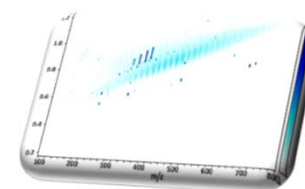
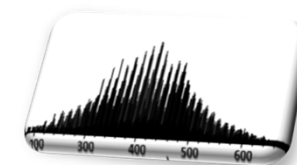
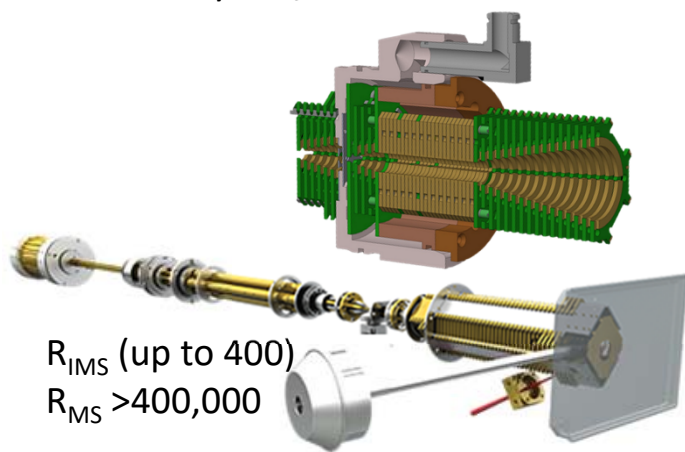


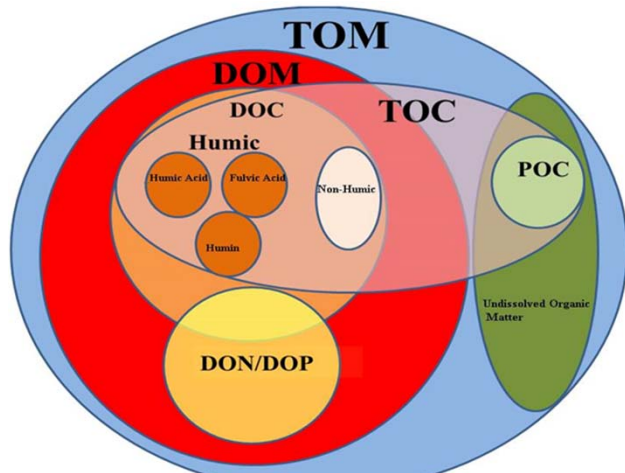
Unraveling the Structural Complexity and Diversity of Dissolved Organic Matter using TIMS-FT-ICR MS

Dennys Leyva, Lilian V. Tose, Jacob Porter, Jeremy Wolff, Rudolf Jaffé and Francisco Fernandez-Lima



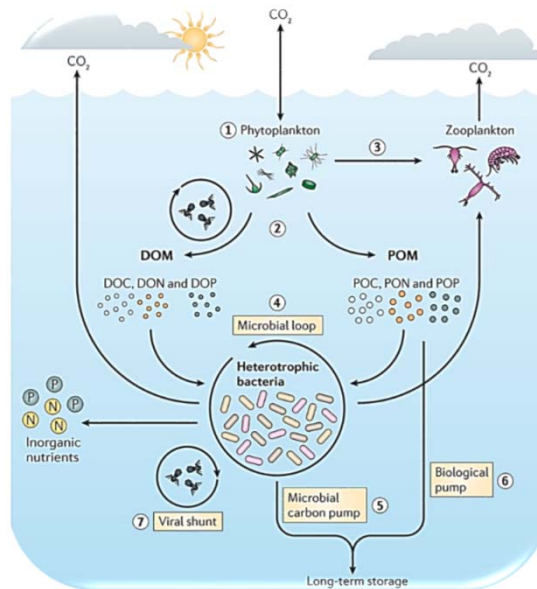
Motivation

Dissolved Organic Matter (DOM): Complex mixture from bacteria, algal and high plant organic matter degradation.

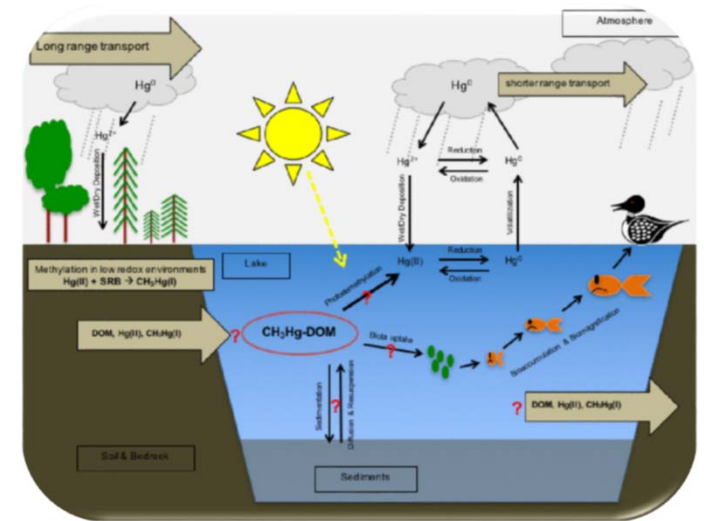


T. Pagano, M. Bida and J. E. Kenny, *Water*, 2014, 6, 2862.

Role in aquatic environments



Nature Reviews Microbiology 12, 686-698 (2014)

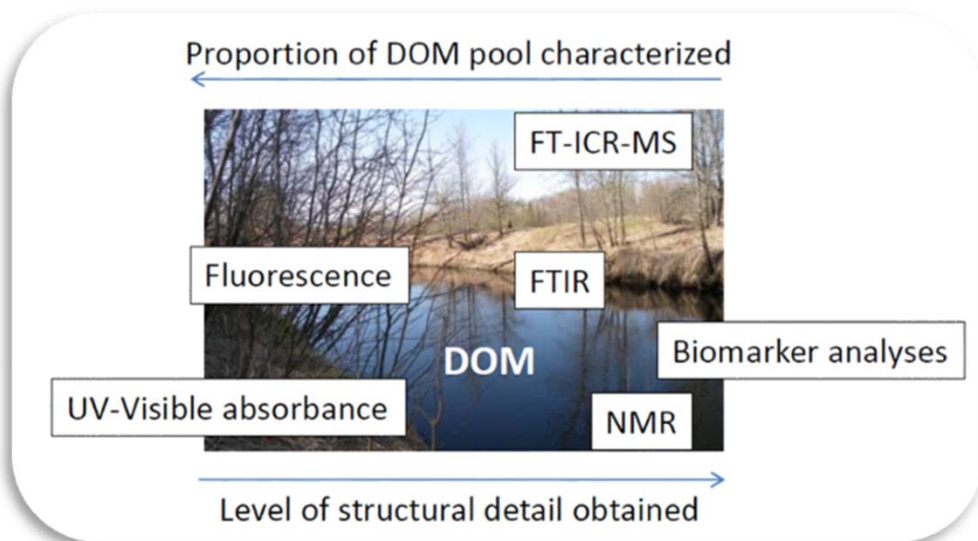


Bulletin of Environmental Contamination and Toxicology (2018) 100:14–25

A full understanding of DOM impact in climate change, ecology, and toxicology, requires a comprehensive knowledge of their molecular composition and structure.

