementation. Choose from two mass spectrometry options; each is a comprehensive LC-MS solution from sample processing to reporting of toxicology analytes.



Tox Explorer Collection for HRAM MS



Tox Explorer Collection for Triple Quad MS

Vanquish Flex Binary UHPLC system



Productivity and quality in chromatography

With confident retention time stability and maximum pressure up to 1,000 bar, the Thermo Scientific™ Vanquish™ Flex UHPLC systems increase productivity without compromising quality of results. The platform shares all the Vanquish system values, such as design focused on uptime, robustness, and quality, enabling every toxicology laboratory to address critical screening and quantitation challenges.

- Designed for maximum flexibility.
- Achieve separations with better reproducibility.
- Increase productivity with high sample throughput.
- Integrates seamlessly with Thermo Scientific mass spectrometers with dedicated software.

UHPLC columns

Fast separations with excellent resolution and sensitivity

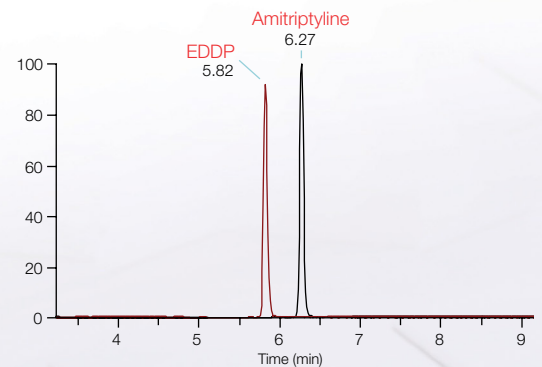
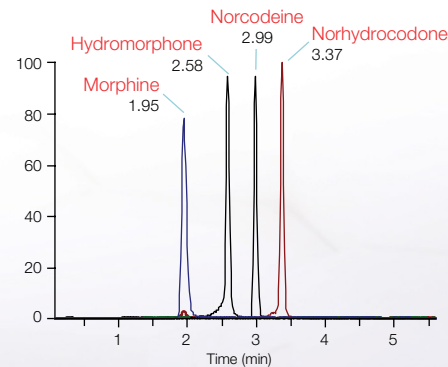
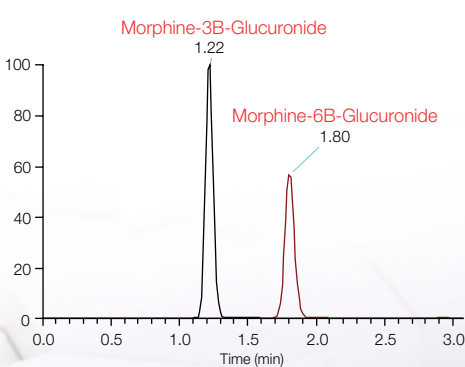
Based on Core Enhanced Technology, Thermo Scientific™ Accucore™ columns provide fast, high-resolution separations without the elevated backpressures required by sub-2 µm particles.

- Robust columns offer wide separation chemistry for all UHPLC applications.
- Highly efficient, rugged separations are ideal for LC-MS applications.



Chromatographic resolution

The 15-minute LC-MS/MS method of the Tox Explorer Collection is capable of separating drugs of abuse isomers. Screening is based on retention times for each specific isomer.



Commonly found isomers in drugs of abuse screening

Tox Explorer Collection for Orbitrap Exploris 120 mass spectrometer



Tox Explorer Collection for Orbitrap Exploris 120 MS

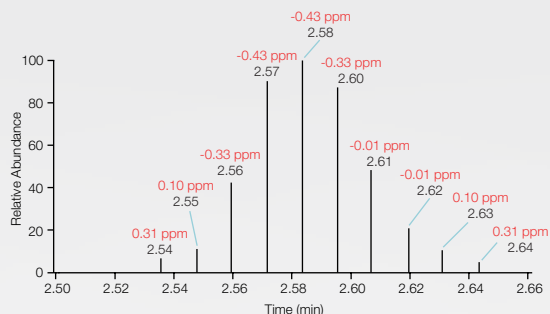
High resolution qualitative and quantitative performance

Navigate with ease using fast scan rates, polarity switching, and an over 1,800-compound database with corresponding high-resolution spectral library. A standardized method and onsite set-up and training ensures less time spent on method development.

