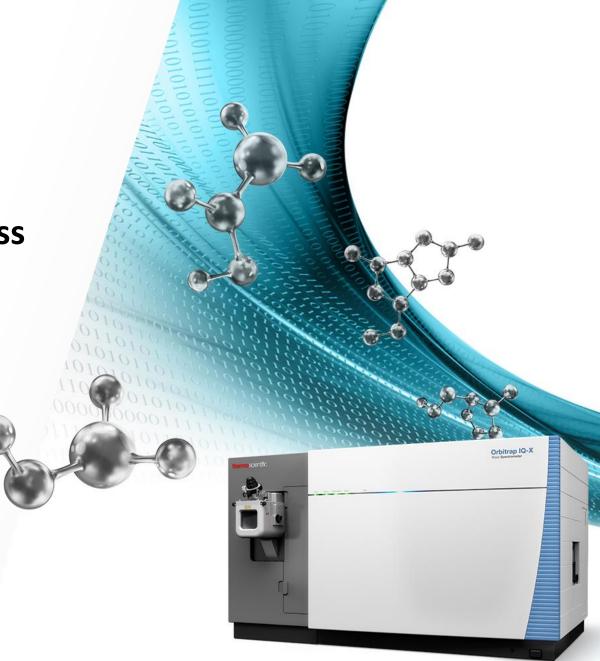
Thermo Fisher

High-Resolution Data-Dependent Mass Spectrometry Guided by Real-Time Library Search for the Detection and Characterization of PFAS

Brandon Bills Product Manager

Thu 7 8/3/2023



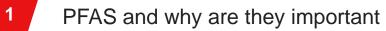
The world leader in serving science

Conflict of interest disclosure

The presenter and fellow authors are employees of Thermo Fisher Scientific whose instrumentation and software were used to acquire and process the data.

Outline

3



2 Analytical challenges for unknown PFAS

Mass filtering to focus analysis

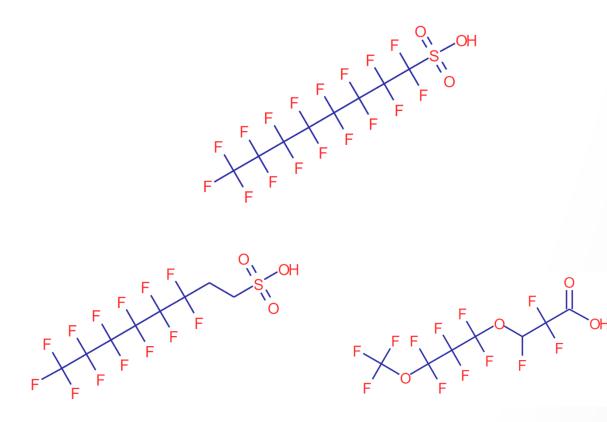
Real-Time Library Search for additional information

Data processing



PFAS

Per- and PolyFluoroAlkyl Substances contain fully fluorinated methyl or methylene carbon atoms





Proprietary & Confidential | brandon.bills@thermofisher.com | June-2023

Problem overview

- Ubiquitous PFAS have been found in water, soil, air, food, and biological samples
- Long Lasting PFAS can take years to degrade in the body
- ¹Health Impacts PFAS has been found to have links to:
 - Endocrine disruptions of the thyroid
 - Reduced immune system functionality
 - Low birth weights
 - Liver Toxicity
- ²Emerging Regulations March 2023 the EPA proposed a new regulation on six PFAS in drinking water

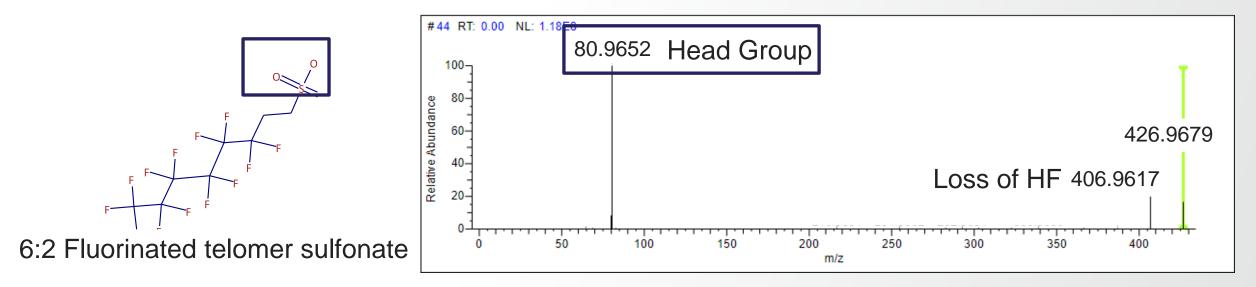
- 2. EPA. Per- and Polyfluoroalkyl Substances (PFAS) Proposed PFAS National Primary Drinking Water Regulation. March 2023
- 5 Proprietary & Confidential | brandon.bills@thermofisher.com | June-2023



