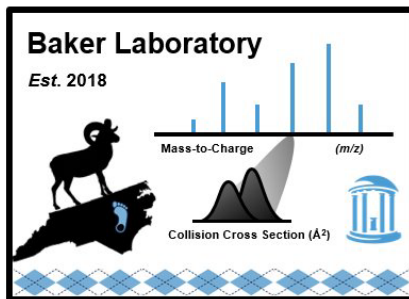
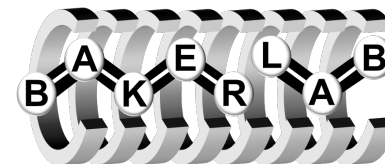


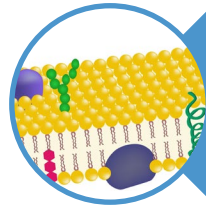


# Development and Application of a Multidimensional Lipid Database Containing Liquid Chromatography, Ion Mobility Spectrometry, and Mass Spectrometry Separation Characteristics in Skyline



**Amie Solosky**  
Skyline User Meeting  
ASMS 2024

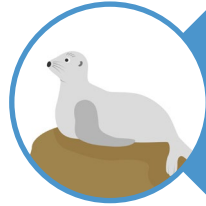




Lipid Overview



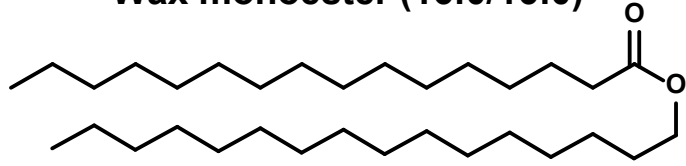
Database  
Creation



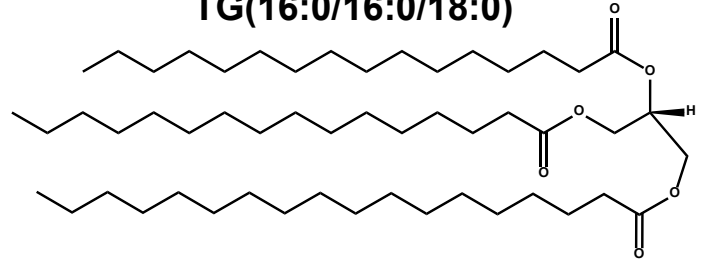
Sea Lion  
Application

# What is a lipid?

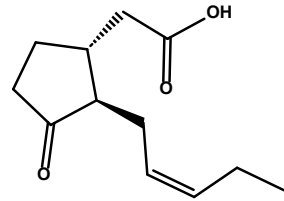
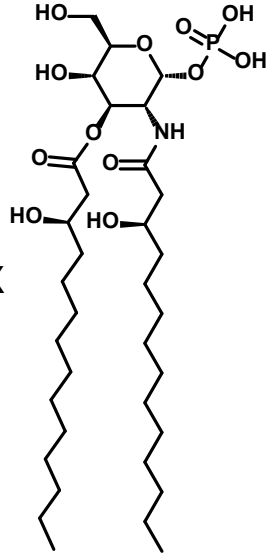
Wax monoester (16:0/16:0)



TG(16:0/16:0/18:0)



