Lipids Analysis

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Lipids

- Lipids are mostly very hydrophobic
- Most are conjugates of fatty acids of a variety of chain lengths, which have different degrees of unsaturation, cis-trans isomers, and chiral centers
- The conjugating frame to which the fatty acids bind can be quite hydrophilic
- This results in a very wide (evergrowing) number of lipid species

Analysis of fatty acids

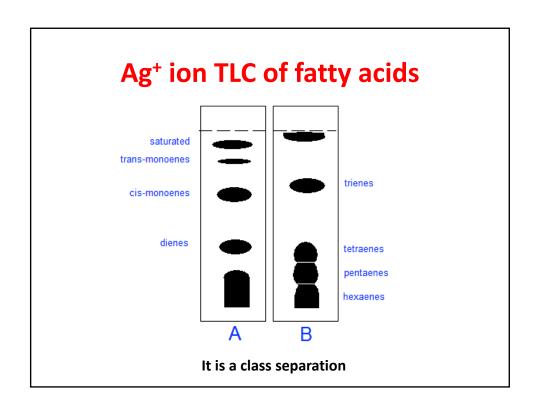
- Fractional crystallization
- Thin layer chromatography (TLC)
 - Argentation TLC (to separate according to number of double bonds)
- Gas liquid chromatography
 - Packed columns
 - Capillary columns
- LC-MS
- SWATH-MS
- · Differential ion mobility
- DESI-MS

Fractional crystallization

- Still used in industry
- Crystallization is used to determine whether adulteration of butter fat by other lower quality fats has occurred
- Unsaturated fats are more soluble at lower temperatures
 - Division into "stearins" and "oleins"
 - For fatty acids, make lead salts and cool in diethyl ether or ethanol – the saturated FAs crystallize out first

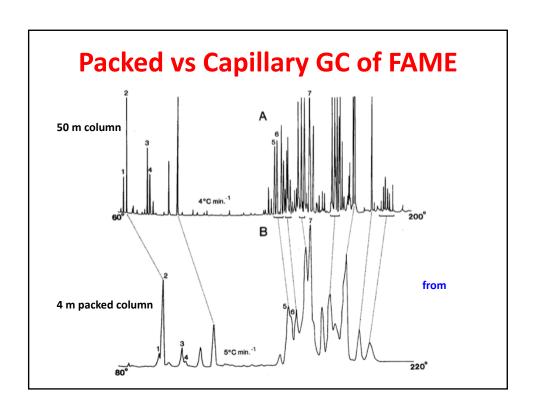
Thin-layer chromatography

- Fatty acids or methylated fatty acids separated on alumina or silica gel TLC
- When $AgNO_3$ is incorporated into the silica slurry before making the TLC plate, the observed separation is dependent on the degree of unsaturation (π -bonding)
 - Saturated
 - Mono-unsaturated
 - Di-unsaturated, etc.



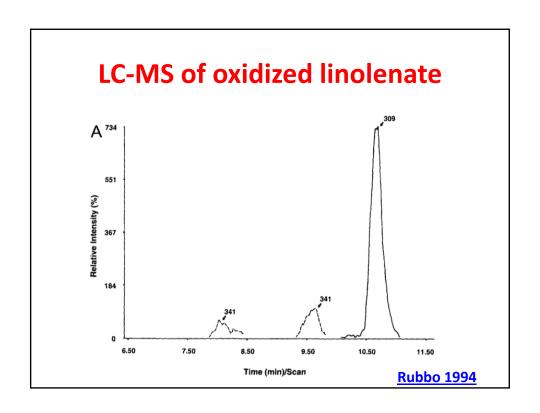
Gas-liquid chromatography

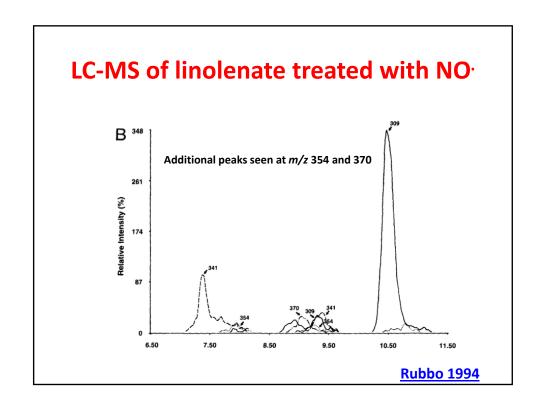
- 1952 Martin and James start GC by separating volatile fatty acids (C₁-C₆)
 - Quickly extended it to long chain FAs by methylating them
 - Used 5-6 feet x ¼ inch glass or stainless steel packed columns
- 1955 Patent for capillary, open tubular columns awarded
 - Did not enter commercial use until the mid-1970s