^{1.} Agency for Toxic Substances and Disease Registry. Toxicological Profile for Perfluoroalkyls. May 2021

Analytical Challenges

- Often present at lower concentrations in complex matrices
- Over 12,000 PFAS compounds known
- Minimal commercial standards available
- Collision based MS² doesn't always give enough information to piece together the structure



Thermo Scientific[™] Orbitrap[™] IQ-X[™] Tribrid[™] mass spectrometer

- Orbitrap resolution up to 500K standard or 1M with upgrade
- Ion Trap for on-the-fly CE optimization as well as CID and MSⁿ fragmentation
- Highly customizable method editor to tailor the method to the analytes of interest
- Real-Time Library Search to selectively target compounds in real time using a spectral library



Customizable methods focuses analysis on PFAS



HRAM data helps differentiate isobars

A mass filter to prioritize compounds with a negative mass defect

Assisted CE optimizes settings on the fly

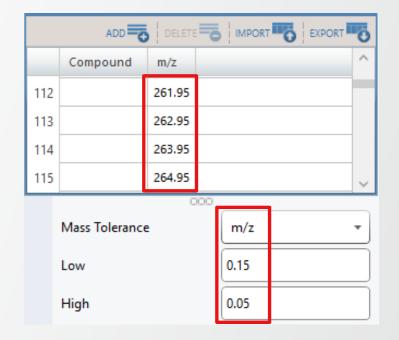
RTLS checks against a library in real time then triggers scans on likely PFAS

ddMS³ and CID fragmentation gets additional structural information on unknown PFAS compounds

A mass filter to prioritize a negative mass defect

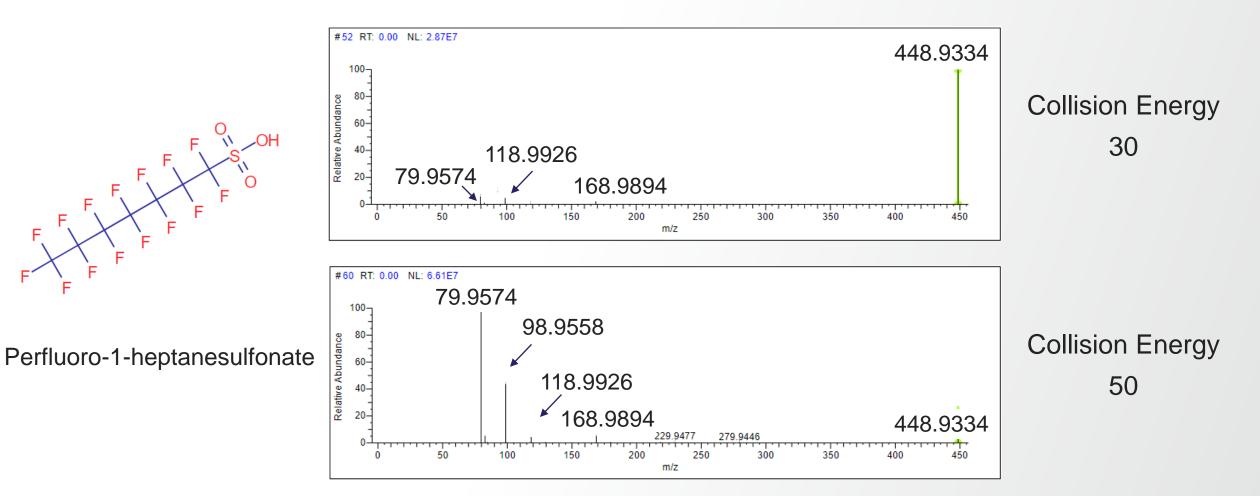
- Highly fluorinated compounds tend to have m/z below the integer values
- An inclusion list focuses ddMS² on those mass ranges during the analysis