Lipid X



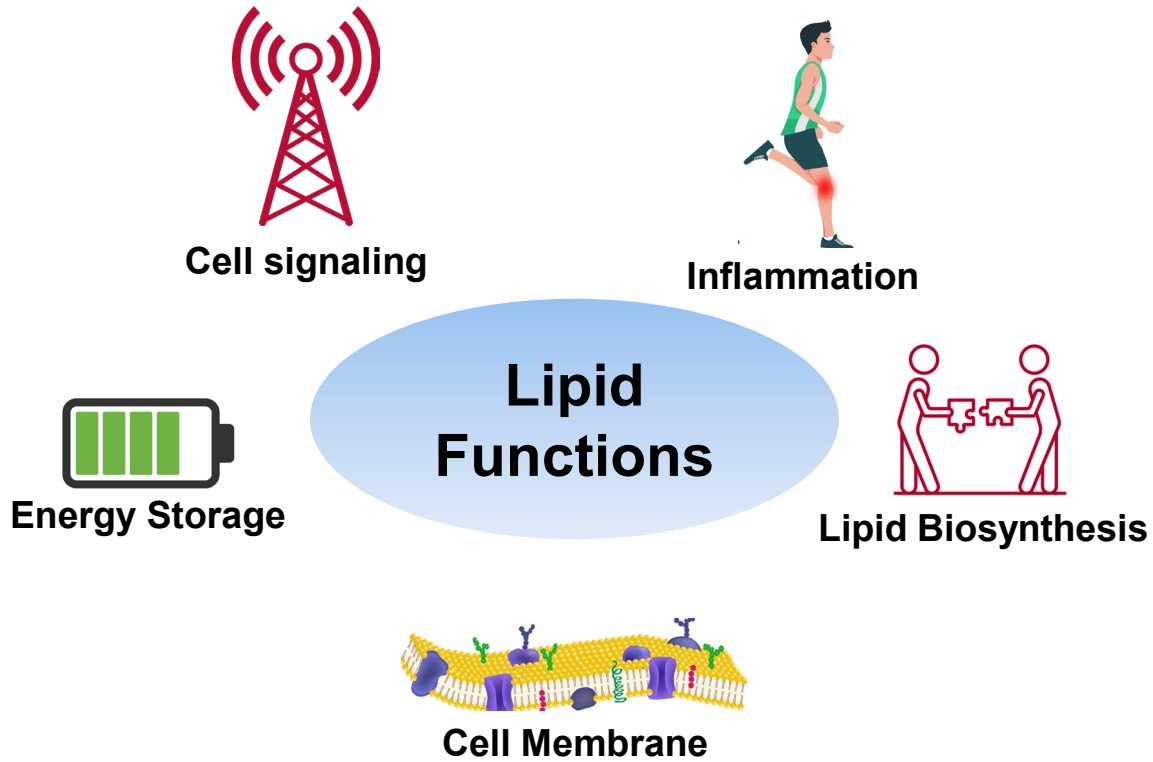
Jasmonic acid

Common denominator?



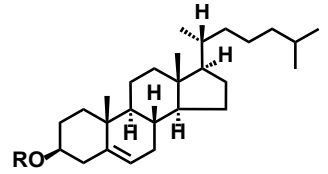
Lipids do not dissolve in water!

# Lipidomics - why are lipids so important?

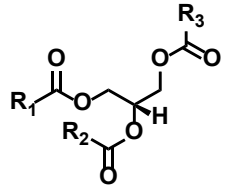


# Lipid structure

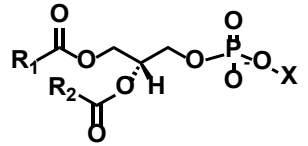
## Sterol Lipids



## Glycerolipids



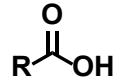
## Glycerophospholipids



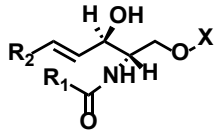
## Lipid Speciation



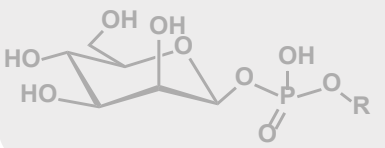
## Fatty Acids



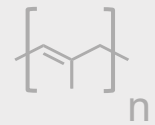
## Sphingolipids



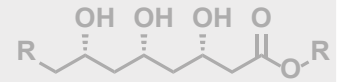
## Saccharolipids



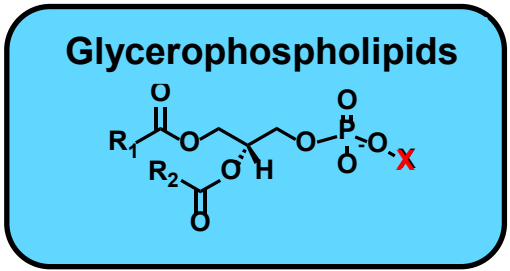
## Prenol Lipids



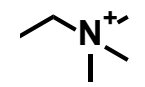
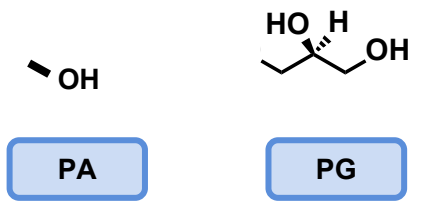
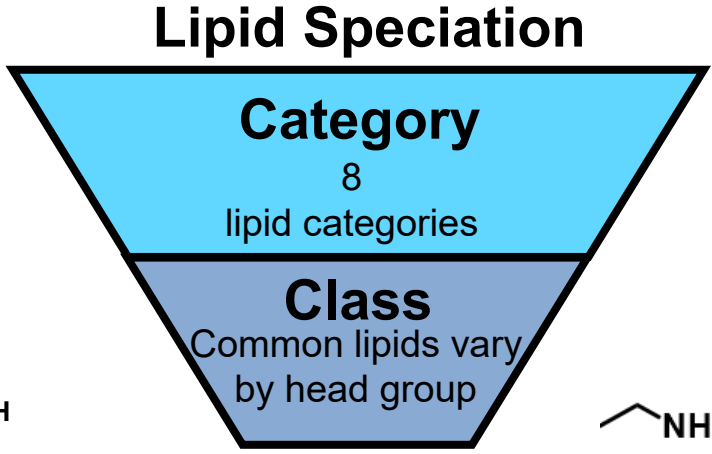
## Polyketides