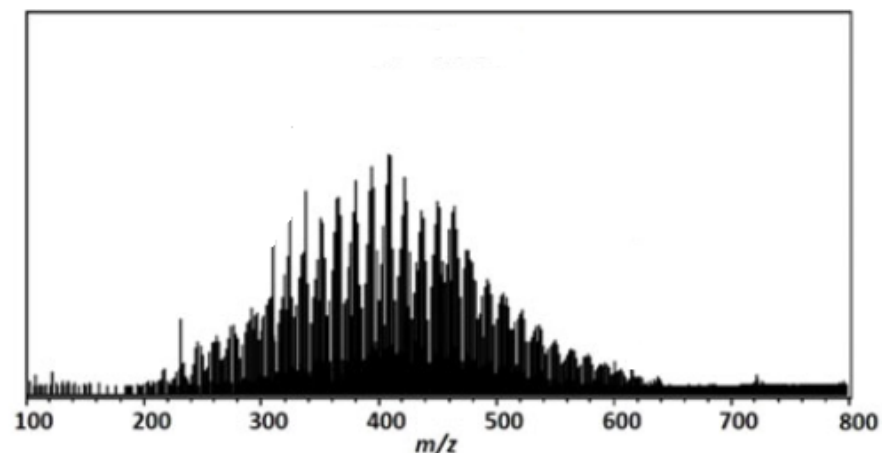
Analytical approaches

Bulk vs molecular level characterization

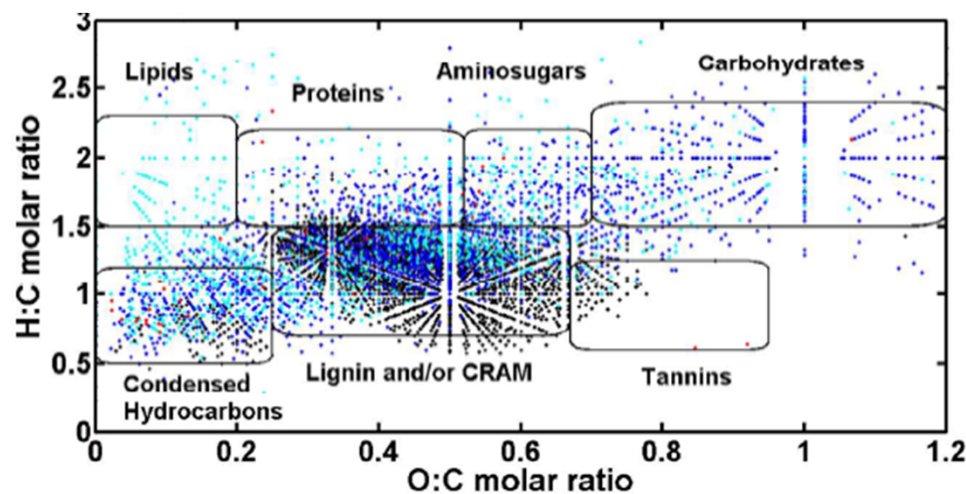


Challenges in DOM structure elucidation

- High structural heterogeneity
- Wide range of molecular weights
- **Isomeric diversity**



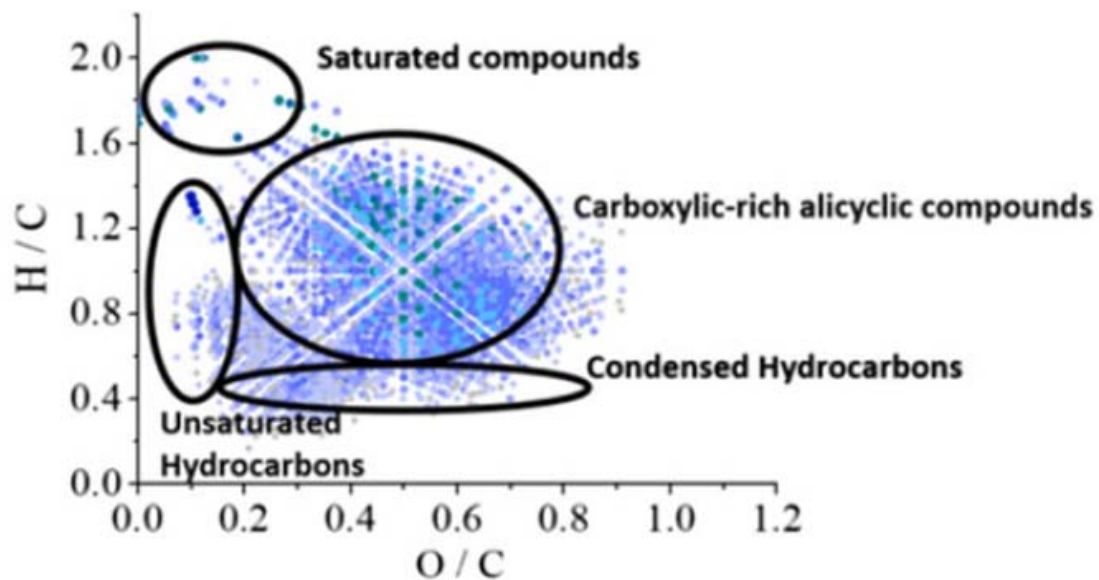
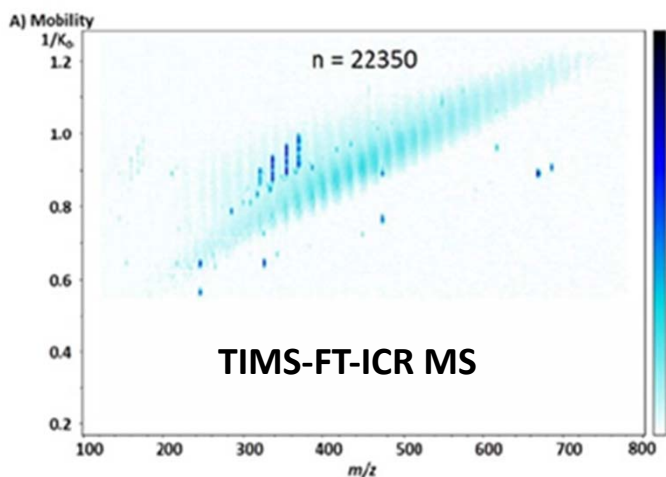
DOM MS spectrum



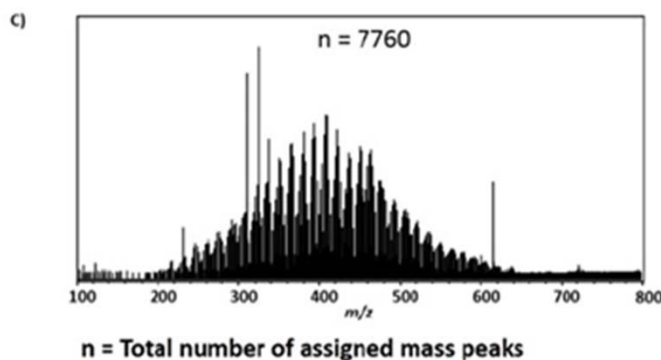
Diversity in molecular classes

Coupling TIMS with FT-ICR MS in SRFA analysis

How can TIMS help to understand the complexity of DOM?