PFAS compound	m/z					
Perfluorohexanoic acid	312 <mark>.9728</mark>					
Perfluoro-1-butanesulfonamide	297.9590					
HFPO-DA	328.9677					
6:2 Fluorinated telomer sulfonate	426.9679					
N-MePFOSAA	569.9673					



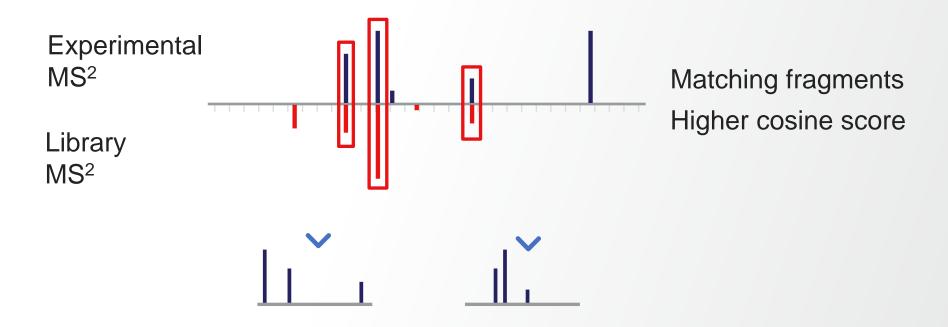
Assisted CE

- The ion trap can check up to 5 collision energies in parallel with Orbitrap acquisition
- The CE that fragments at least 80% of the precursor is used for the ddMS² acquisition



Real-Time Library Search

- Real-Time Library Search
 - ddMS² data is compared to a library in real time to find similar spectral features and generate a cosine score



Likely PFAS trigger additional experiments like ddMS³ or CID

Bills et al. Anal. Chem. **2022,** 94, 9, 3749–3755

Experimental details

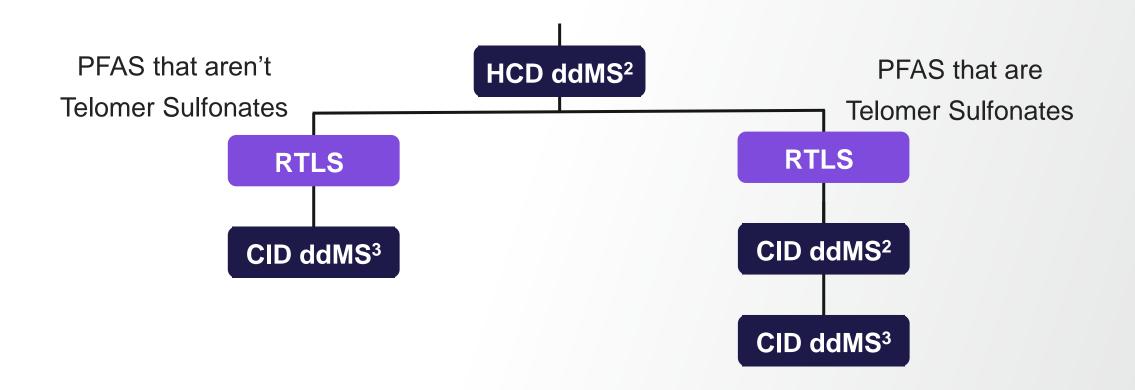
 A mixture of 30 PFAS compounds (PFAC30PAR from Wellington Laboratories) were run both diluted in solvent at 10 ppb and spiked into plasma extract at 1 ppb

Material	Details						
LC	Thermo Scientific [™] Vanquish [™] Flex UHPLC system with PFAS HPLC Kit						
Column	Thermo Scientific™ Accucore™ C18 HPLC Column						
Solvent A	100% water + 10 mM ammonium acetate						
Solvent B	75:20:5 MeOH : ACN : Water + 10mM Ammonium acetate						
Sample Vials	Polypropylene Vials and caps						