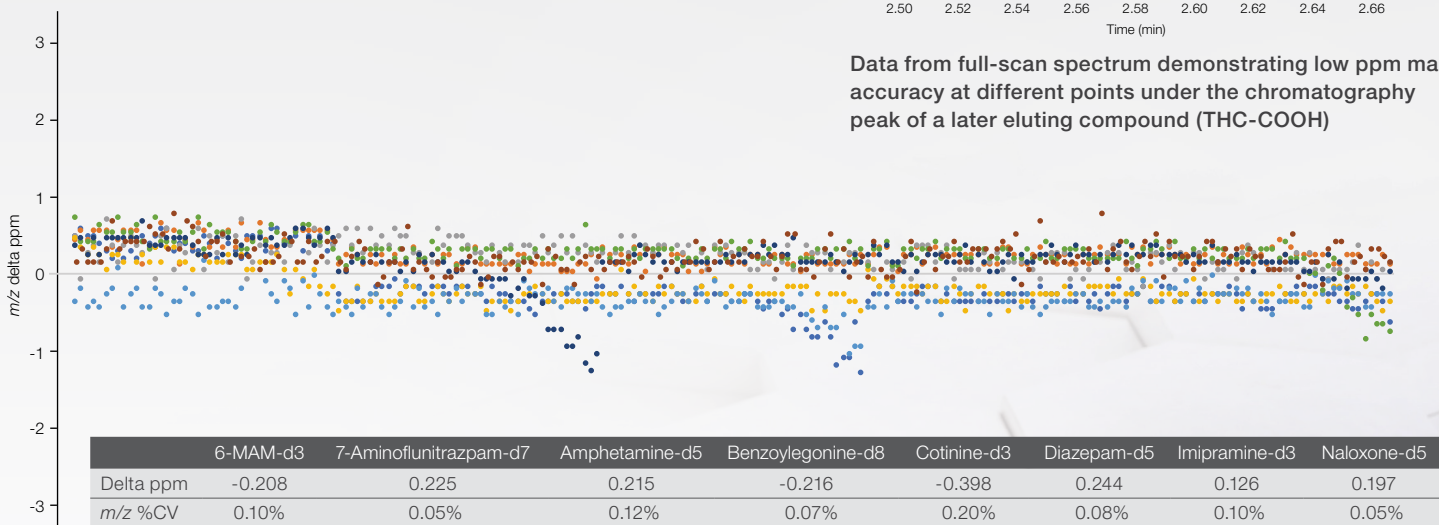
- **Pos/neg switching:** standard
- **Column:** Accucore Phenyl Hexyl, 100 × 2.1 mm, 2.6 μm UHPLC column
- **Software:** Thermo Scientific™ TraceFinder™ 5.1 software
- **CDB and library:** 1,800 compounds; experimentally obtained with true standards
- **Mass range:** m/z 40–3,000
- **Onsite installation:** included— instrument install and configuration set-up
- **Onsite training:** included— 2-days on-site install, support and Tox Explorer Collection application training
- **Method:** 15-min w/RT for 1,800 compounds (HRAM tubing config.)
- **LC:** Vanquish Flex Binary UHPLC system
- **MS:** Thermo Scientific™ Orbitrap Exploris™ 120 mass spectrometer
- **HRAM resolution:** Full Scan 60K, ddMS² 15K

High resolution and mass accuracy

Thermo Scientific™ Orbitrap™ technology platforms offer high resolution mass spectrometry and unparalleled mass accuracy without loss of sensitivity and provides positive/negative polarity switching in a single analytical run, saving time and offering greater confidence in data output.



Data from full-scan spectrum demonstrating low ppm mass accuracy at different points under the chromatography peak of a later eluting compound (THC-COOH)



Stable and precise mass accuracy is shown for eight internal standards. Masses were within 1ppm; all were below 1% CV for over 200 injections across over 2 days without needing system recalibration