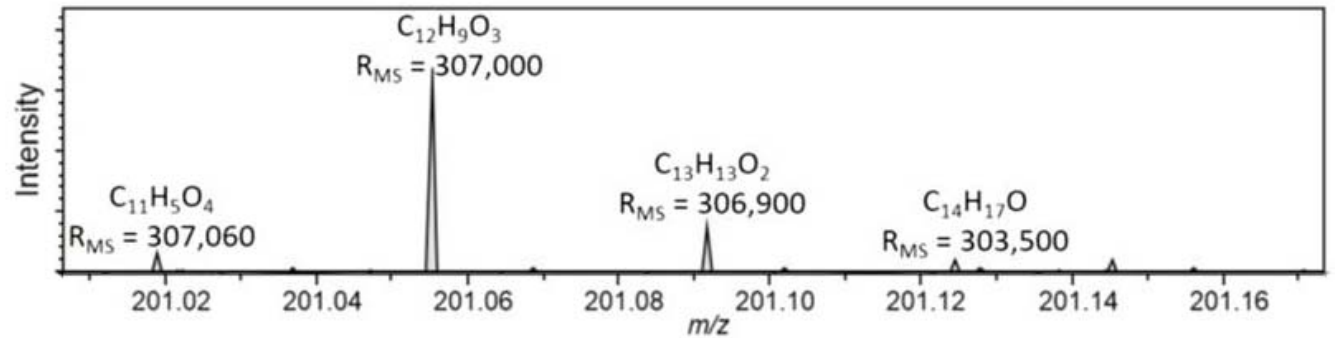
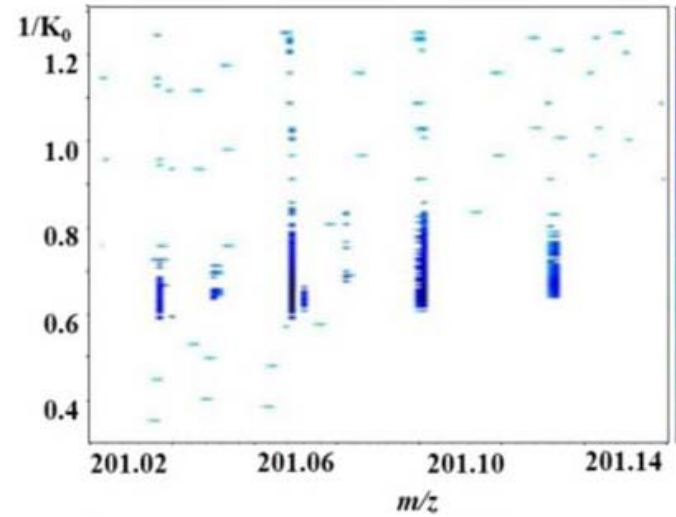
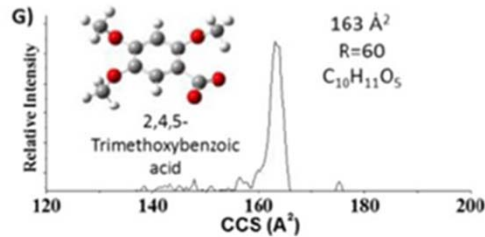
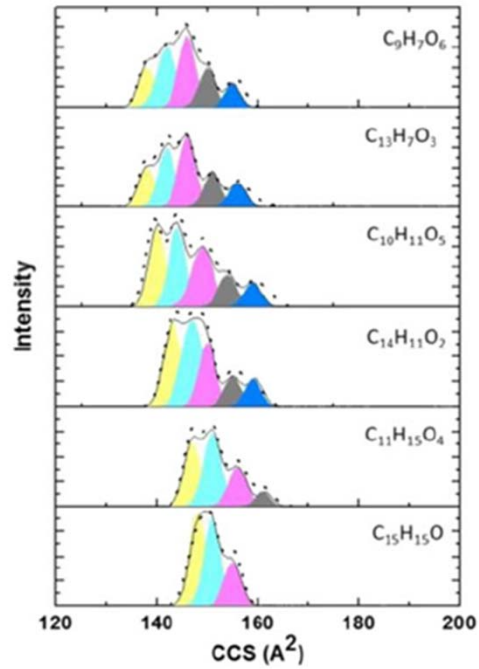


3050 chemical formulas assigned in SRFA standard



Isomeric content in SRFA standard

E)



IMS-MS (top) and MS (bottom) projections from the SRFA analysis at m/z 201

We provided, for the first time, a lower cutoff estimation of the number of molecular isomers in the SRFA standard.