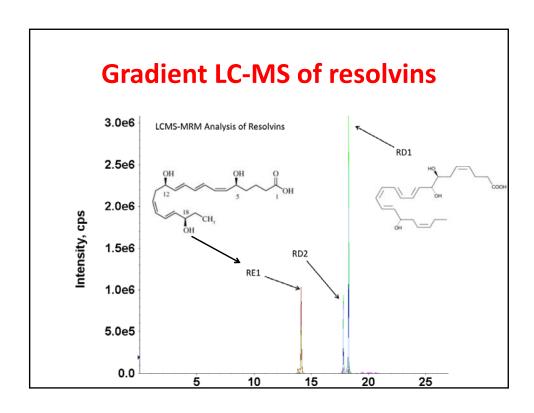


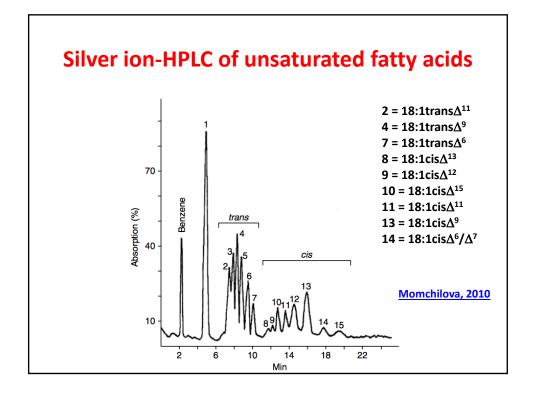
(HP)LC

- Reverse-phase LC
 - Can be used for lipid class separation based on hydrophobicity
 - Again, Ag⁺ can be introduced into the medium to enhance the separation of unsaturated fatty acids
 - Very difficult to detect lipids spectroscopically
 - LC-MS is the preferred method







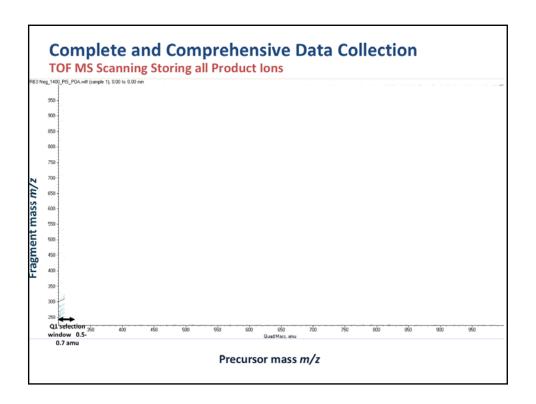


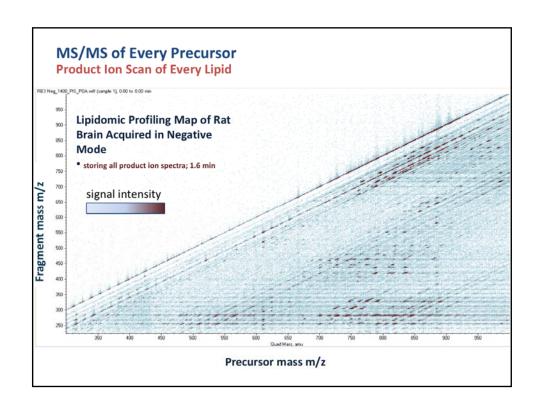
Modern lipidomics

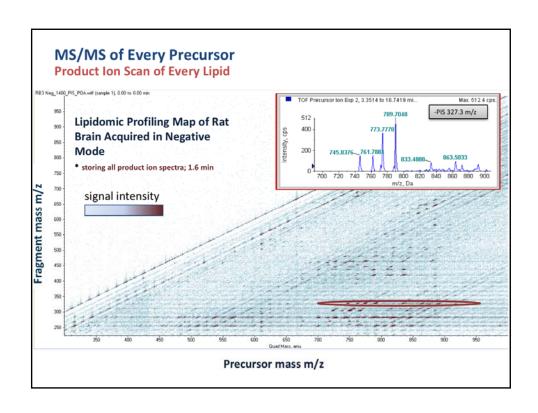
- Use of the SWATH-MS approach
- Preceded by total lipid extraction using a two-phase partition by adding CHCl₃:MeOH
 - Bligh-Dyer and Folch extractions
 - Crucial to do so in an atmosphere of argon and in the presence of butylated hydroxytoluene to prevent oxidation