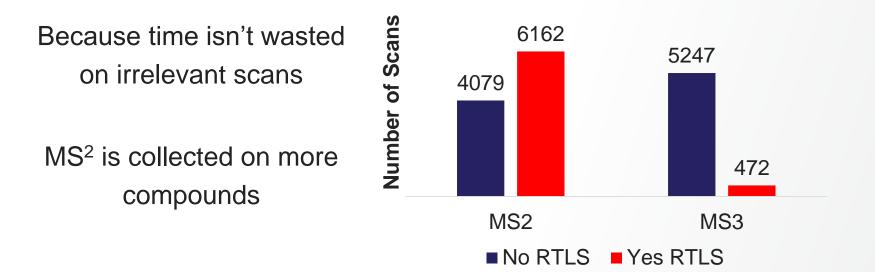
RTLS Method

- A library was curated in mzVault without 7 of the compounds
- A method was created using a mass defect filter and Real-Time Library Search in similarity mode



More relevant scans with RTLS

- Without guidance, ddMS³ is collected on every compound regardless of relevance
- By only triggering compounds on likely PFAS, ddMS³ is limited to relevant compounds
- For example, look at how the instrument spends its time when analyzing a single injection of plasma spiked with a mixture of 30 PFAS compounds

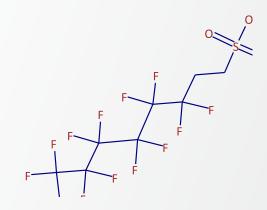


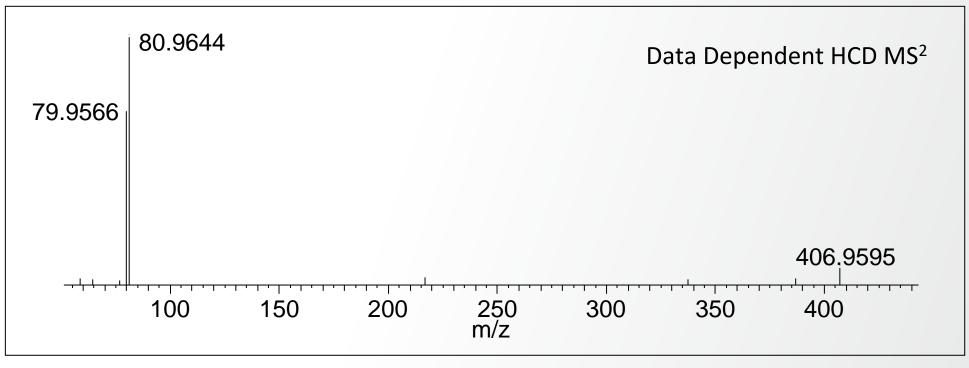
Instead of thousands of MS³ collected on non-PFAS

there is an easier to review data set with MS³ collected only on likely PFAS

Data collected on Standards

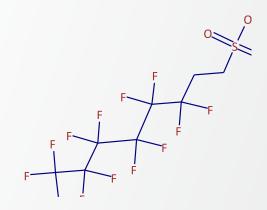
- All 30 standards in neat solvent were detectable and triggered ddMS² and 29 triggered ddMS³ data
- CID MS²/MS³ was collected on 6:2 Fluorinated telomer sulfonate despite not being in the library

