



Knowledge that will change your world

1st UAB Metabolomics Workshop
July 22-25, 2013

Future of Metabolomics

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Issues in metabolomics research

- **Standards and methods standardization**
- **Improved databases and MS/MS**
- **Spatial distribution**
 - Imaging mass spectrometry
 - DART
 - Spotting methods
- **Localized metabolomics**
 - Head space GC-MS
 - Breath and other body odors
 - ?? urine “odor”
 - iKnife-MS
 - Metabolome of the “smoke” created with cauterized surgical knife
- **Derivatives and Isotope labeling**

Standards

- The majority of compounds that are detected have not been fully characterized and don't exist in pure standard form
- The LipidMaps endeavor (www.lipidmaps.org) has gone a long way to improve the state of that field
- A heavy investment in chemical synthesis of other metabolites is sorely needed

Standardized methods

- Importance of standardized operating procedures and recording keeping of any deviations from them – use of LIMS
- NMR yields absolute, quantitative data, but for a limited number of metabolites
- Whereas LC-MS is very sensitive, quantification is a challenge
- For primary untargeted metabolomics data collection, the Qtof instrument is both the fastest and most robust
 - The UK Phenomics Center is using a single manufacturer's Qtof - [harmonization](#)

Capillary electrophoresis-MS and metabolomics

- *“Capillary electrophoresis and MS is a marriage made in heaven, but not on earth”* (Richard Smith)
- Can CE-MS provide for metabolomics what CE has done for DNA deep sequencing?
- CE has many forms and can accommodate a wide variety of even hydrophilic compounds
- The interface is the key component

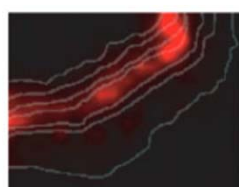
Can we improve the chromatography?

- The sheer numbers of metabolites demands a better chromatographic solution
 - CE-MS is one answer
- Taking a leaf from the gas chromatography community, LC needs to move towards open tubular columns
 - The back pressure of smaller and smaller particles is limiting
 - So, get rid of the particles

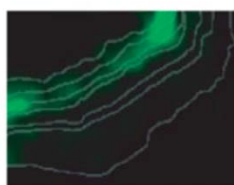
Understanding where metabolites are

- When we collect the metabolites from tissue or from cells, we destroy their localization information
- We assume that all metabolites are in contact with each other
- But there are as many as 15 distinct compartments in cells
- Some cells rely on neighboring cells for specific metabolites

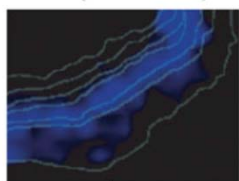
Differential lipid distribution in the retina



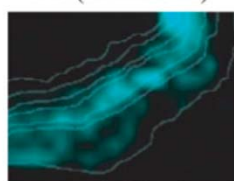
m/z 756.5
PC (16:0/16:0)



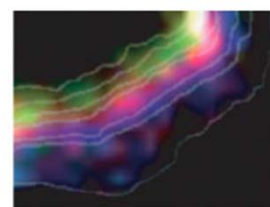
m/z 782.7
PC (16:0/18:1)



m/z 797.4
PC (18:0/22:6)-N(CH₃)₃



m/z 769.5
PC (16:0/22:6)-N(CH₃)₃



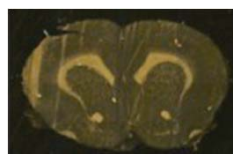
merged

MALDI-QIT-TOF
Hayasaka et al. 2008
8 μm sections sprayed
with matrix (in MeOH)
using an airbrush

Imaging of lipids in tissue

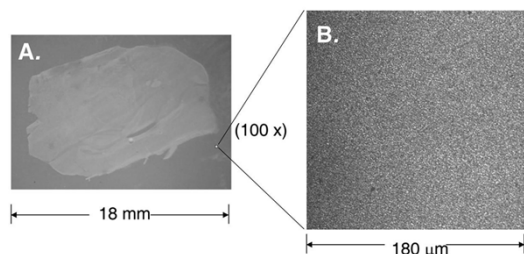


Fresh frozen tissue



Frozen tissue sectioning

Cut and transfer 10-20 micron section to conductive glass slide



DHB (MALDI matrix) is transferred by sublimation. This produces a very uniform coating (as shown in B) and high resolution images