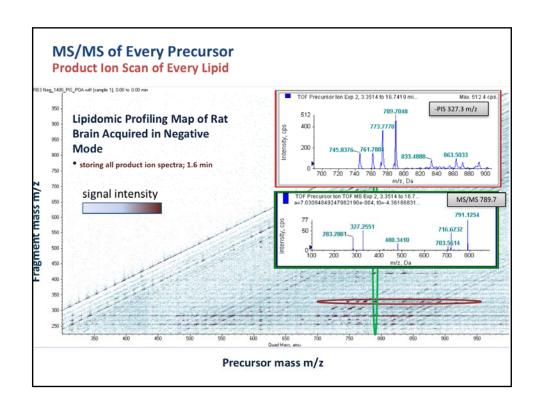
SWATH-MS

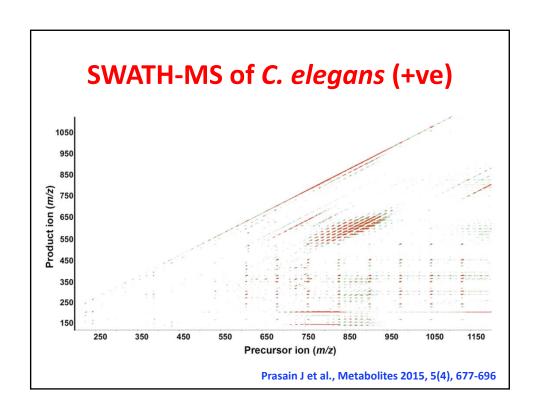
- Based on an infusion strategy on a 5600 TripleTOF
 - lons are filtered 1.2 m/z at a time in the quadrupole over a m/z range of 200-1200
 - The filtered ions are collisionally dissociated and fragment ions analyzed by the TOF analyzer
 - MSMS spectra collected for 500 msec for each m/z, i.e., infusion for 500 sec (8.33 min)