Experimental

Samples



Pantanal National Park, Brazil

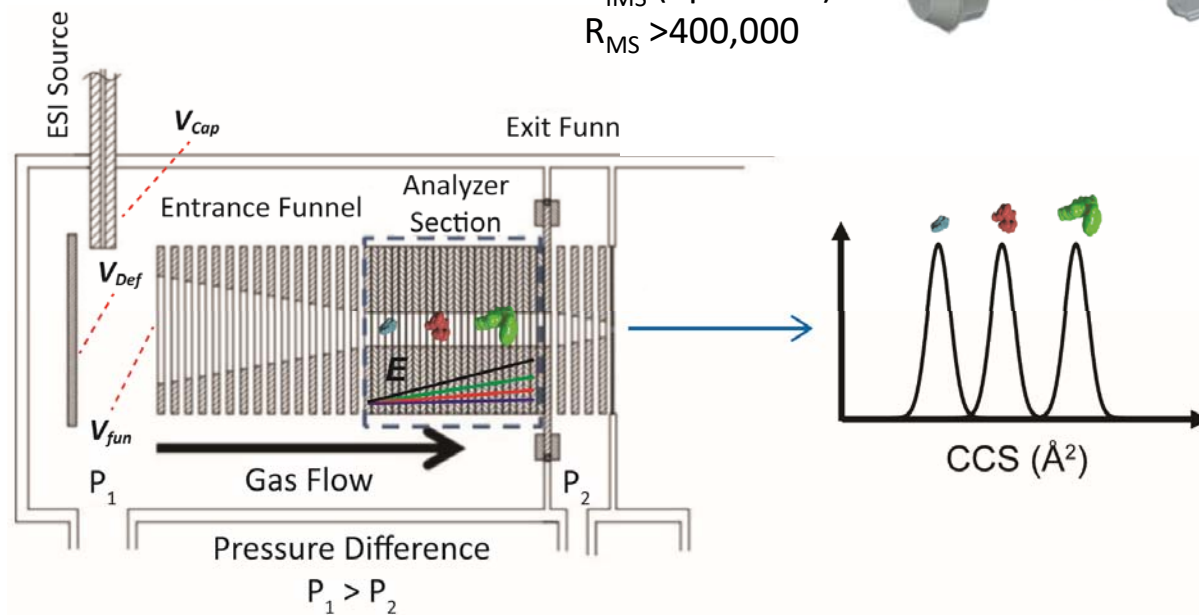
TIMS-FT-ICR MS 7T Solarix



Trapped Ion Mobility Spectrometry



R_{IMS} (up to 400)
 $R_{MS} > 400,000$



$$k = \frac{v_g}{E} = \frac{A}{(v_{elution} - v_{out})}$$

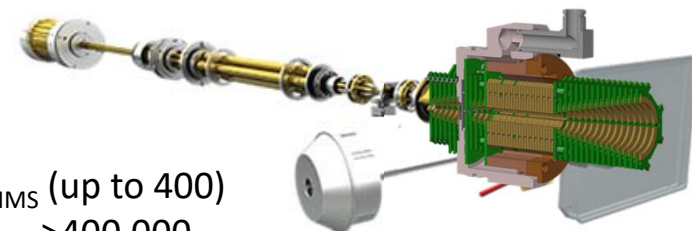
- Nonlinear stepping scan function
- Mobility spectra calibrated using Tuning Mix

FT-ICR MS/MS: quadrupole isolation-CID 15-20 eV.

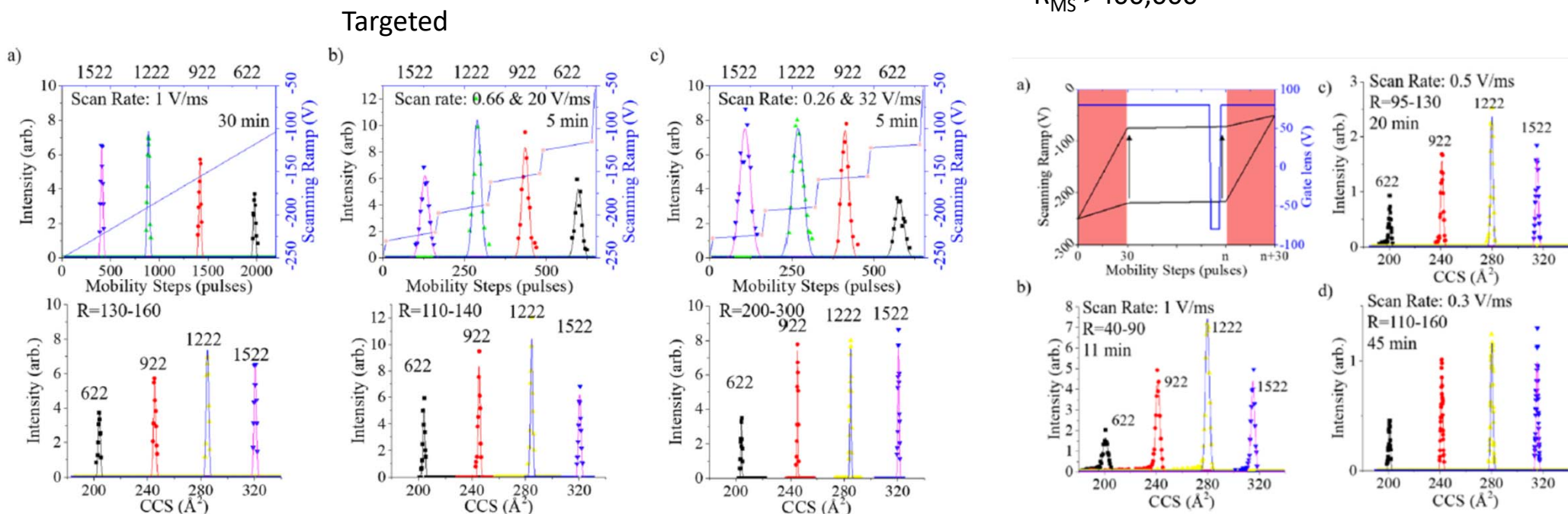
Experimental

Trapped Ion Mobility Spectrometry

Non linear scan functions TIMS-FT-ICR MS



R_{IMS} (up to 400)
 $R_{MS} > 400,000$



P. Benigni; J. Porter; M. Ridgeway; M. Park; F. Fernandez-Lima*. "Increasing analytical separation and duty cycle with non-linear analytical mobility scan functions in TIMS-FT-ICR MS". *Anal Chem.* 2018, 90 (4), 2446–2450.