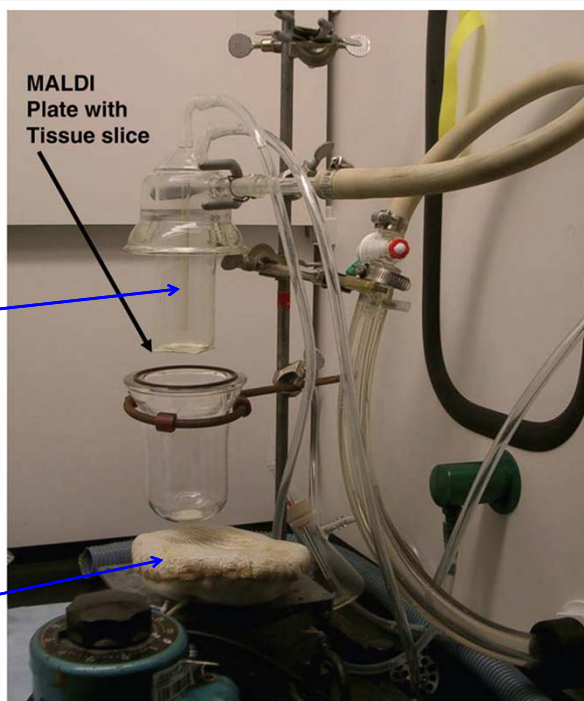
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Sublimation method for applying MALDI matrix to the tissue slice

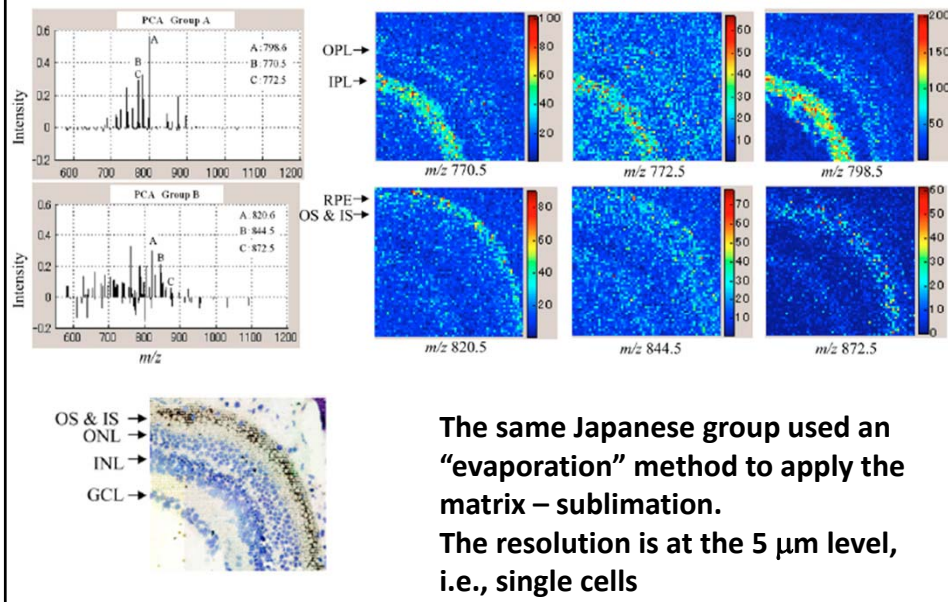
MALDI plate is upside down – ice is placed inside the glass “finger” to condense the DHB

A cold trap (not shown) is needed after the heated chamber to protect the vacuum pump