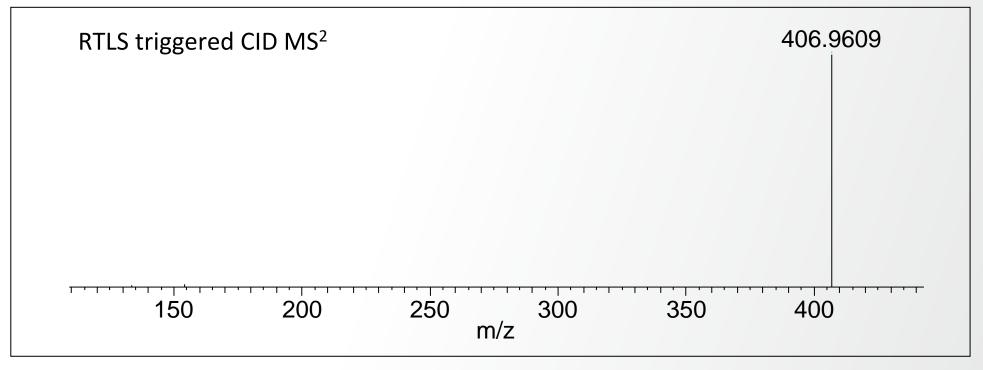




Data collected on Standards

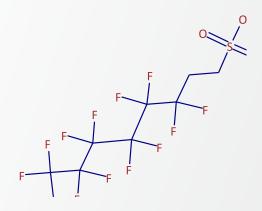
- All 30 standards in neat solvent were detectable and triggered ddMS² and 29 triggered ddMS³ data
- CID MS²/MS³ was collected on 6:2 Fluorinated telomer sulfonate despite not being in the library

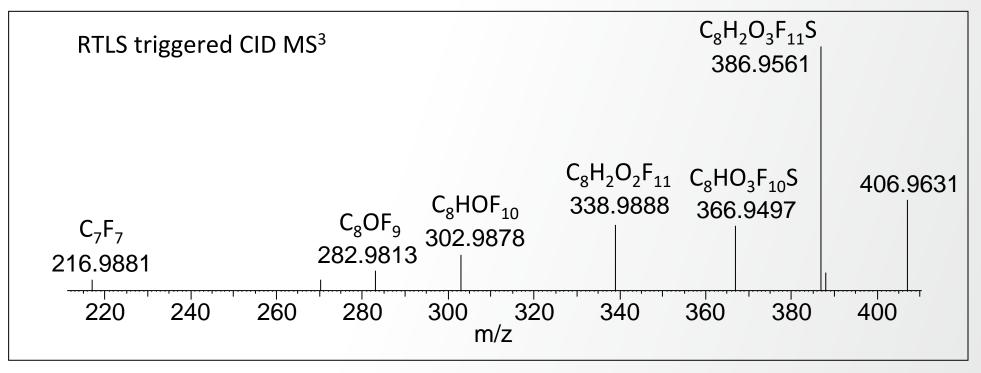




Data collected on Standards

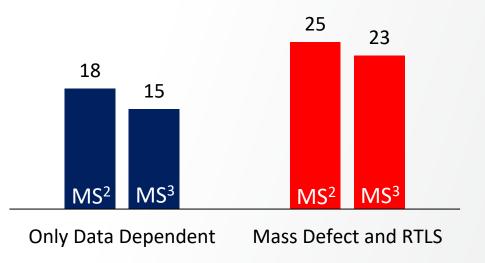
- All 30 standards in neat solvent were detectable and triggered ddMS² and 29 triggered ddMS³ data
- CID MS²/MS³ was collected on 6:2 Fluorinated telomer sulfonate despite not being in the library





Advantages of Mass Defect Filter and RTLS

- In the plasma sample, unguided ddMS³ often targets higher mass matrix compounds
- With a filter for mass defect, ddMS² is limited to compounds that are potential PFAS
- RTLS limited ddMS³ data collection to compounds similar to PFAS in the given library