Experimental

Samples



Pantanal National Park, Brazil

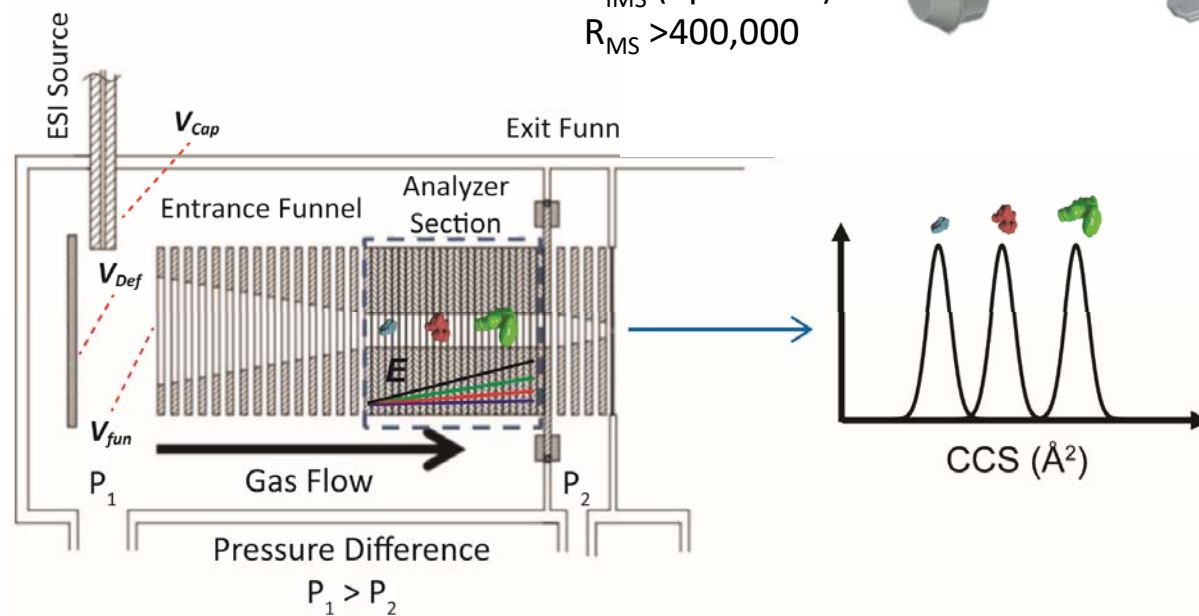
TIMS-FT-ICR MS 7T Solarix



Trapped Ion Mobility Spectrometry



R_{IMS} (up to 400)
 $R_{MS} > 400,000$



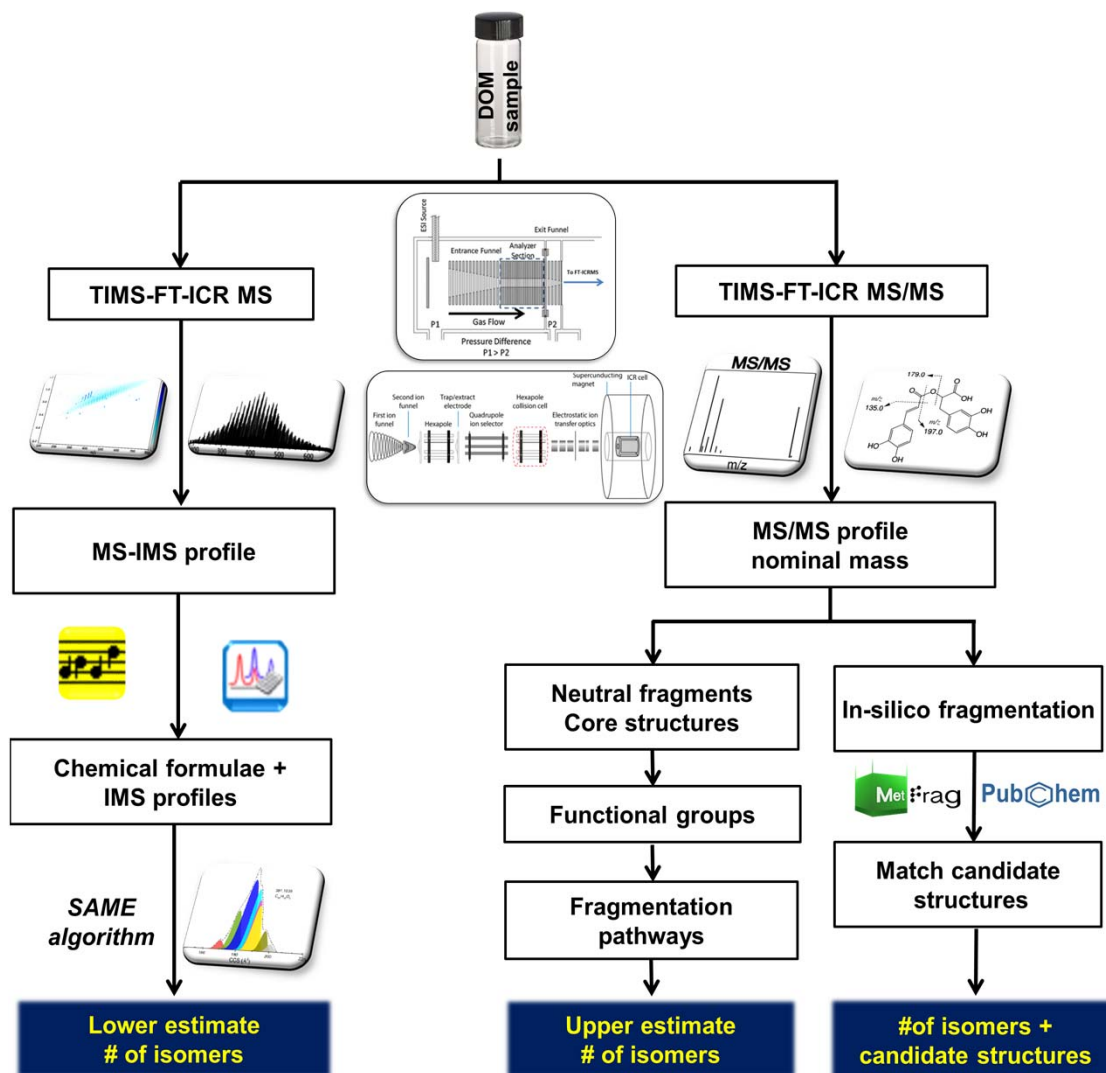
$$k = \frac{v_g}{E} = \frac{A}{(v_{elution} - v_{out})}$$

- Nonlinear stepping scan function
- Mobility spectra calibrated using Tuning Mix

FT-ICR MS/MS: quadrupole isolation-CID 15-20 eV.