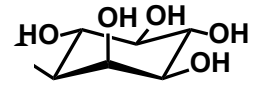
# Lipid classes



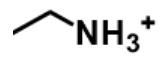
X =



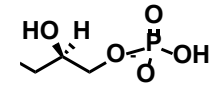
PC



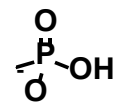
PI



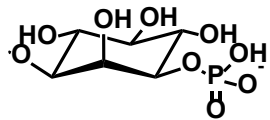
PE



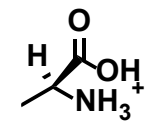
PGP



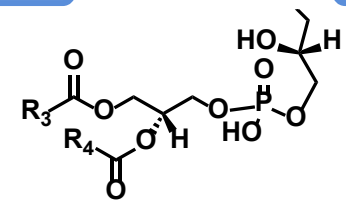
PPA



PIP

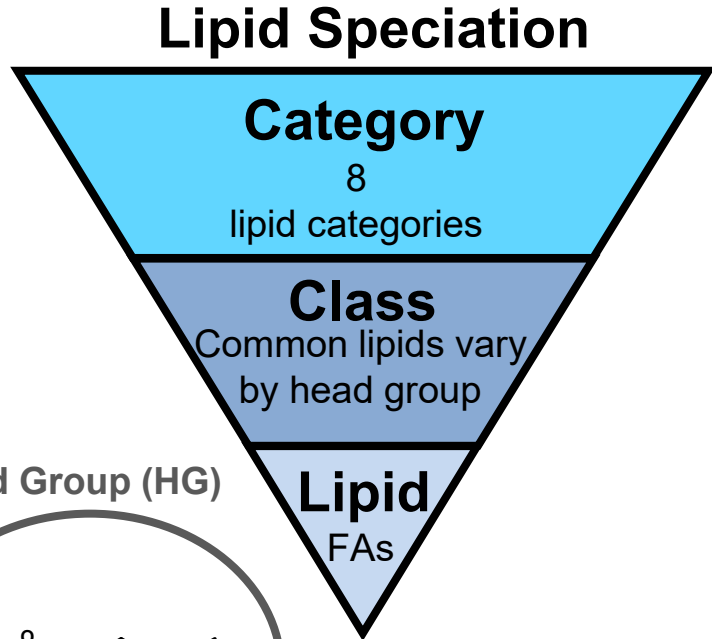
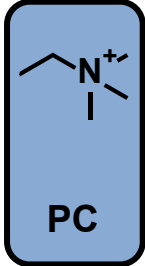
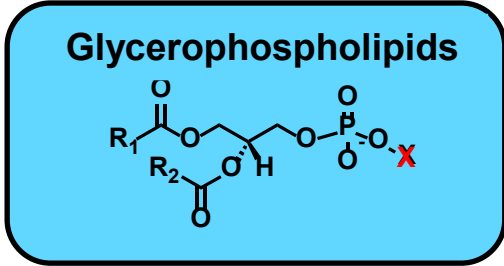


PS



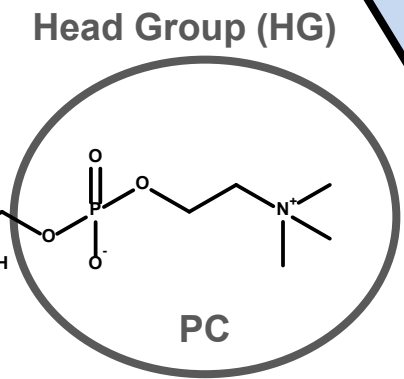
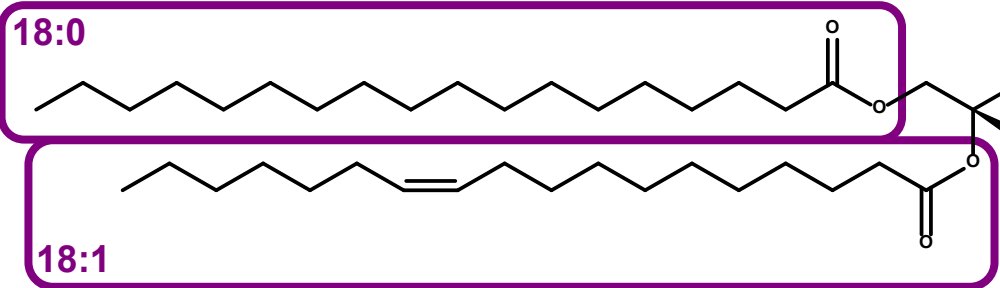
CL

# Lipid nomenclature



PC(18:0\_18:1) → PC(18:0/18:1)

**Fatty Acyl (FA)**



# Targeted vs. untargeted analysis

## Targeted Analysis

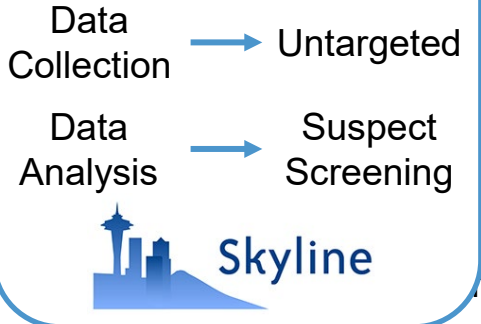
- Pre-determined list of analytes
- Only detect known features
- Quantitative
- Rapid & sensitive
- Regulatory methods