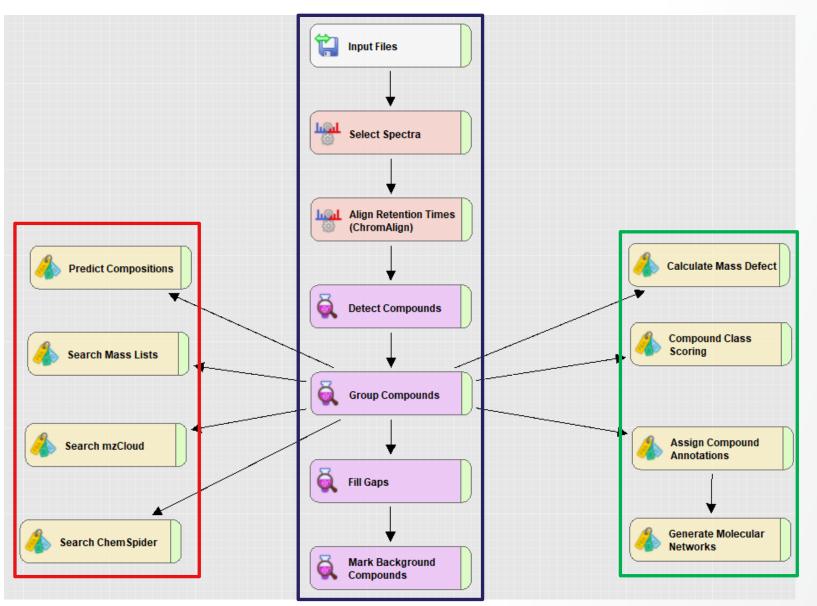


Standards with spectral data

Data Processing in Compound Discoverer



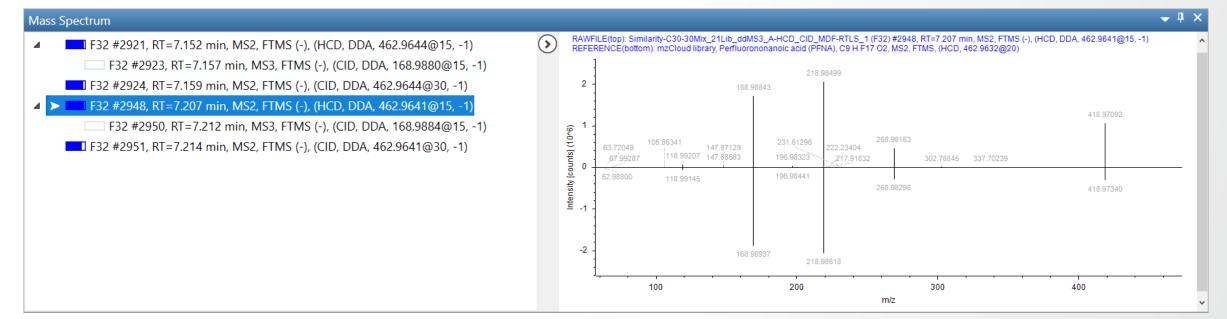
Thermo Scientific[™] Compound Discoverer[™]



- Takes in files, detects compounds, aligns between files, marks background
- Predicts the composition and searches for annotations from multiple sources
- Calculates mass defect, scores their similarity to PFAS, and generates molecular networks

Post-Processing Nodes	
Scripting Node	

Compound Discoverer results



Annotations from mzCloud can be directly compared against experimental results

F	Name	Formula	Annot. Source 🛨	Annot. ∆Mass [ppm]	Calc. MW	m/z 🔺	RT [min]	Area (Max.)	mzCloud Best Match	Mass	Defect	+	Class Cov	verage 🛨	MS Depth
1 中	Perfluorobutanoic acid	C4 H F7 O2		0.42	213.98657	212.97928	3.652	34884364	86.3	- 0.013	-63	0	0.25	6.25	3
2 📼	Perfluoropentanoic acid	C5 H F9 O2		0.54	263.98343	262.97613	4.234	46425737	87.9	-0.017	-63	0	0.25	12.50	3
3 📼	Perfluoro-1-butanesulfonamide	C4 H2 F9 N O2 S		-0.02	298.96625	297.95897	5.474	131675699	90.8	-0.034	-113	-0.015	0.25	12.50	3
4 -⊨	Perfluoro-1-butanesulfonic acid (PFBS)	C4 H F9 O3 S		-0.20	299.95021	298.94293	4.446	123178856	88.3	-0.05	-166	-0.031	0.62	25.00	3
5 ⊨	Perfluorohexanoic acid	C6 H F11 O2		1.19	313.98046	312.97317	4.896	51681327	88.9	-0.02	-62	0.001	0.25	0.00	3

• Results are displayed in a table including annotation, formula, mass defect, and PFAS class coverage