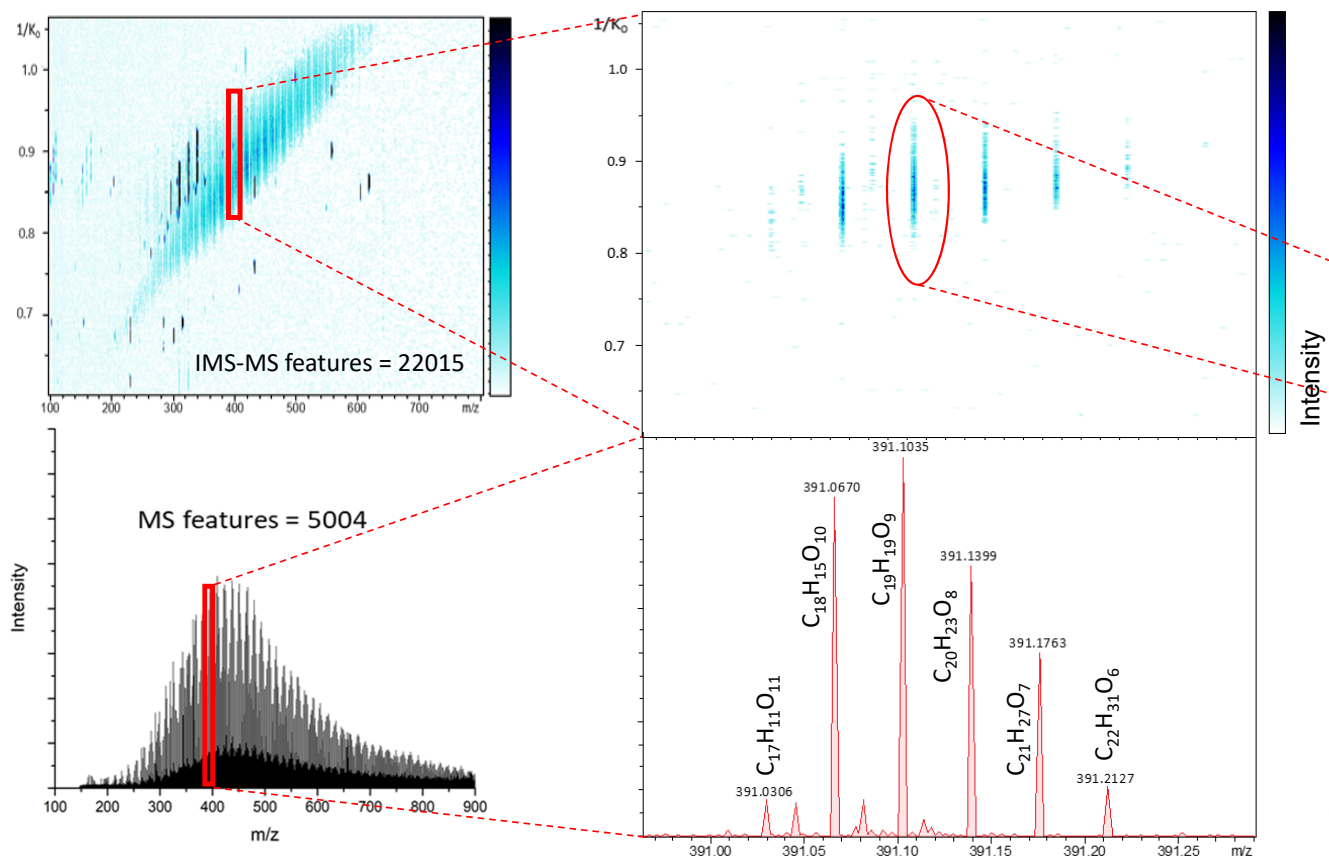
Analytical workflow

How can TIMS-FT-ICR MS help to understand the isomeric diversity of DOM?