## Non-targeted Analysis

- Feature finding
- Requires high-resolution mass spectrometry

### Baker Lab Analysis



Complex multidimensional data

Time-consuming

Relative

Laboratory

### **Unknown unknowns**

Discovery



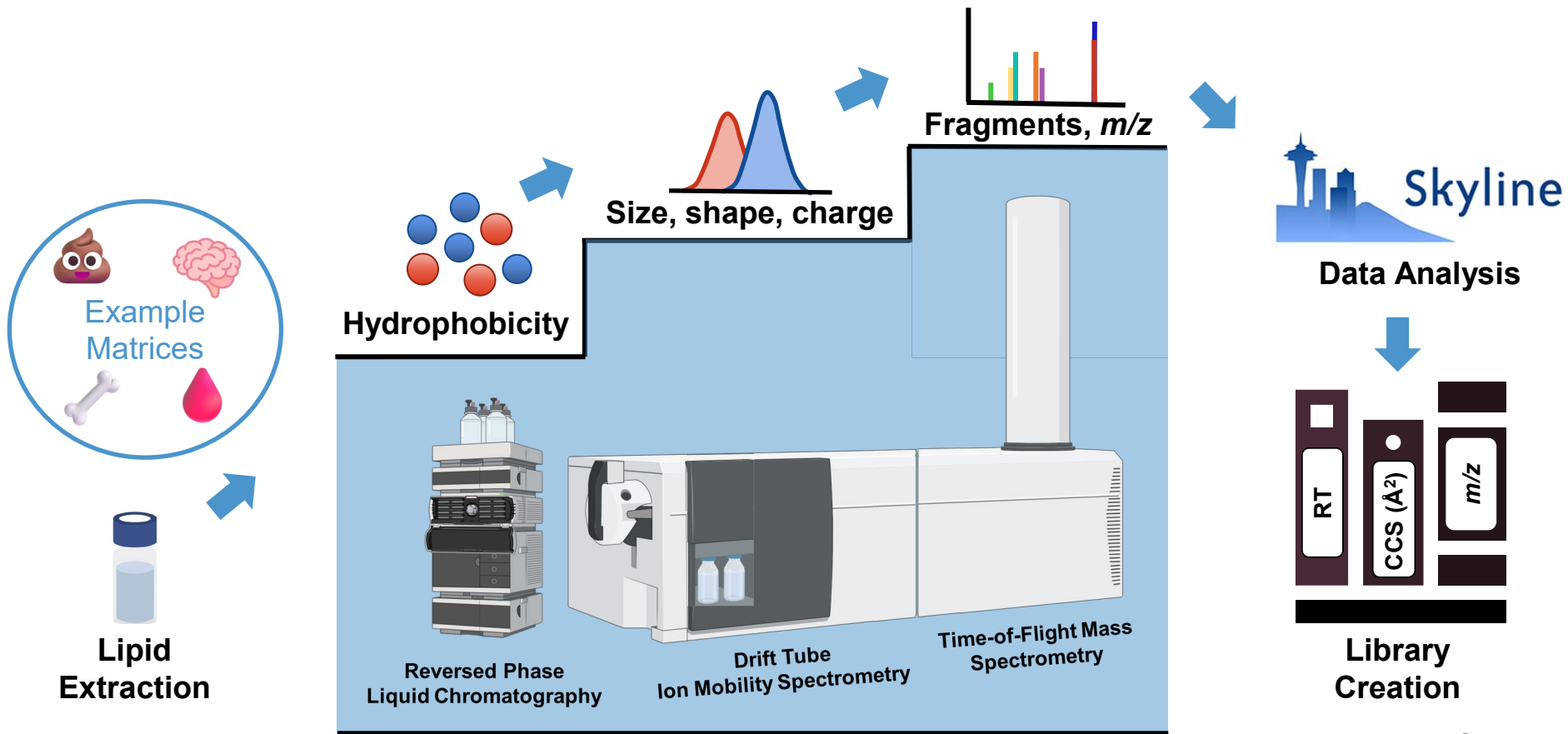
### **Known unknowns**

Suspect Screening



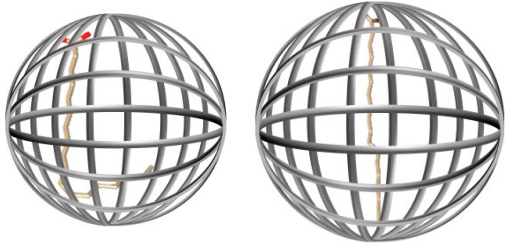
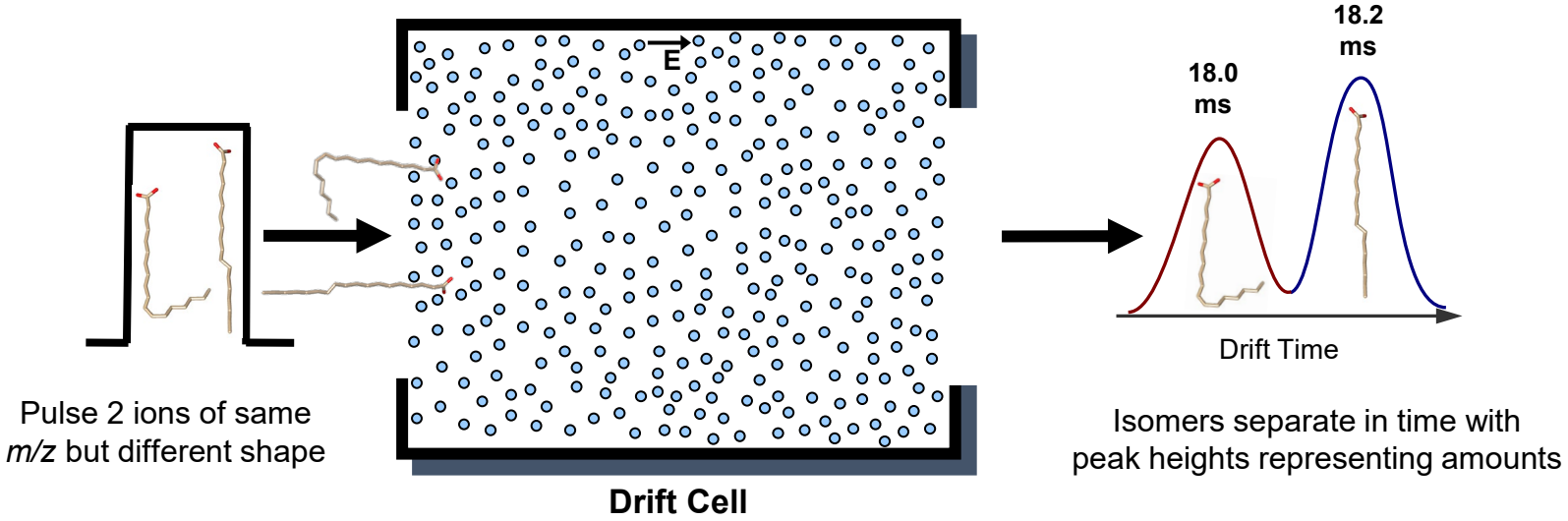


# Multidimensional separation technique for lipids



Multidimensional separation