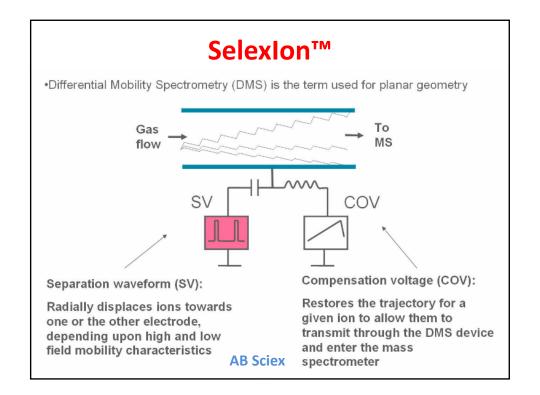


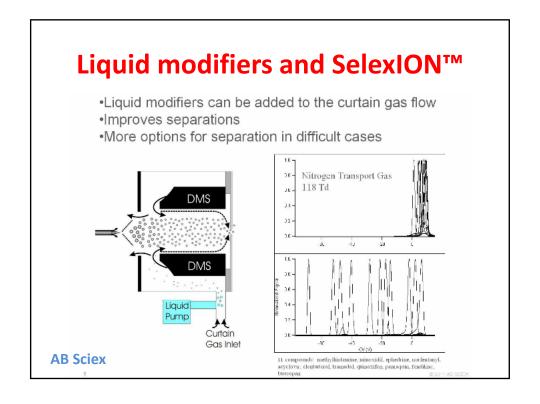


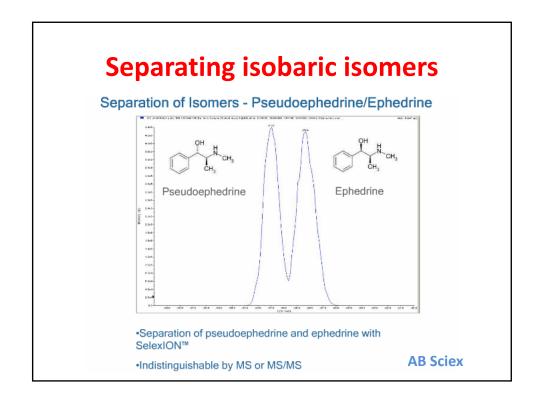


Other MS methods for lipids

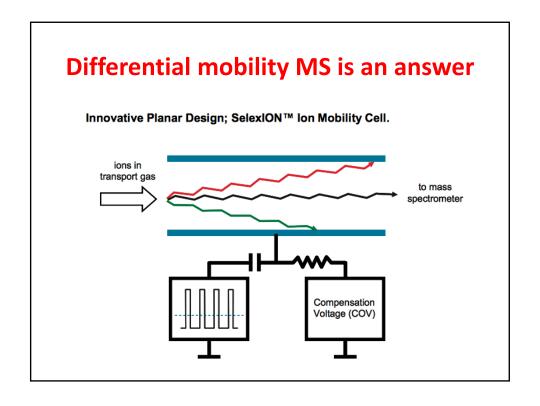
- SWATH-MS is comprehensive no stone unturned
- However, many lipids overlap in mass and there are also isomers with the same mass
- To observe more individual lipids, it is necessary to resolve lipids before analyzing them in the mass spectrometry
- Even then, isomers can be a problem
- · A form of ion mobility may be the answer to this





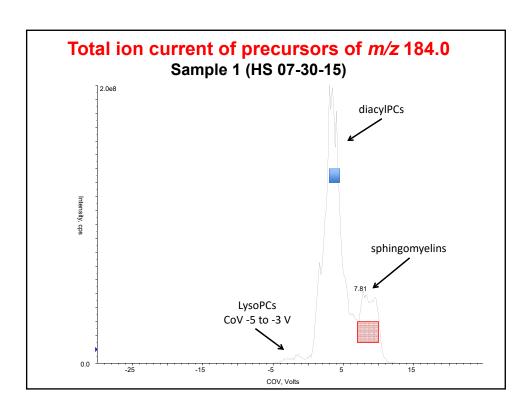


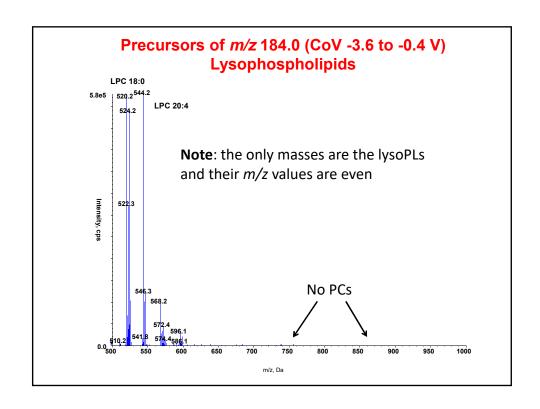
The problem of analyzing lipids Despite the sheer number of lipids, the units comprising them are closely related and therefore they have similar masses • Sphingolipids may only be different in mass by 1 Da from their PC analog 13C-Isotope profiles overlap - Head groups are the Sphingolipid Phospholipid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

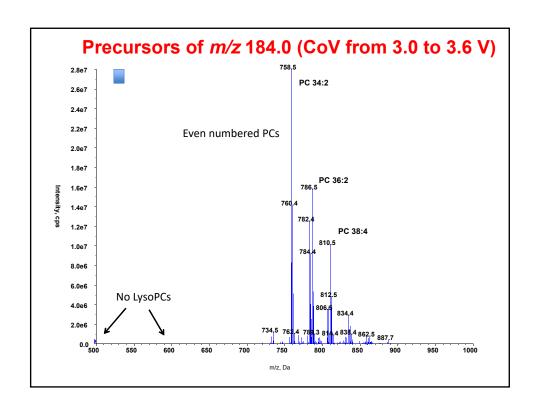


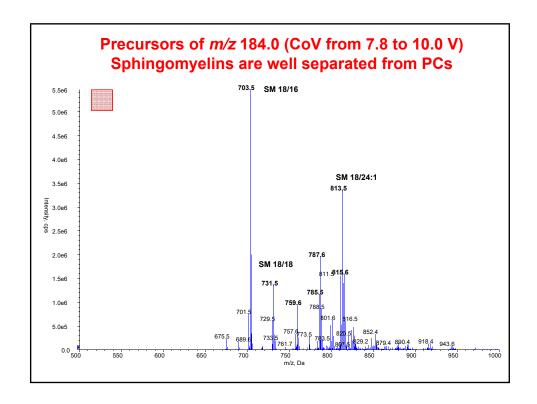
Differential mobility mass spectrometry

- 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, the precursor ions enter the mass spectrometer at different CoVs
- (Note: Further separation is possible using resolvating agents, e.g., isopropanol)