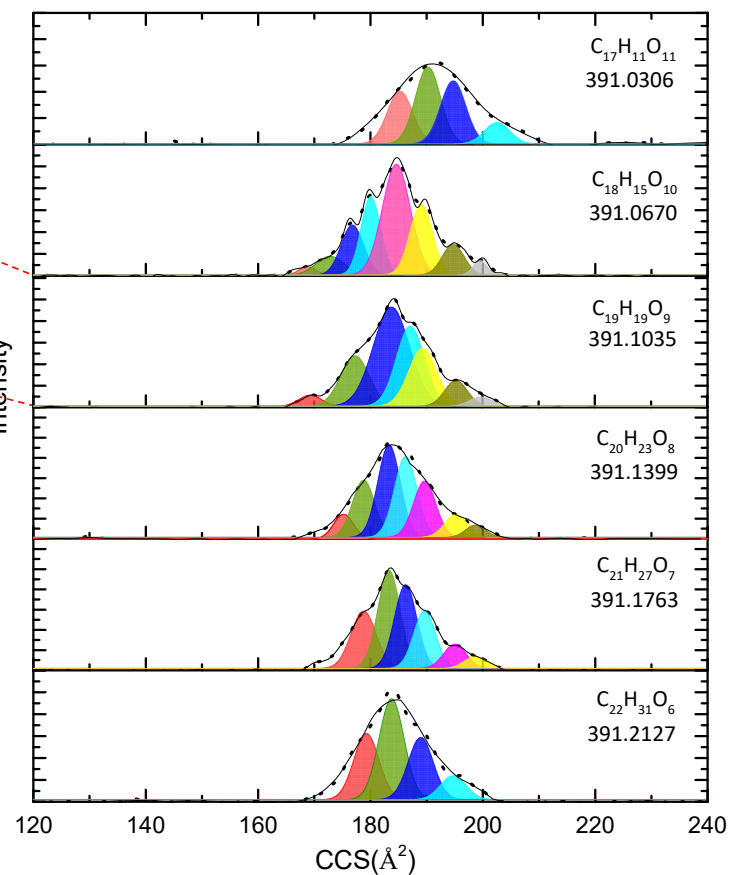
TIMS-FT-ICR MS

More than 3,000 chemical components identified



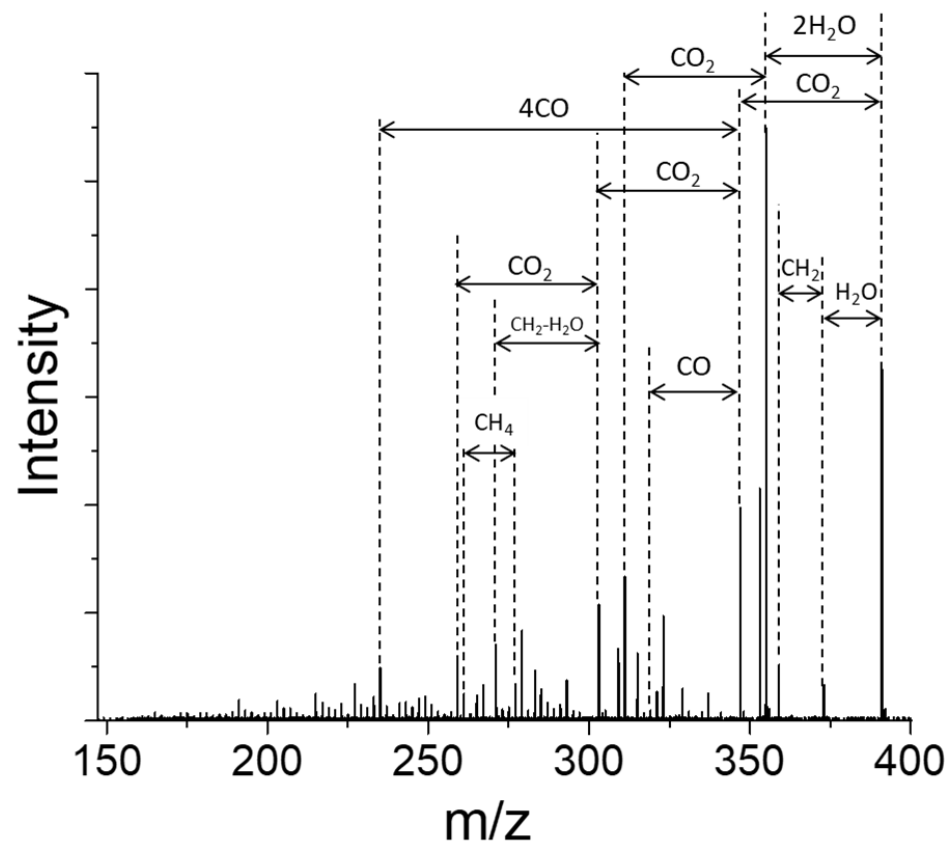
2D IMS-MS and MS projection at m/z 391

6-10 isomers per chemical formula



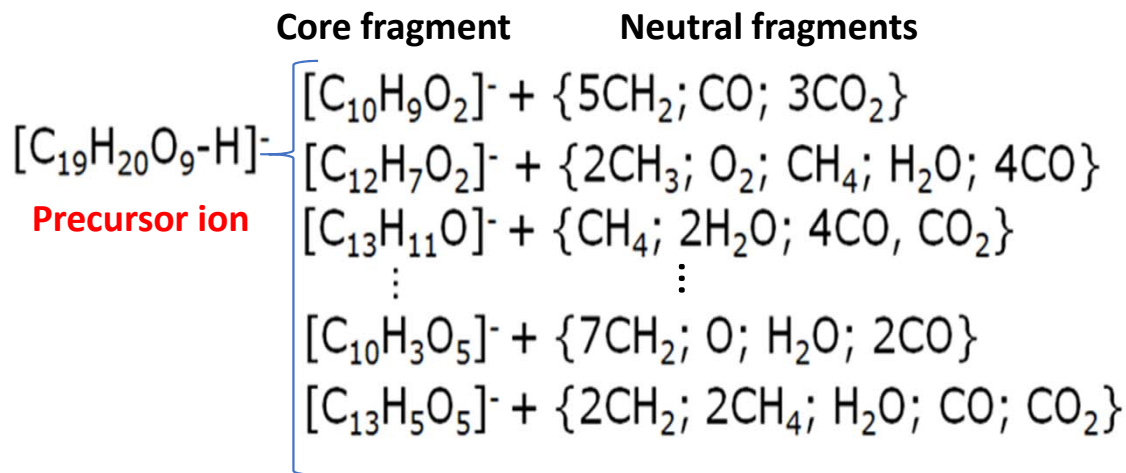
IMS projection at nominal mass m/z 391

FT-ICR MS/MS



FT-ICR MS/MS (q-CID) profile of precursor ion 391 m/z

- Neutral losses associated with functional groups and precursor ion.
- The number of pathways could provide an estimate of the number of structural isomers.



FT-ICR MS/MS (q-CID) of precursor ion 391 m/z

Potential neutral losses							Core fragment m/z	Number of pathways
CH ₂	CH ₃	O	CH ₄	H ₂ O	CO	CO ₂		
5	-	1	-	-	-	3	173.0607 C ₁₁ H ₉ O ₂	23
2	-	-	1	1	4	1		
3	-	-	1	-	1	3		
1	2	-	-	1	4	1		
2	2	-	-	-	1	3		
4	-	-	-	1	2	2		
4	-	2	-	1	4	-		
-	-	-	2	1	6	-		
5	-	2	-	-	1	2		
-	2	1	1	-	4	1		
1	-	1	2	-	4	1		
5	-	3	-	-	2	1		
5	-	4	-	-	3	-		
1	-	-	2	-	3	2		
2	2	1	-	-	2	2		
3	-	1	1	-	2	2		
3	-	2	1	-	3	1		
2	2	2	-	-	3	1		
1	2	1	-	1	5	-		
2	-	1	1	1	5	-		
-	2	-	1	-	3	2		
3	-	-	-	2	5	-		
4	-	1	-	1	3	1		

Precursor ion m/z	Core Fragment m/z	Structural isomers
	161.0607 C ₁₀ H ₉ O ₂	13
	163.0763 C ₁₀ H ₁₁ O ₂	7
	165.0192 C ₈ H ₅ O ₄	3
	165.056 C ₉ H ₇ O ₃	2
	167.0349 C ₈ H ₇ O ₄	1
	171.0814 C ₁₂ H ₁₁ O	23
	173.0607 C₁₁H₉O₂	23
391.1031 C ₁₃ H ₁₉ O ₃	175.0400 C ₁₀ H ₇ O ₃	15
	183.0450 C ₁₂ H ₇ O ₂	40
	183.0814 C ₁₃ H ₁₁ O	25
	185.0607 C ₁₂ H ₉ O ₂	29
	187.0400 C ₁₁ H ₇ O ₃	25
	201.0192 C ₁₂ H ₅ O ₄	25
	202.9984 C ₁₀ H ₁₃ O ₅	15
	205.0140 C ₁₀ H ₅ O ₅	7
	241.0140 C ₁₁ H ₅ O ₅	7

Rapid upper estimate of 260 structural isomers

In-silico fragmentation



MetFrag

In silico fragmentation for computer assisted identification of metabolite mass spectra



Candidate retrieval finished
Got 128 candidates

Database Settings

Database: PubChem

include references:

Parent Ion: 391.1035

[M-H]⁻

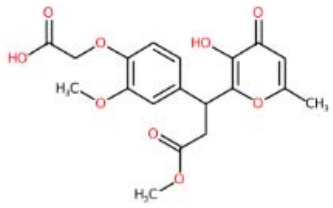
Calculate

Neutral Mass: 392.11079

Search ppm: 5

Formula: C₁₉H₂₀O₉

128 potential candidates found

#	Molecule	Identifier	Mass	Formula	FinalScore	Details
1	 2-[4-[(1R)-1-(3-hydroxy-6-methyl-4-oxo-pyran-2-yl)-3-methoxy-3-oxo-propyl]-2-methoxy-phenoxy]acetic acid	97424766 97424765 71826391 InChIKeyBlock1 = <u>HQBLWEZWLSOZAE</u>	392.111	C ₁₉ H ₂₀ O ₉	1.0	Peaks: 158 / 304 Fragments Scores Download

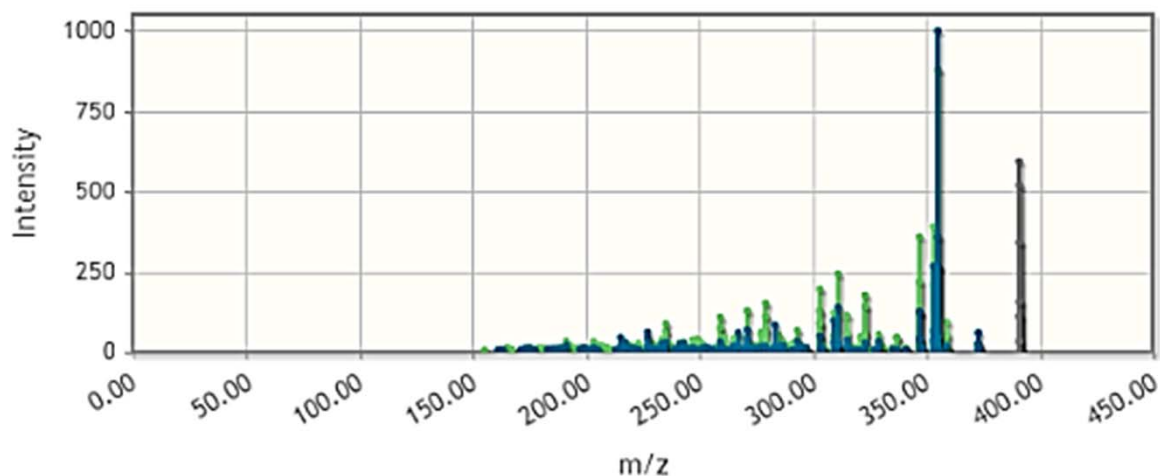
Journal of Cheminformatics, 2016, 8:3

In-silico fragmentation

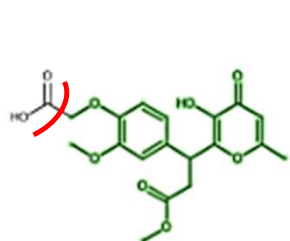
Fragments View

Select area to zoom in. Double click to return.
Click on apex of explained peak to select fragment.