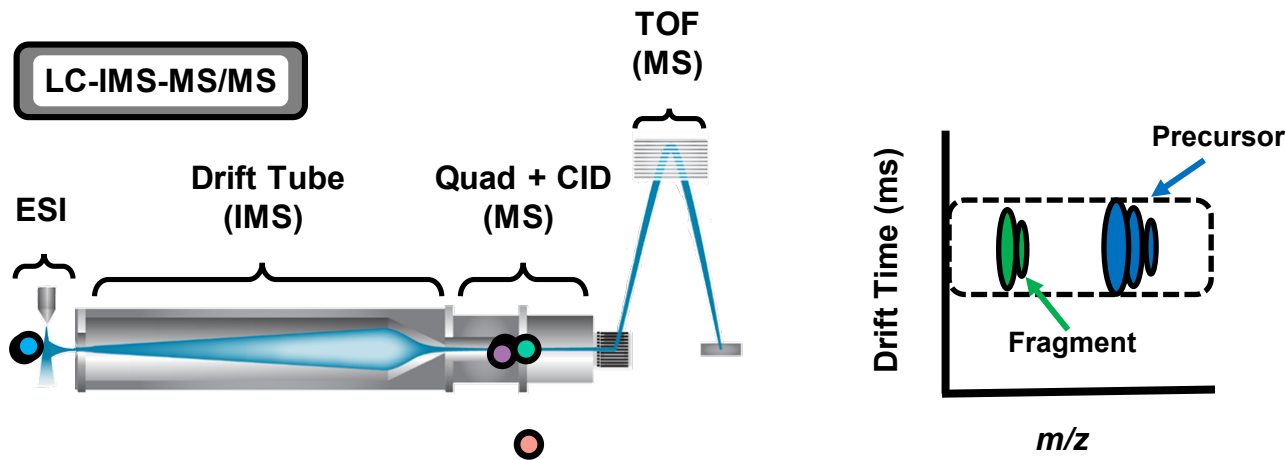
# Drift Tube Ion Mobility Spectrometry (DTIMS)



Higher drift time  
= Larger Collision Cross Section (CCS, Å<sup>2</sup>)

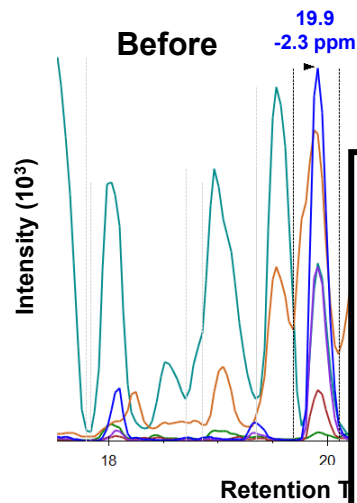
# Fragmentation

- DIA All Ions** – alternating precursor and fragment scans without precursor isolation



Drift time of fragments = drift time of precursor

# Drift time filtering to increase confidence



PE(18:0\_20:4)