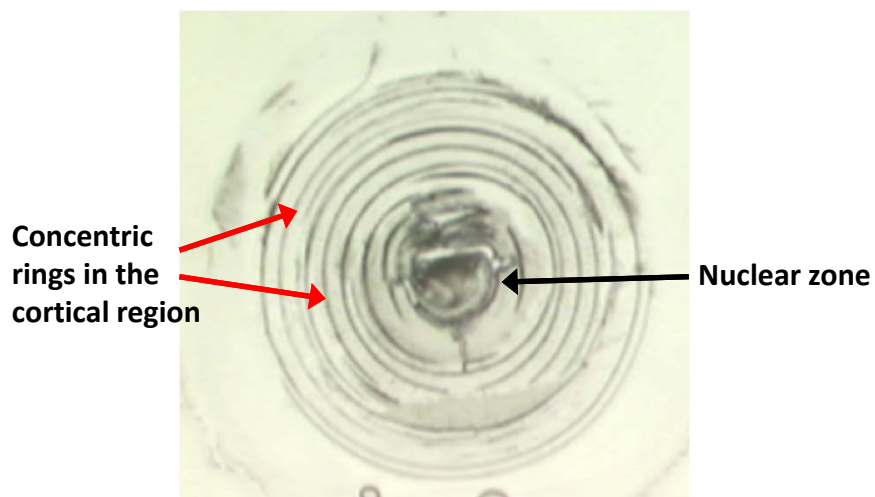
Sand bath to heat the DHB



Using a higher resolution approach

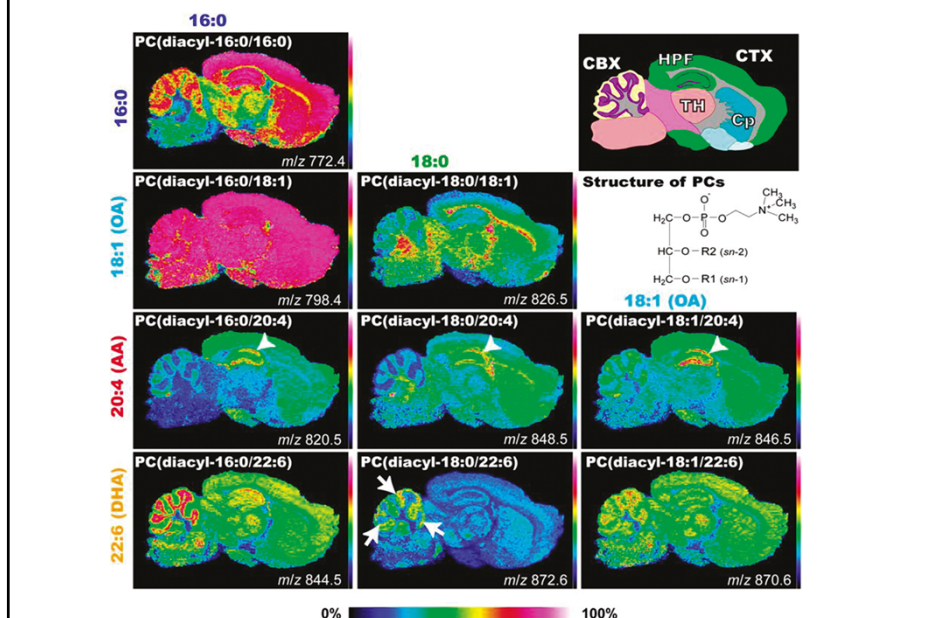


Frozen, thin section of zebrafish lens

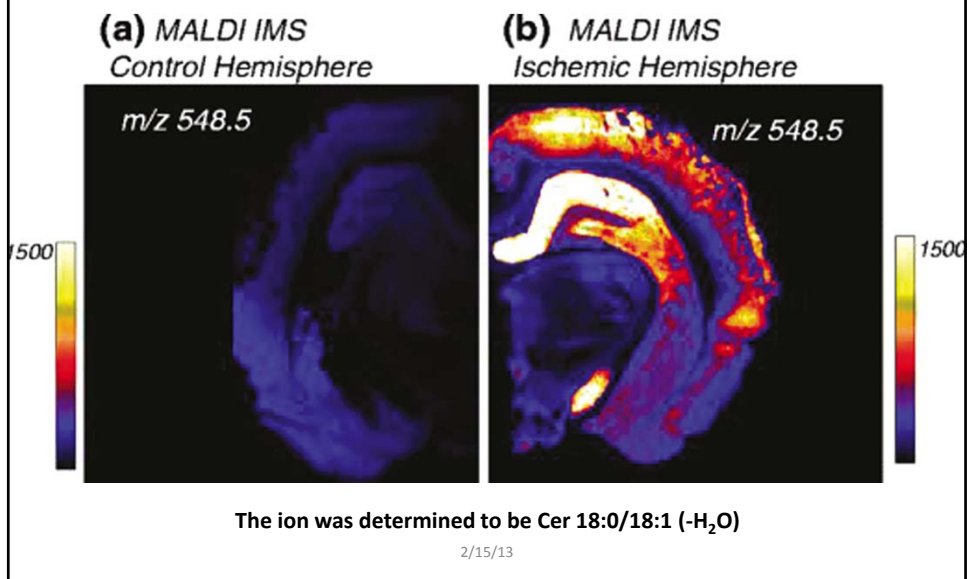


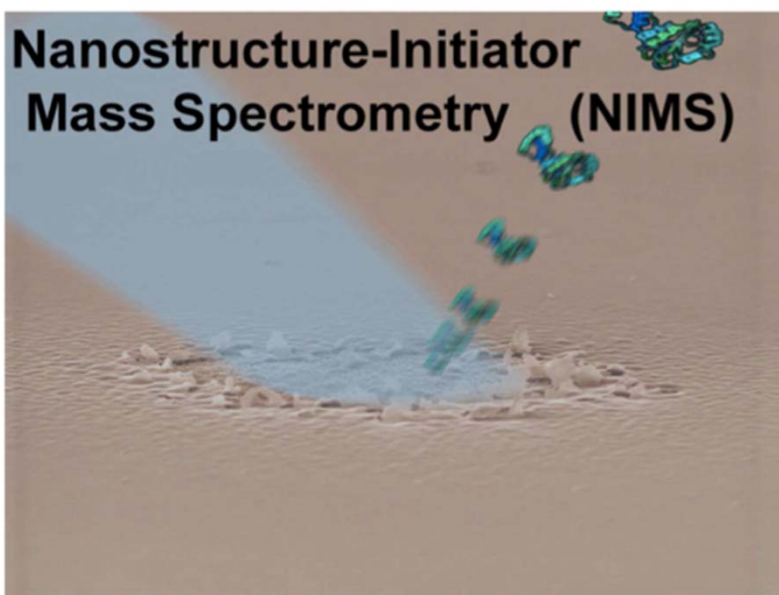
Miranda Collier, 2013

PC species distribution in the brain



Ceramides and ischemia by IMS

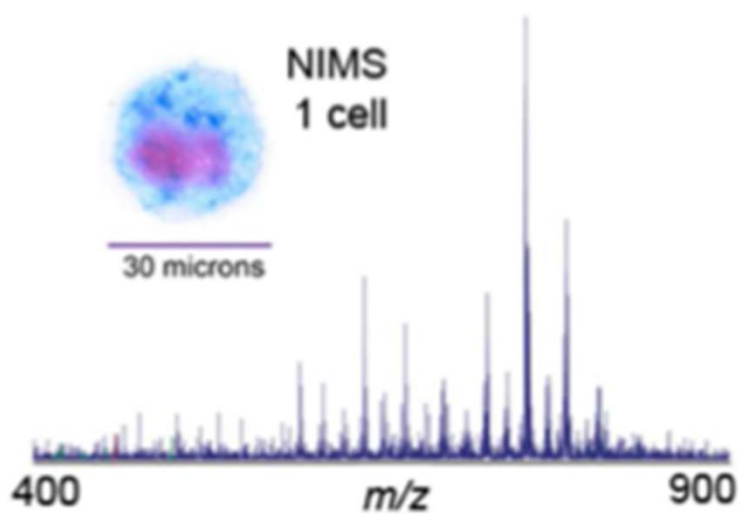




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Greving et al. 2011

Spectrum from a single cell

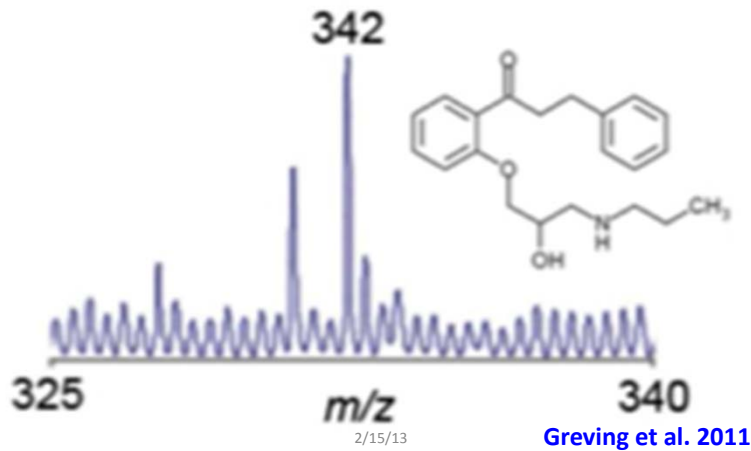


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Greving et al. 2011

Extreme sensitivity

propafenone (650 ymol)



Metabolism and time