■ matched
■ not matched
■ excluded



Fragments

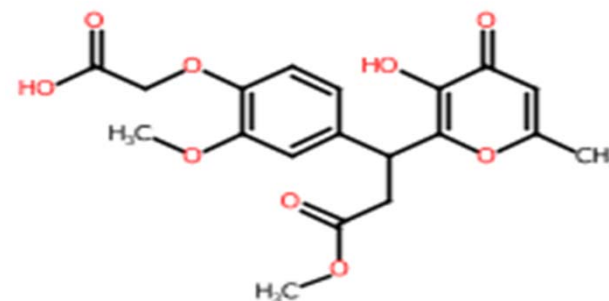


Fragment 153

Peak m/z: 347.113304
Fragment Mass: 347.11369 Da
Fragment Formula: $[C_{18}H_{19}O_7]^-$

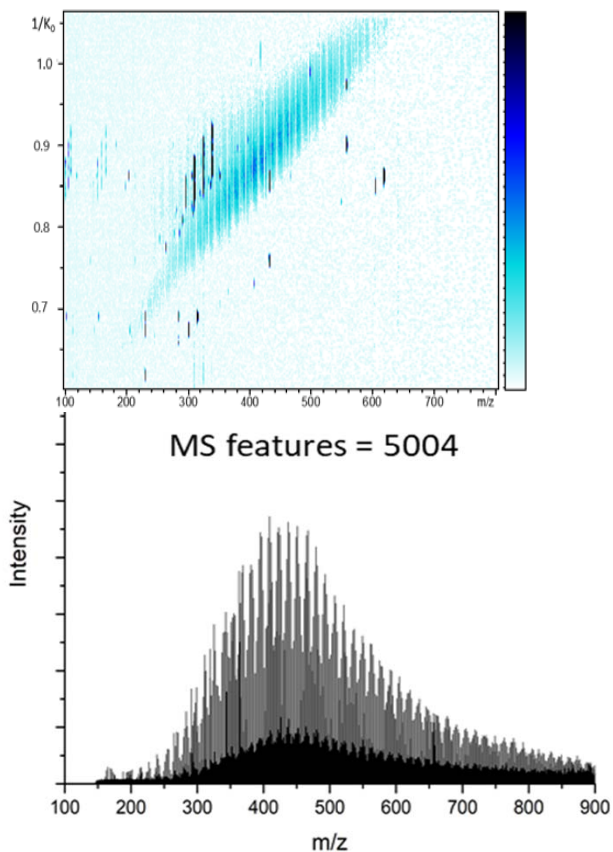
96 candidate structures based on accurate masses of precursor and fragments.

Candidate structure

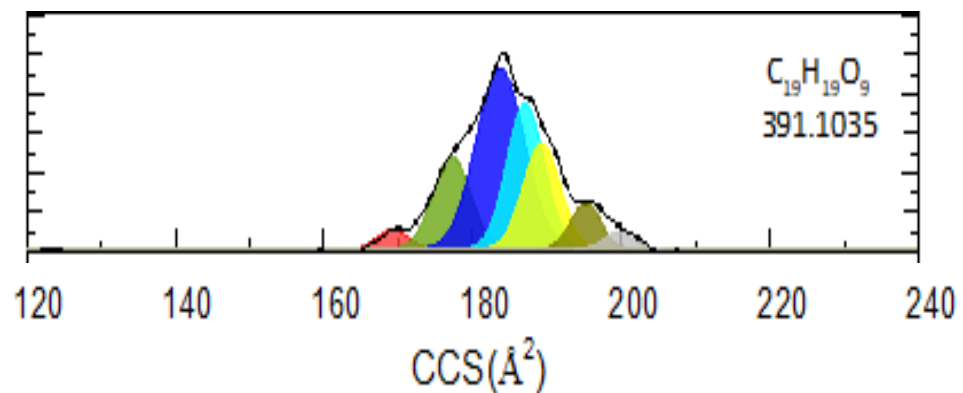


Conclusions

3,066 chemical components identified.



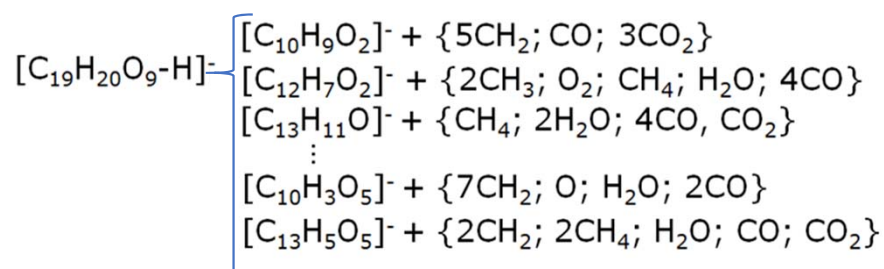
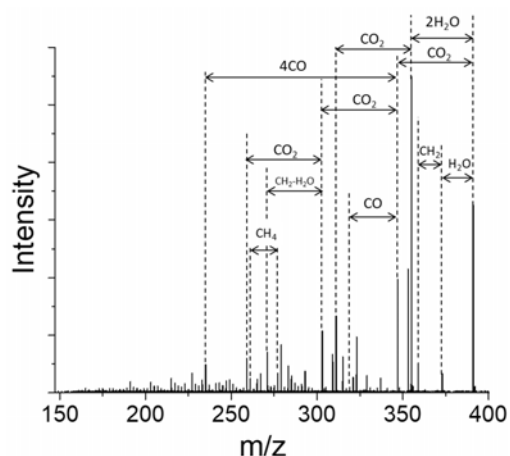
DOM Isomeric content per chemical formula based on TIMS.



Faraday Discussions, 2019, in press,

Conclusions

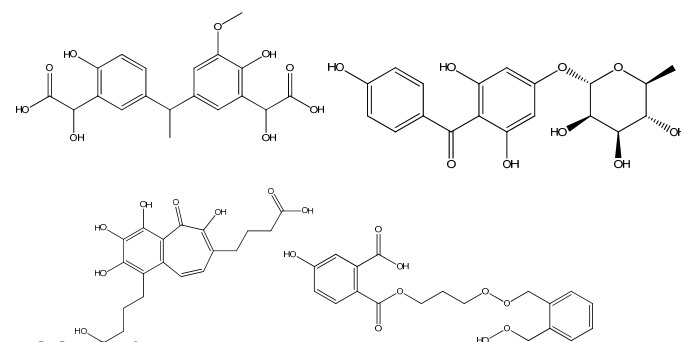
DOM isomeric content based on unique neutral loss fragmentation pathways/core fragments.



96 candidate structures from Metfrag-Pubchem



#	Molecule	Identifier	Mass	Formula	FinalScore	Details
9		58275149 InChIKeyBlock1 = YRPLVKYCIHGAGD	392.111	C ₁₉ H ₂₀ O ₉	0.8567	Peaks: 145 / 304 Fragments Scores Download



Faraday Discussions, 2019, in press,

Acknowledgements



NSF-CREST Program award HRD-1547798

NSF Division of Chemistry, CAREER award CHE-1654274



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Shirley Hernandez

