SWATH-MS, Ion Mobility and LC-MS for lipidomics

Jeevan Prasain jprasain@uab.edu 6-2612

1

SWATH-MS (Sequential Window Acquisition of all TheoreticalMass Spectra (in Triple-TOF system)

MSMS^{ALL}- No stone unturned

Data-independent workflow with a capability of acquiring high resolution MS/MS data for all detectable ions (*m/z* 200-1200) in a single run (6 min)

Strength of MS/MS^{ALL}

- Comprehensive, no requirement of a priori assumptions
- Combination of top-down and bottom-up methods
- Digital records of MS and MS/MS (high mass resolution)
- Data can be searched retrospectively
- Comparison (statistical) lipid changes between/among groups

Prasain et al. Metabolites, 2015

3

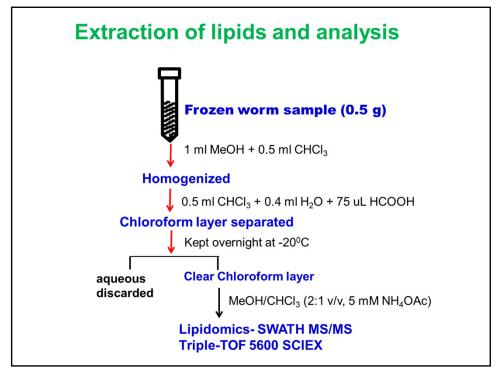
High speed, high resolution, sensitive detection and accuracy are crucial for lipid analysis

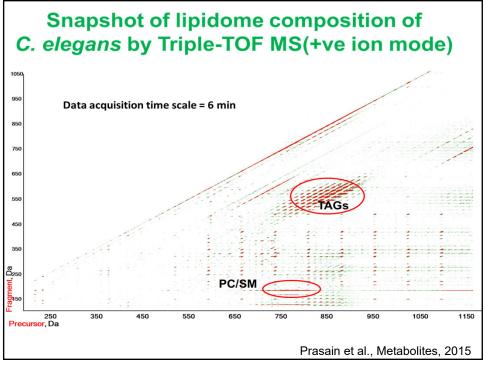
Sciex 5600 Triple-TOF

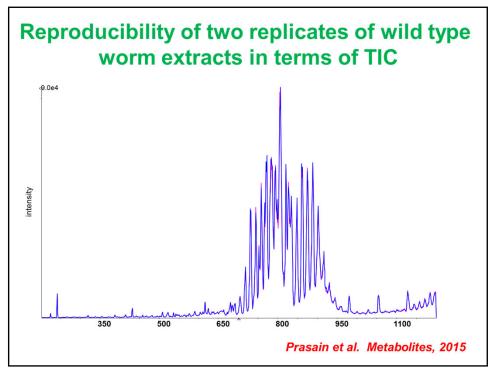
- Over 30,000 mass resolution
- <5 ppm mass accuracy
- Very fast acquisition of MSMS spectra (10 ms)
- Precursor and neutral loss analyses are possible performed post hoc