Tox Explorer Collection for TSQ Quantis Plus mass spectrometer

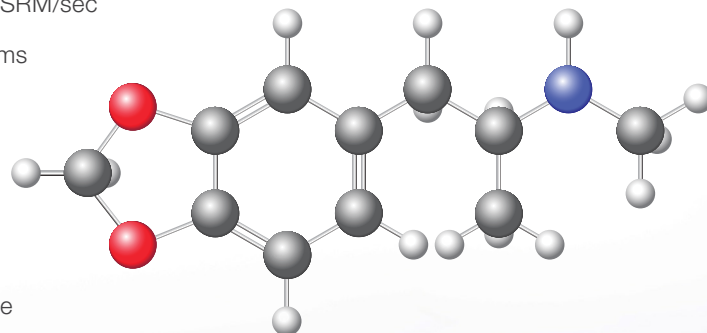


Tox Explorer Collection for
TSQ Quantis Plus MS

Accessible quantitative performance

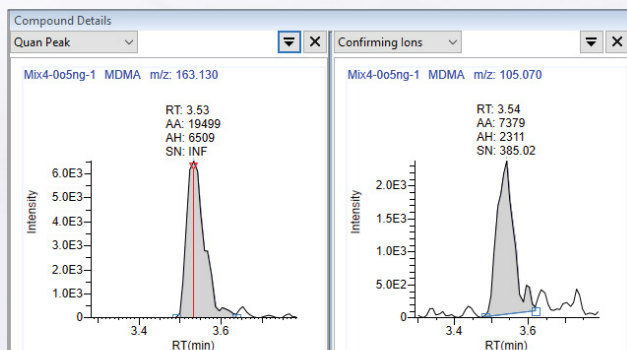
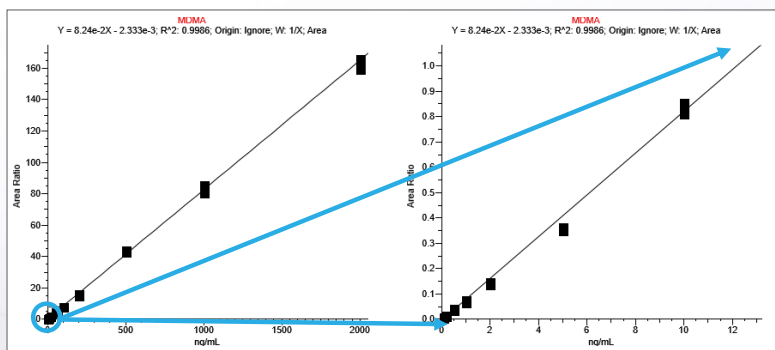
Achieve scientific and business goals by accessing over 1200 compounds with superior acquisition speeds and polarity switching. Quantitative precision and accuracy delivered through innovative product designs enable TSQ Plus mass spectrometers to address the ever-changing drug landscape with consistency and reliability.

- **Column:** Accucore Phenyl Hexyl, 100 × 2.1 mm, 2.6 μm UHPLC column
- **Software:** TraceFinder 5.1 software
- **CDB and library:** 1,200 compounds; experimentally obtained with true standards
- **Mass range:** m/z 2–3,000
- **Onsite installation:** included— instrument install and configuration set-up
- **Onsite training:** included— 2-days on-site install, support and Tox Explorer application training
- **Method:** 15-min w/RT for 1,200 compounds (QQQ tubing config.)
- **LC:** Vanquish Flex Binary UHPLC system
- **MS:** Thermo Scientific™ TSQ Quantis™ Plus mass spectrometer
- **SRM acquisition:** 600 SRM/sec
- **Pos/neg switching:** 5 ms



Targeted screening and quantitation

Incorporating the TSQ Quantis Plus mass spectrometer for drugs of abuse testing enables sensitive and reproducible analytical measurement needed for confirmation for compounds covering a range of compound classes and polarities.



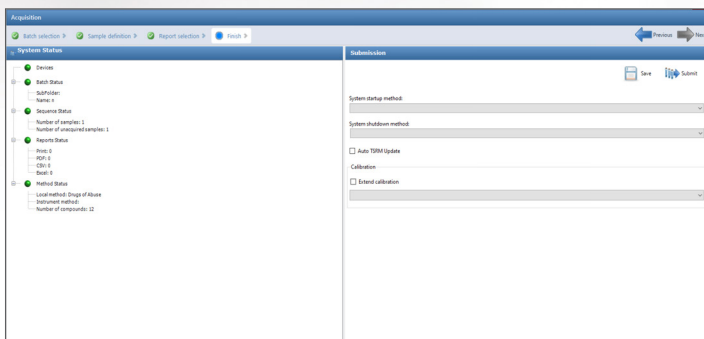
Limits of confirmation (LOC) were determined for MDMA with concentrations ranging from 0.1 to 2000 ng/mL (left) as well as chromatogram results from MDMA at its LOC of 0.5 ng/mL showing quantitation and confirming peaks (right). Calibration curves for quantitation were linear with R^2 values of >0.99 and replicates all <30%.