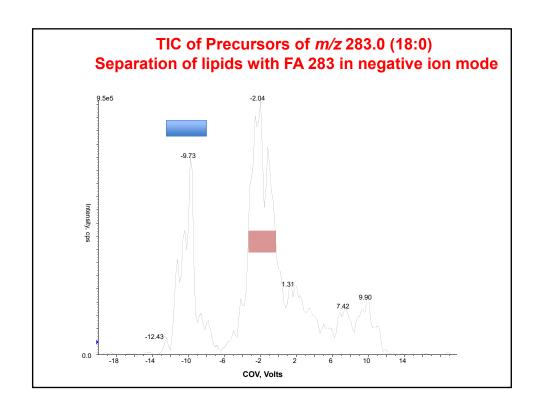


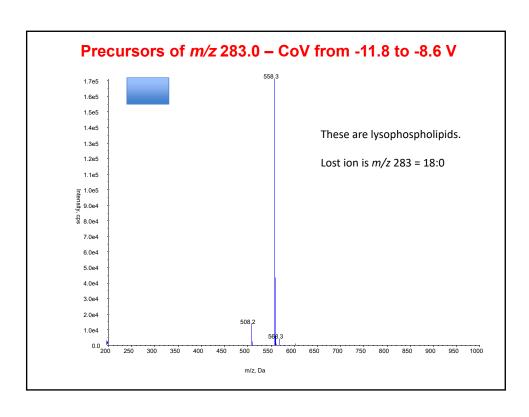


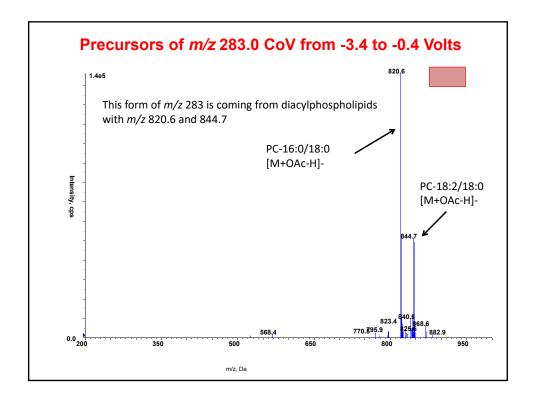


Origins of negatively charged product ions

Precursors of *m/z* 283 (stearate, 18:0) studied at different compensation voltages

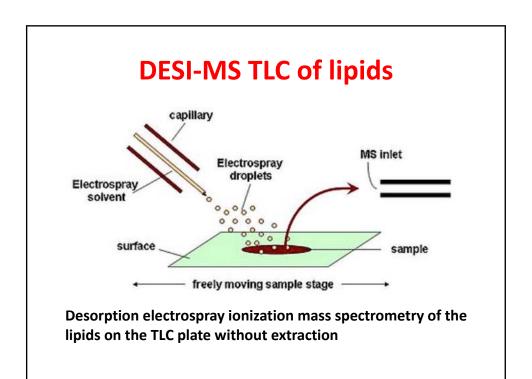


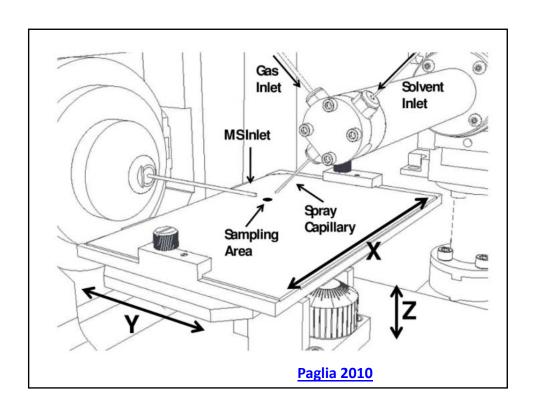


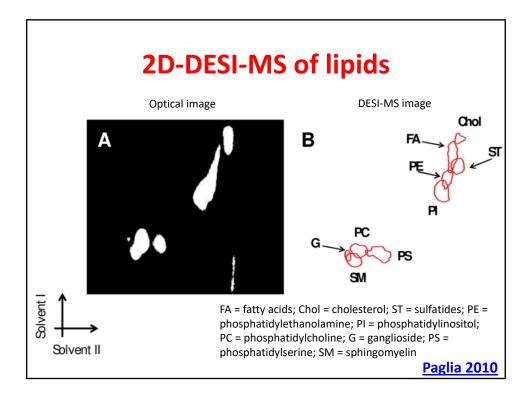


Summary

- Differential mobility mass spectrometry (DMS) is an important new tool in the study of lipids
 - It overcomes many of the problems that beset the analysis of lipids with overlapping masses
- Further separation (not exploited yet) comes from differential resolvation with specific solvents
- Metabolon has introduced a kit for the analysis of lipids where the extraction solvent contains ~1,000 deuterated lipid internal standards enabling absolute quantification of a wide range of lipids using DMS







References

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