[M-H]  
FM-4-LIP

nature protocols

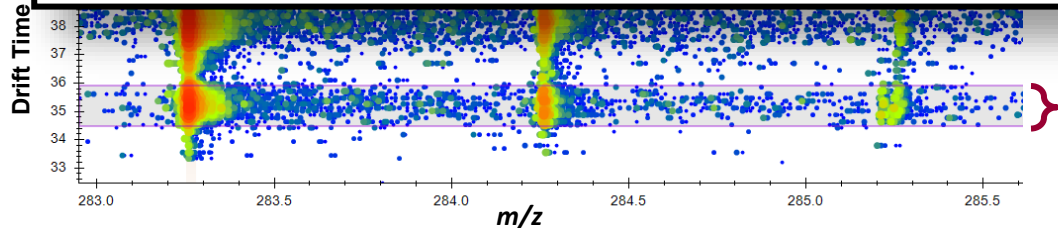
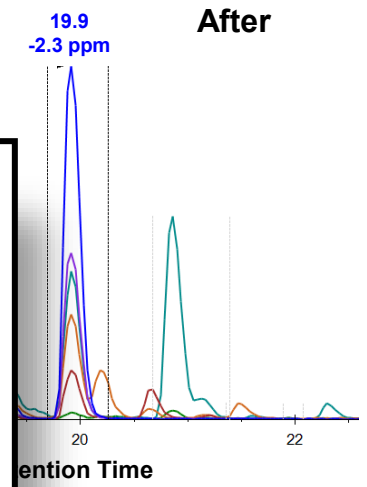
PROTOCOL EXTENSION

<https://doi.org/10.1038/s41596-022-00714-6>

Check for updates

## Utilizing Skyline to analyze lipidomics data containing liquid chromatography, ion mobility spectrometry and mass spectrometry dimensions

Kaylie I. Kirkwood<sup>1</sup>, Brian S. Pratt<sup>2</sup>, Nicholas Shulman<sup>2</sup>, Kaipo Tamura<sup>2</sup>, Michael J. MacCoss<sup>2</sup>, Brendan X. MacLean<sup>2</sup> and Erin S. Baker<sup>1,3</sup>✉

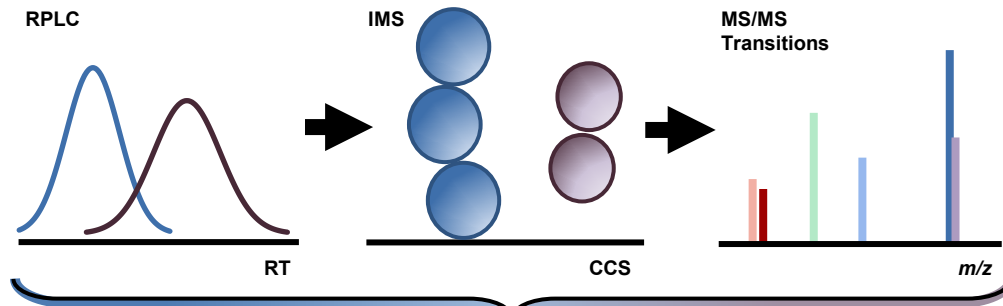
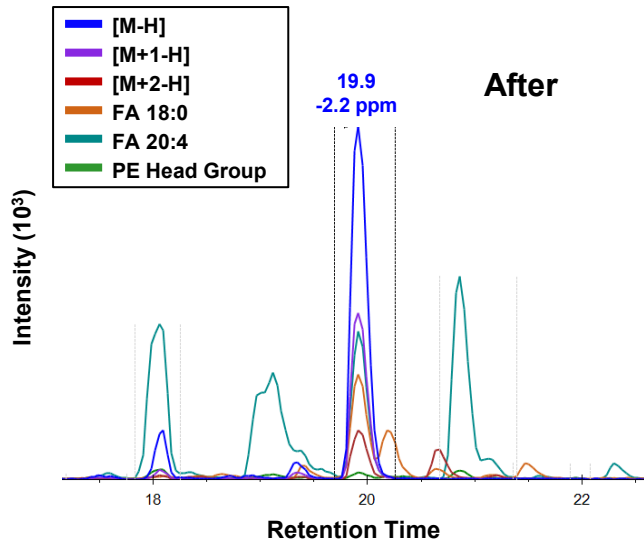


Signal included in chromatogram

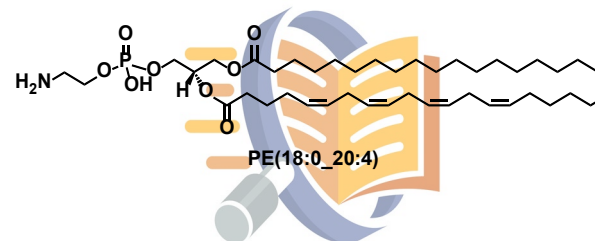
# Library match



PE(18:0\_20:4)