- Not only should metabolites appear in the right place, there is also the question of the importance of the timescale
- Metabolism defects in the heart may be only seconds away from death – rogue wave in metabolism??
- Irreversible damage to the brain may occur in minutes
- Go/No-go decisions for a cell to divide or apoptose may occur in a similar timescale

Can existing techniques provide answers?

- **Mass spec is mostly a destructive method**
 - However, it can measure volatiles rapidly on a sec timescale (as we'll see shortly)
- **NMR can provide spatial and descriptive information on living or unextracted materials**
 - However, sensitivity and speed of data acquisition are extremely limiting
- **Future in other spectroscopic techniques?**

Components of breath

- **Two types of components**
 - **Droplets in exhaled breath**
 - Can be condensed (proteins, lipids, metabolites)
 - **LC-MS analysis**
 - **True volatiles (requires GC-MS)**
 - **Non-condensable gases**
 - H₂, N₂, O₂, CO₂
 - **Organics**
 - Alcohols, aldehydes, alkanes

Detecting smells

- Unlike dogs, humans have relatively few olfactory receptors (dogs have lousy taste!)
 - We can detect certain “bad smells” (and cover these by washing and deodorants and fragrances)
- The reality is that we all emit gases that can be diagnostic
 - GC-MS using a headspace technique can be used
 - Breath hydrogen is diagnostic for lactase insufficiency