TraceFinder software for fast actionable results



Simplify high-throughput screening and quantitation

Driven by an intuitive set of acquisition and data processing tools, TraceFinder software provides rapid access to results and increases output across the entire laboratory. Enhance productivity with built-in intelligence such as four-click acquisition and saved templates. A single software platform for method development, acquisition, and processing increases efficiency and confidence. Finally, benefit from increased application productivity using customizable views for reporting, flagging, and screening criteria.



Simplify routine quantitation

Ease of use

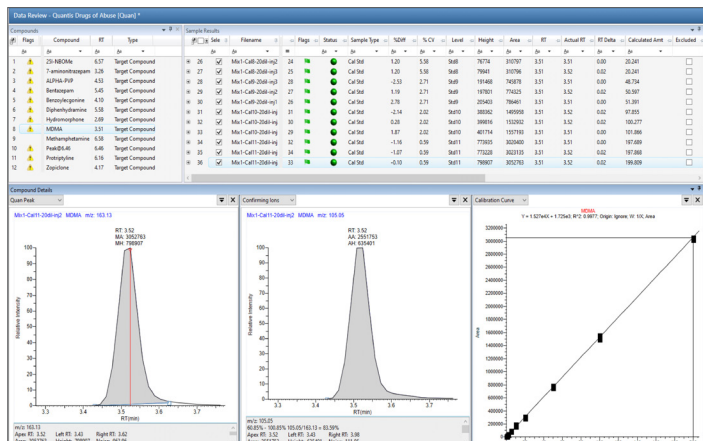
- Four-click acquisition
- Saved templates
- Workflows
 - Quantitation, screening and unknown analysis
 - Small and large molecule analysis

Compound Name	Peak Label	Peak Workflow	Associated Target Peak	Chemical Formula	MS Data	Precision m/z	Product m/z	Adduct	Polarity
1 Ketamine	T1.238.28912	TargetPeak		C15H23NO	m/z	238.28912	0.000	238.28912	Hydrogen
2 Ketamine	T1.238.28912-125	Confounding	T1.238.28912	C15H23NO	m/z	238.28912	125.01541	238.28912	None
3 Ketamine	T1.238.28912-126	Confounding	T1.238.28912	C15H23NO	m/z	238.28912	126.01541	238.28912	None
4 Ketamine	T1.238.28912-178	Confounding	T1.238.28912	C15H23NO	m/z	238.28912	178.02211	238.28912	None
5 Ketamine	T1.238.28912-207	Confounding	T1.238.28912	C15H23NO	m/z	238.28912	207.02717	238.28912	None
6 Ketamine	T1.238.28912-220	Confounding	T1.238.28912	C15H23NO	m/z	238.28912	220.02627	238.28912	None
7 Ketamine	T1.238.28912-121	Fragment	T1.238.28912	C15H23NO	m/z	238.28912	121.0154	238.28912	None
8 Ketamine	T1.238.28912-126	Fragment	T1.238.28912	C15H23NO	m/z	238.28912	126.0154	238.28912	None
9 Ketamine	T1.238.28912-178	Fragment	T1.238.28912	C15H23NO	m/z	238.28912	178.0221	238.28912	None
10 Ketamine	T1.238.28912-207	Fragment	T1.238.28912	C15H23NO	m/z	238.28912	207.0271	238.28912	None
11 Ketamine	T1.238.28912-220	Fragment	T1.238.28912	C15H23NO	m/z	238.28912	220.0263	238.28912	None
12 1,4-Methoxyphenylamine	T1.181.13154	TargetPeak		C11H13NO	m/z	181.13154	0.000	181.13154	Hydrogen
13 1,4-Methoxyphenylamine	T1.181.13154-1182	Confounding	T1.181.13154	C11H13NO	m/z	181.13154	118.01354	181.13154	None
14 1,4-Methoxyphenylamine	T1.181.13154-1182	Confounding	T1.181.13154	C11H13NO	m/z	181.13154	118.01354	181.13154	None
15 1,4-Methoxyphenylamine	T1.181.13154-1184	Confounding	T1.181.13154	C11H13NO	m/z	181.13154	118.01354	181.13154	None
16 1,4-Methoxyphenylamine	T1.181.13154-121	Confounding	T1.181.13154	C11H13NO	m/z	181.13154	121.0254	181.13154	None
17 1,4-Methoxyphenylamine	T1.181.13154-122	Confounding	T1.181.13154	C11H13NO	m/z	181.13154	122.0254	181.13154	None
18 1,4-Methoxyphenylamine	T1.181.13154-118	Fragment	T1.181.13154	C11H13NO	m/z	181.13154	118.0135	181.13154	None
19 1,4-Methoxyphenylamine	T1.181.13154-118	Fragment	T1.181.13154	C11H13NO	m/z	181.13154	118.0135	181.13154	None
20 1,4-Methoxyphenylamine	T1.181.13154-118	Fragment	T1.181.13154	C11H13NO	m/z	181.13154	118.0135	181.13154	None
21 1,4-Methoxyphenylamine	T1.181.13154-121	Fragment	T1.181.13154	C11H13NO	m/z	181.13154	121.0255	181.13154	None
22 1,4-Methoxyphenylamine	T1.181.13154-122	Fragment	T1.181.13154	C11H13NO	m/z	181.13154	122.0255	181.13154	None
23 1,4-Methoxyphenylamine	T1.181.13154-122	Fragment	T1.181.13154	C11H13NO	m/z	181.13154	122.0255	181.13154	None