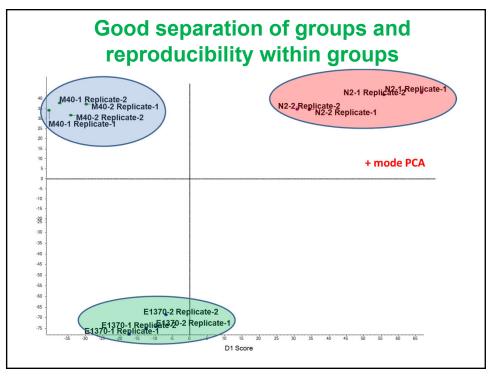


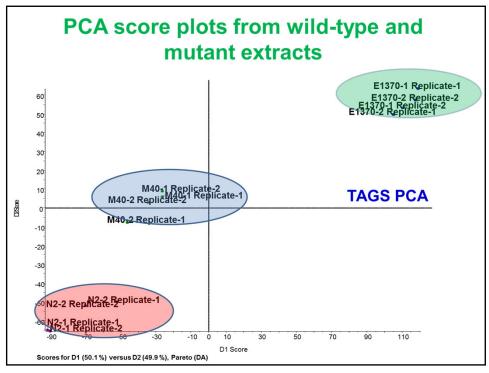
AB Sciex Triple TOF 5600

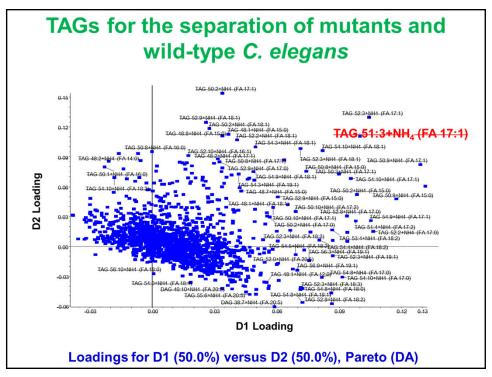


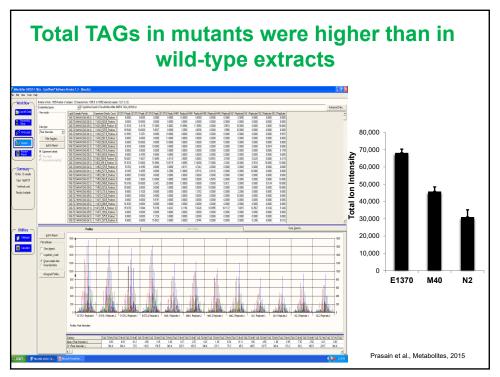


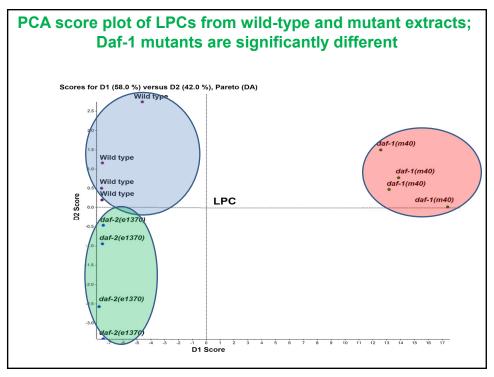












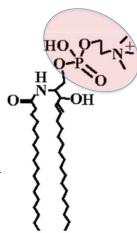
MS/MS^{ALL} limitations

- · Poor selectivity
- MS/MS fragment ions may contain fragment ions from other precursor ions such as isobaric species to precursor ion of interest
- Limited to infusion or flow injection acquisition

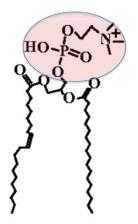
13

The problem of analyzing lipids

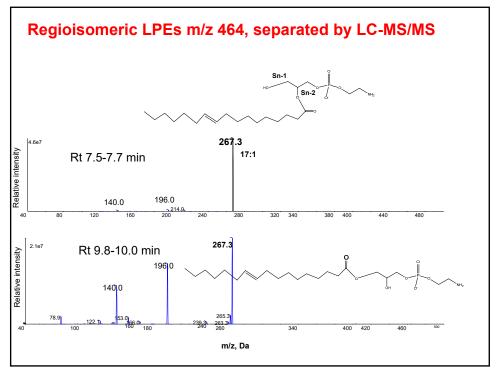
- Despite the sheer number of lipids, the units comprising them are closely related and therefore they have similar masses
- Sphingolipids may only be different in mass by 1 Da from their PC analog
 - 13C-Isotope profiles overlap
 - Head groups are the same

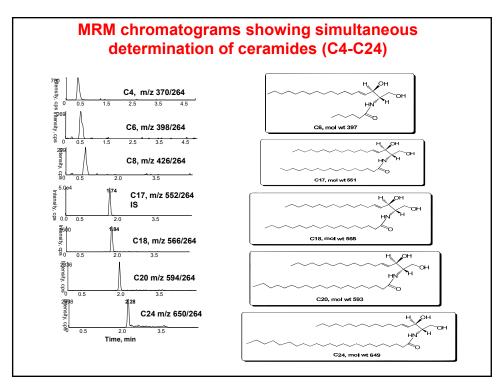






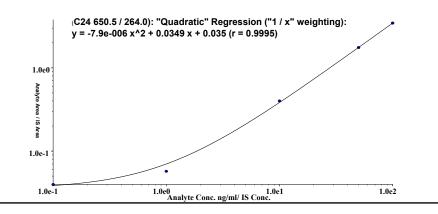
Phosphocholine





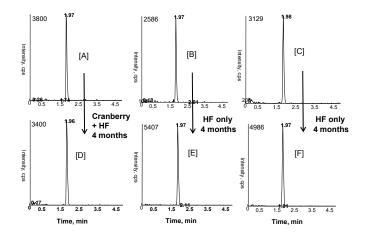
A linear response for Cer C24:0 was observed over a range of 0.1-100 ng/ml with correlation coefficient greater than 0.99

Sample Name	Analyte Peak Name	Calculated Concentration (ng/mL)	Accuracy (%)
Ceramide Standard 100 ng/ml	C24 650.5 / 264.0	100	100
Ceramide Standard 50 ng/ml	C24 650.5 / 264.0	49.8	99.6
Ceramide Standard 10 ng/ml	C24 650.5 / 264.0	10.5	105
Ceramide Standard 1 ng/ml	C24 650.5 / 264.0	0.634	63.4
Ceramide Standard 0.1 ng/ml	C24 650.5 / 264.0	0.132	132



17

Cranberry fruit powder treatment reduced the HF induced increased levels of Ceramide C20 in rats



[A]-[C] represent base line plasma ceramide C20 (594/264) from three animals [D] after 4 months treatment with cranberry (1 g/kg b. w. and high fat diet