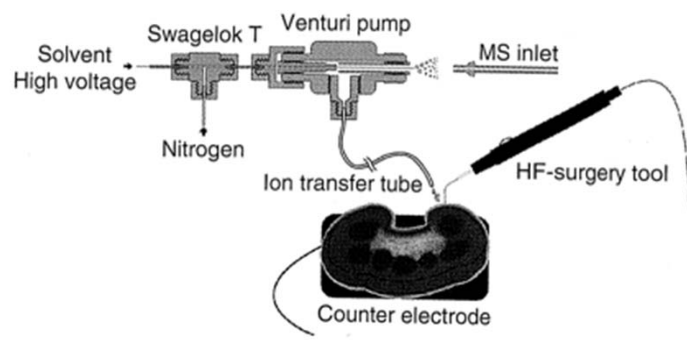
Portable Metabolomics Devices



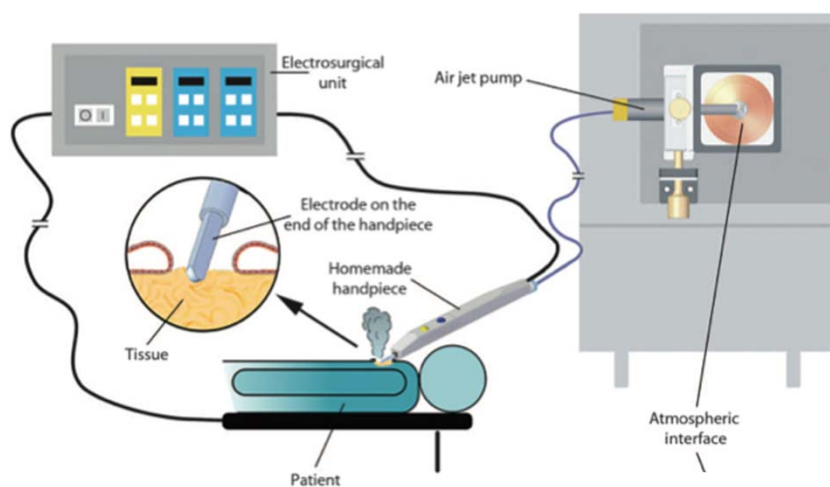
David Wishart

iKnife mass spectrometry

Electrospray post ionization mass spectrometry of electrosurgical aerosols. [Guenther et al., JASMS 2011](#)

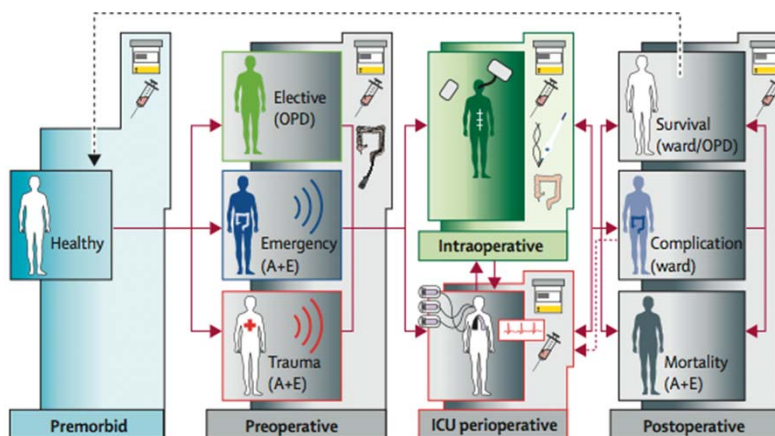


The iKnife – real time metabolomics



[Balog et al. Nature Med 5; 194r93](#)