Efficiency and confidence

One software platform

- Method development, acquisition, processing
- Compound databases and spectral libraries
- Auditing/administration



Application productivity

Customization

- Custom report generation
- Customized data viewing
- Smart sample flagging
- Screening criteria
- Auditing/administration

mzVault mass spectral library



Confident identification of toxicology compounds

When using Orbitrap technology, the Thermo Scientific™ mzVault™ mass spectral library seamlessly integrates with the acquisition and processing TraceFinder software while providing an extensive repository of valuable information on critical analytes that enables confident identification of small molecules.

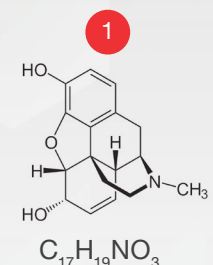
- Includes exact mass of compounds
- Fragmentation data at different collisional energies helps identify unknown compounds

Workflow for adding compounds to the database

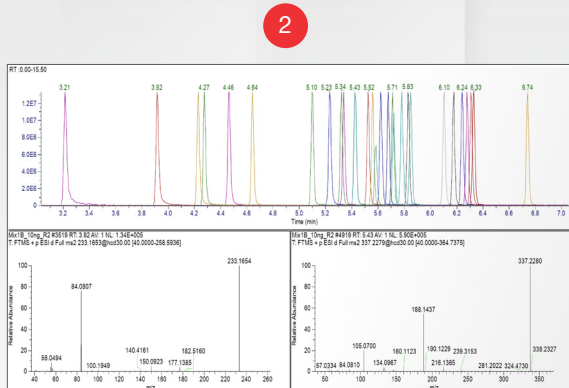
Addition of new compounds to a database can be done with ease because manual optimization of individual precursors is not required. New compounds can easily be added to the existing method by collecting full scan data with targeted MS² spectra. Once the retention time is known, the information is simply added to the compound database and library. There is no need for

revalidation because the cycle time/dwell time does not change in the method.

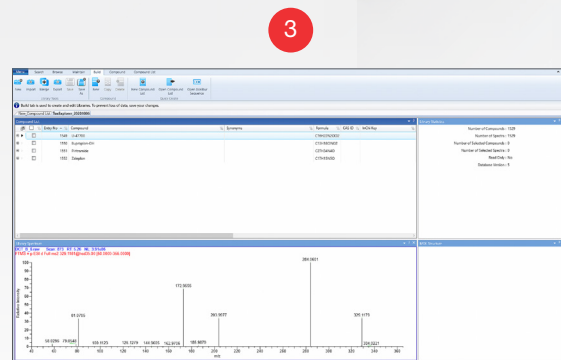
Users can also add their own spectra to the existing library or create their own library. The customer will need to update the retention times of the new analytes that are added to the database.