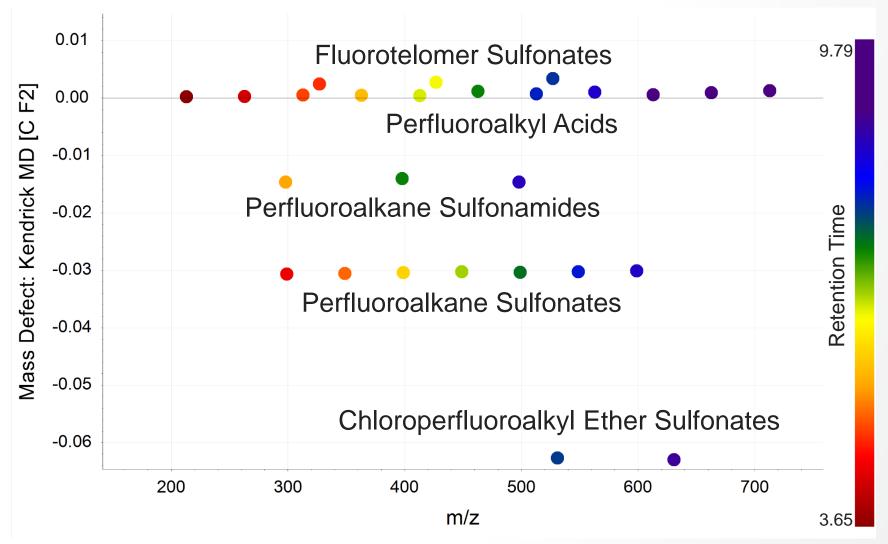
Kendrick Mass Defect

- Kendrick mass normalizes the mass based on mass of a repeating subunit (CF₂ in this case)
- Plotting compounds by Kendrick mass defect and molecular weight shows homologous series

Kendrick Mass = Observed Mass $\times \frac{50.0000}{49.9968}$

Kendrick Mass Defect = Nominal Mass - Kendrick Mass

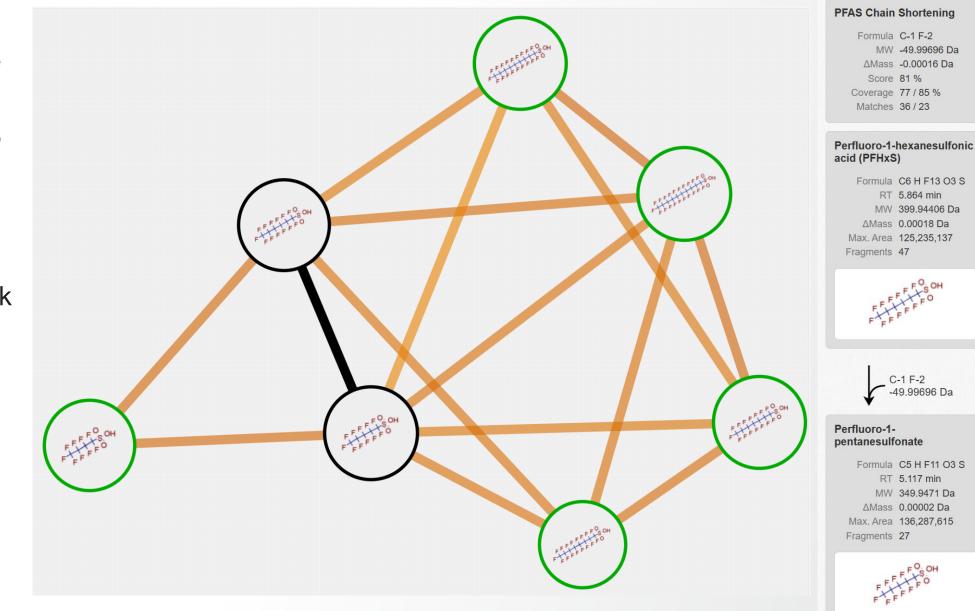
Homologous Series



- 5 homologous series PFAS were graphed
- Compounds in a series have the same Kendrick mass defect and are spaced in increments of 50 amu
- The RT is color coded as a third dimension to provide a simple verification of increasing RT based on PFAS chain length.

Molecular networks for homologous series

- Compounds are linked based on their relations to other compounds
- Clicking on a link displays information about both compounds and how they are related



Conclusions

- PFAS is a complex compound class that poses numerous analytical challenges
- The Orbitrap IQ-X offers tools such as Real-Time Library Search to focus data acquisition on relevant compounds
- ddMS³ and CID provides unique information to help identify unknown structures
- Compound Discoverer helps process and visualize PFAS results



Acknowledgments



Thermo Colleagues Sunandini Yedla Juan Sanchez Tim Stratton Ralf Tautenhahn

Ed George Cynthia Grim Vlad Zabrouskov

Thank you

I would be happy to answer questions