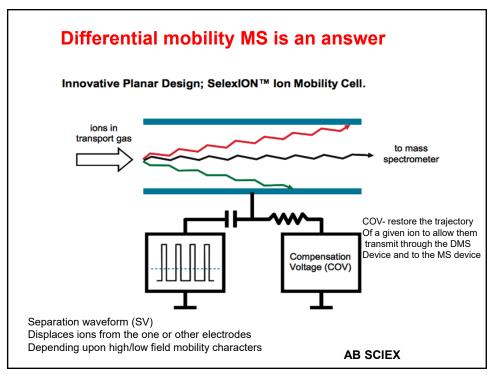
[E] & [F] after 4 months treatment with high fat diet only

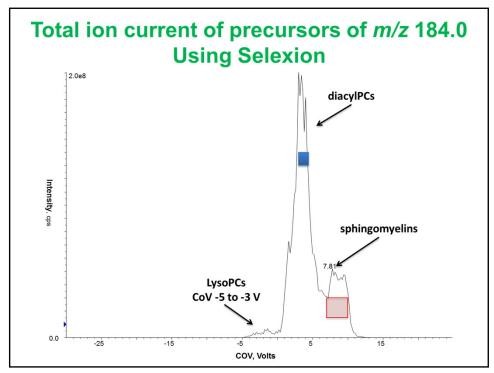
Prasain and Barnes, Food Frontiers 2020

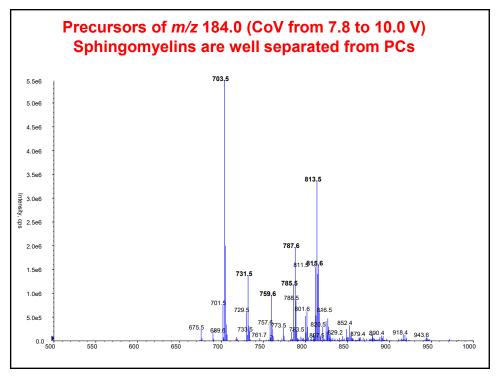
Differential mobility mass spectrometry

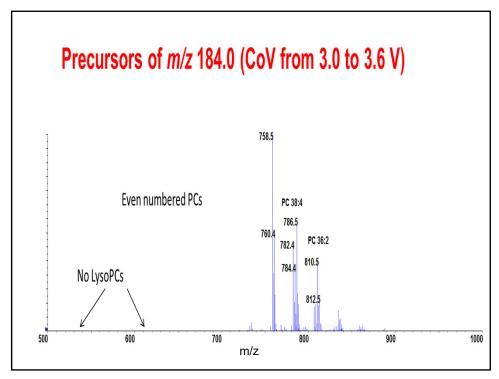
- When a fragment ion may have multiple precursor ions, the precursor ions may be separable by DMS before they enter the mass spectrometer
- By scanning with the compensating voltage (CoV), the precursor ions enter the mass spectrometer at different CoVs
- (Note: Further separation is possible using resolving agents, e.g., isopropanol)

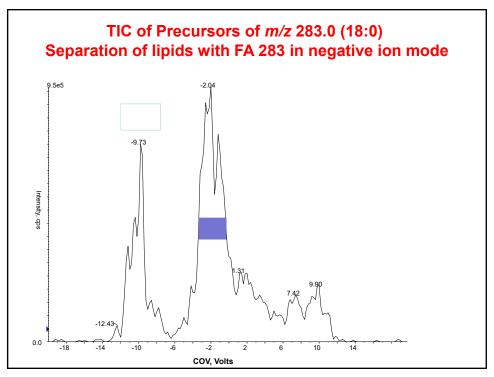
19

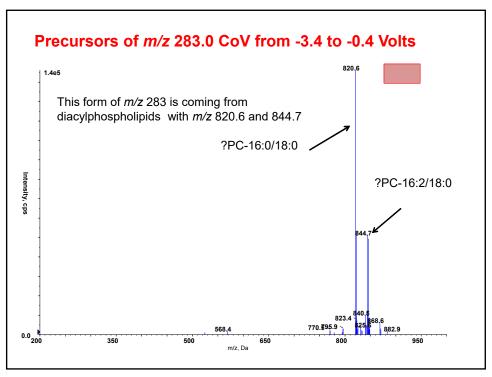


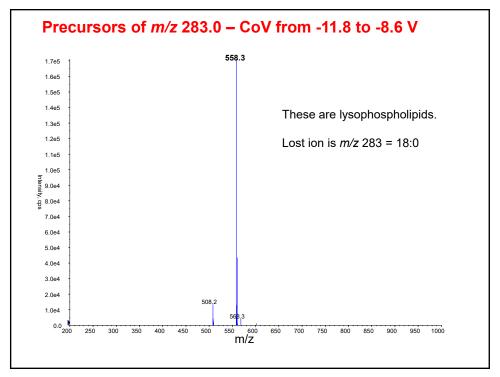












Conclusions

- Shotgun lipidomics approaches are high throughput and applicable to perform profiling as well as quantitative analysis of various lipids in biological samples.
- Differential ion mobility is useful for reducing or separating isobaric interferences
- LC-MS/MS method operating in multiple reaction ion monitoring mode (MRM) can be used for identification and simultaneous quantification of lipids.