Known data:
m/z and empirical formula to calculate isotope pattern



Run standard and obtain RT and MS² spectrum (includes fragments)

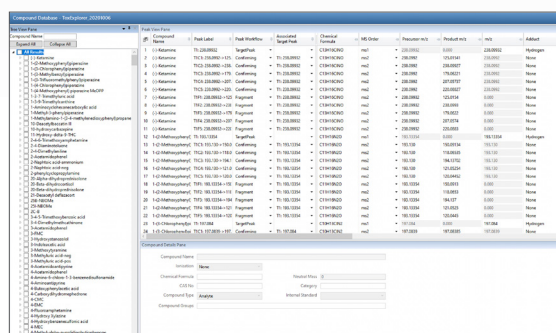


Import MS spectrum in mzVault

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Mass [m/z]	Formula [M]	Species	CS [I]	Polarity	Start [min]	End [min]	(NCE)	MSX ID	
238.2032				Positive	3.44	4.41			0) Ketamine
193.13354				Positive	2.85	3.85			1) 2-Methoxyphenylpiperazine
197.08400				Positive	3.56	4.56			1) 3-Chlorophenylpiperazine
191.15428				Positive	1.96	2.96			1) 3-Toluylphenylpiperazine
231.11036				Positive	4.12	5.12			1) 3-Trifluoromethylphenylpiperazine
197.08400				Positive	3.47	4.47			1) 4-Chlorophenylpiperazine
193.13354				Positive					1) 4-Methoxyphenylpiperazine MeOPP
545.23812				Positive					1) Deacetylbacatin III
255.11280				Positive	4.44	5.44			1) Hydroxycarbapazine
331.22677				Positive	8.39	9.39			1) Hydroxy-delta-9-THC
209.08001				Negative	2.58	3.58			1) 3,7-Timethylnic acid
195.08765				Positive	2.15	3.15			1) 3,5-Timethylaniline
144.10191				Positive	0.45	1.45			1) Anisocyclohexanecarboxylic acid
177.13863				Positive	0.30	1.30			1) Methyl-3phenylpiperazine
194.11796				Positive	3.25	4.25			1) Methylamino-1-(3,4-methylenedioxyphenyl)piperazine
363.21660				Positive	4.86	5.86			2) Alpha-dihydrocodeine
365.23225				Positive	5.04	6.04			2) Beta-dihydrocodeine
363.21660				Positive	5.00	6.00			2) Beta-dihydrocodeine
420.21195				Positive	6.50	7.50			2) Theacoptin difluoracetate

Export compound database to inclusion list



Export to compound database

Ordering information

Tox Explorer Collection for Orbitrap Exploris 120 mass spectrometer	
Catalog number	WFLOW-62002
Type	Tox Explorer Collection
Mass spectrometer interface	Orbitrap Exploris 120 mass spectrometer (Sold separately)*
Liquid chromatography system	Vanquish Flex Binary UHPLC system (Sold separately)*
Methods	LC-MS/MS w/RT (HRAM tubing config.)
Software	TraceFinder 5.1 software
Description	Tox Explorer Collection for Orbitrap Exploris 120 mass spectrometer
For use with (application)	High resolution QUALITATIVE and QUANTITATIVE performance in: Forensic toxicology, clinical research toxicology, employee drug testing, and anti-doping applications
Includes	Standardized LC-MS/MS method, (U)HPLC column, experimentally obtained database with retention times, and LC-MS/MS instrument set-up and Tox Explorer application training by dedicated experts
Unit size	Each
Tox Explorer Collection for TSQ Quantis Plus mass spectrometer	
Catalog number	WFLOW-62003
Type	Tox Explorer Collection
Mass spectrometer interface	TSQ Quantis Plus mass spectrometer (Sold separately)*
Liquid chromatography system	Vanquish Flex Binary UHPLC system (Sold separately)*
Methods	LC-MS/MS w/RT (QQQ tubing config.)
Software	TraceFinder 5.1 software
Description	Tox Explorer Collection for TSQ Quantis Plus mass spectrometer
For use with (application)	Accessible QUANTITATIVE performance in: Forensic toxicology, clinical research toxicology, employee drug testing, and anti-doping applications
Includes	Standardized LC-MS/MS method, (U)HPLC column, experimentally obtained database with retention times, and LC-MS/MS instrument set-up and Tox Explorer application training by dedicated experts
Unit size	Each

* The Orbitrap Exploris 120 (HRAM) or TSQ Quantis Plus (triple quadrupole) mass spectrometers and Vanquish Flex Binary UHPLC system can be ordered separately in case you already own these instruments and are looking to purchase only the method, data package, and training components of Tox Explorer Collection.

 Learn more at thermofisher.com/ToxExplorer

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