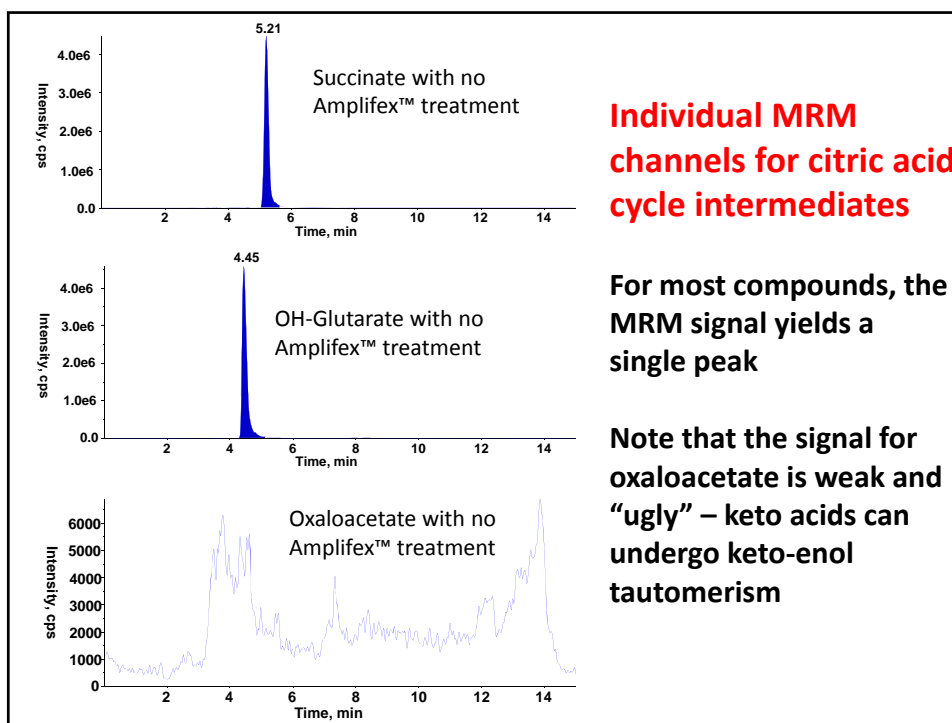
Clinical applications of metabolomics



Concepts of Jeremy Nicholson

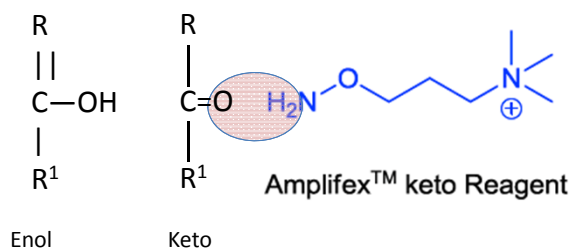
Back to derivatization in metabolomics

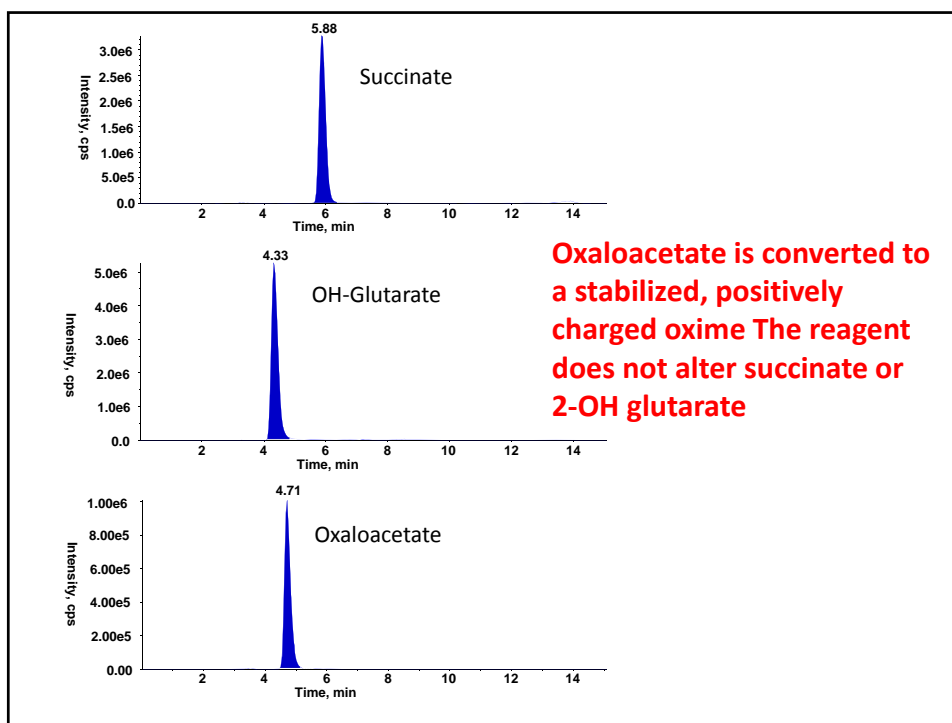
- A great advantage of LC-MS is its ability to measure compounds “as is”
- However, there are many compounds that either weakly form ions or are found in multiple chemical forms (e.g., steroids, ketosteroids, keto acids, fatty acids)
- The reagents used to overcome this problem can be isotopically labeled (iMetab)



Modifiers to the keto group

- Methoxylamine – CH_3ONH_2 (from GC-MS)
- Biotin hydrazide – RN.NH_2
- Amplifex™-Keto Reagent (ketosteroids)





Isotopic derivatization

- investigators are exploring reagents that can be coupled to -NH_2 and -COOH groups
 - c.f., ICAT, iTRAQ and TMT reagents for proteomics
- These may allow one sample of a pair to be labeled with a “light” reagent and another sample with the same reagent, but one incorporating ^2H , ^{13}C or ^{15}N atoms and therefore being “heavy”.
 - The two samples can then be mixed and run at the same time
 - Multiplexed reagents are also possible
- Software to sort this out has not yet been written