Library Match



# Baker Lab lipid library composition

## Lipid Library Magicians ✨



Dr. Kirkwood-Donelson

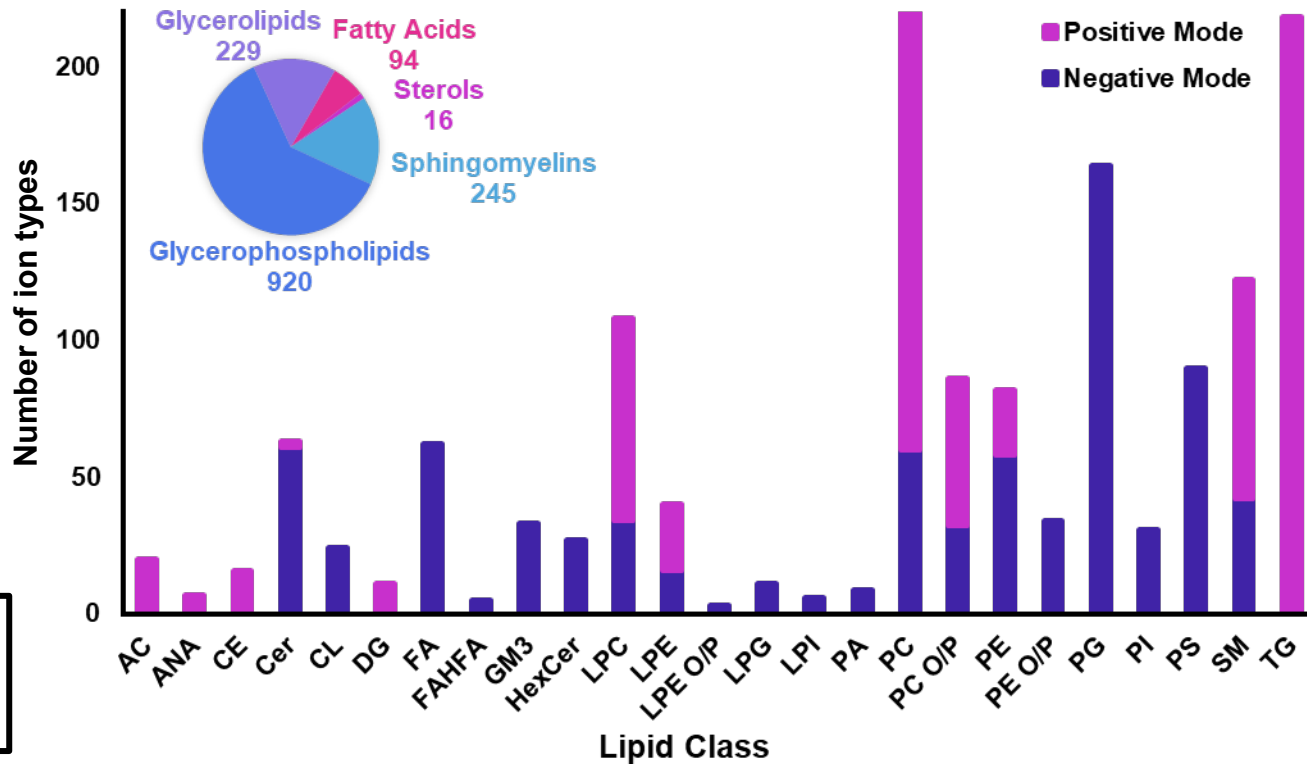


Dr. Odenkirk

**1504 precursor ion types:**

- 877 *unique* lipids

- 23 stable isotope labeled



# You can do it too!

LipidCreator (in silico)



Transition lists can come from any source: feature finding, online databases, literature, Panorama



LipidCreator 1.2.1 - Opened with Skyline

File Options Help

Home Glycerolipids **Glycerophospholipids** Sphingolipids Sterol lipids Lipid Mediators

Step 1: Precursor selection

Type:  Regular  Lyso  Cardiolipin  Plasmalogen

Head group: BMP, CDPDAG, DMPE, MMPE, PA, **PC**, PE, PEt, PG, PI, PIP, PIP2, PIP3, PS

Fatty acyl chain: 18, No. DB: 0, No. Hydroxy: 0

Fatty acyl chain: 18, No. DB: 1, No. Hydroxy: 0

FA  FAp  FAc

Positive adducts:  +H<sup>+</sup>,  +2H<sup>+</sup>,  +NH<sub>4</sub><sup>+</sup>

Negative adducts:  -H<sup>+</sup>,  -2H<sup>+</sup>,  +HCOO<sup>-</sup>,  +CH<sub>3</sub>COO<sup>-</sup>

Step 2: MS/MS selection: Manage heavy isotopes, MS2 fragments, Filters

Step 3: Assembly registration: Modify lipid, Add phospholipids

Lipid list

Category	Building Block 1	Building Block 2	Building Block 3	Building Block 4	Adducts	Filters	Options	Edit	Delete
Glycerophospholipid	HG: PC	FA:18; DB: 0; OH: 0	FA:18; DB: 1; OH: 0		H <sup>+</sup>	with precursors, with heavy			

Review Lipids

[Home](#) / Databases

## Databases

---

### Links to Baker Databases

#### Multidimensional RPLC-IMS-CID-MS Lipid Database

[Baker-Lab-Lipid-Database\\_03\\_28\\_2024](#)

[Download](#)

3/28/2024 Release Notes:

- This database contains information for 877 unique lipids from 1504 precursor  $[M+H]^+$ ,  $[M+NH_4]^+$ ,  $[M+Na]^+$ ,  $[M-H]^-$ ,  $[M-2H]^{2-}$ ,  $[M+HCOO]^-$  and  $[M+CH_3COO]^-$  ion types and their associated CID fragments (DOI: coming soon).
- This is an updated version of the Skyline [library](#) developed by Kirkwood et al., 2022 (DOI: [10.1021/acs.jproteome.1c00820](https://doi.org/10.1021/acs.jproteome.1c00820)).
- This database has 361 additional lipids, including heavy labeled internal standards by Avanti Polar Lipids
- A new matrix was added: California Sea Lion plasma



<https://tarheels.live/bakerlab/databases/>



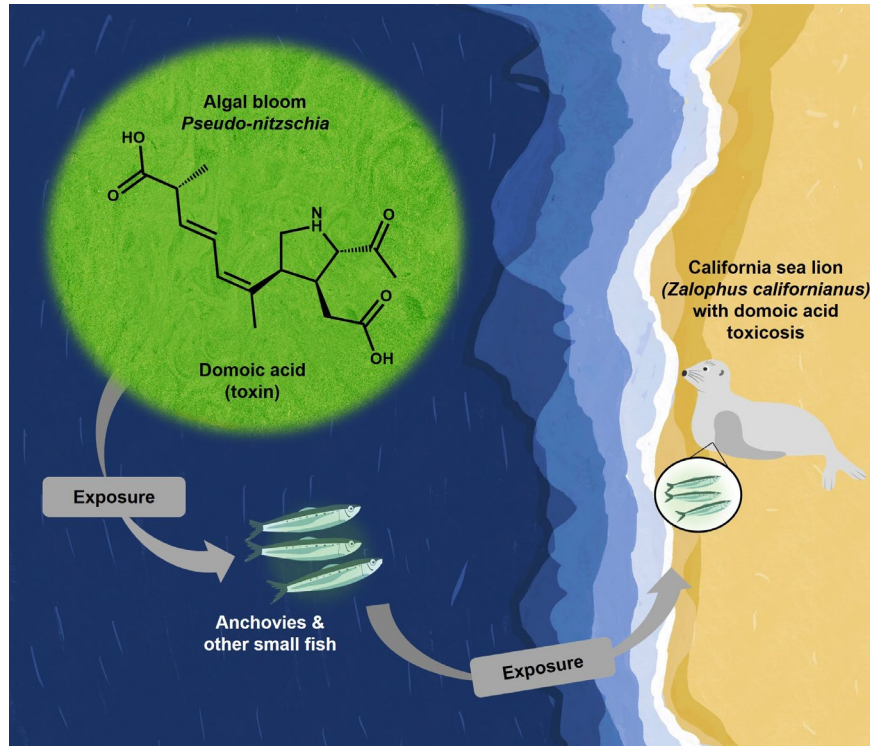
# California sea lions (*Zalophus californianus*)



<https://www.pier39.com/sealions/>



# DAT & Sea lions (*Zalophus Californianus*)



## Symptoms:

- Seizures
- Heart issues
- Fatal

## Challenges:

No diagnostic tools

Half-life of domoic acid is < 48 hours

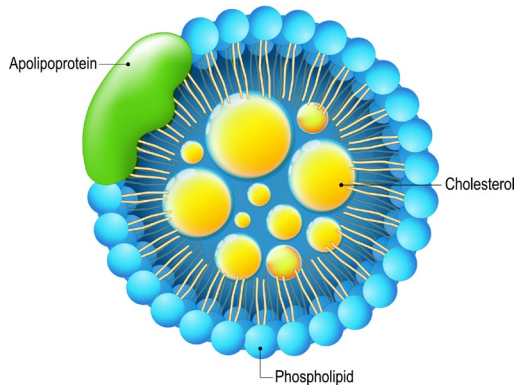
# Previous proteomic study



**Dr. Ben Neely,  
NIST**

## APOLIPOPROTEINS

(proteins that bind lipids to form lipoproteins)



> PLoS One. 2015 Apr 28;10(4):e0123295. doi: 10.1371/journal.pone.0123295. eCollection 2014.

## Proteomic Analysis of Plasma from California Sea Lions (*Zalophus californianus*) Reveals Apolipoprotein E as a Candidate Biomarker of Chronic Domoic Acid Toxicosis

Benjamin A Neely <sup>1</sup>, Jason A Ferrante <sup>2</sup>, J Mauro Chaves <sup>1</sup>, Jennifer L Soper <sup>3</sup>, Jonas S Almeida <sup>4</sup>, John M Arthur <sup>5</sup>, Frances M D Gulland <sup>3</sup>, Michael G Janech <sup>6</sup>

**Hypothesis:** Apolipoprotein dysregulation indicates that lipids are informative of DAT

# Experimental design (lipidomics)

## Rescued Sea Lion Samples

14 DAT\*  
17 non-DAT  
13 males, 18 females  
All ages



\*Diagnosed based on histological exam of brain tissues & clinical signs

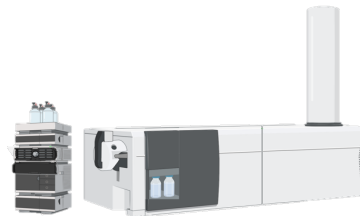
## Lipidomic Workflow

### Experimental

sea lion plasma



- Modified-Folch lipid extraction



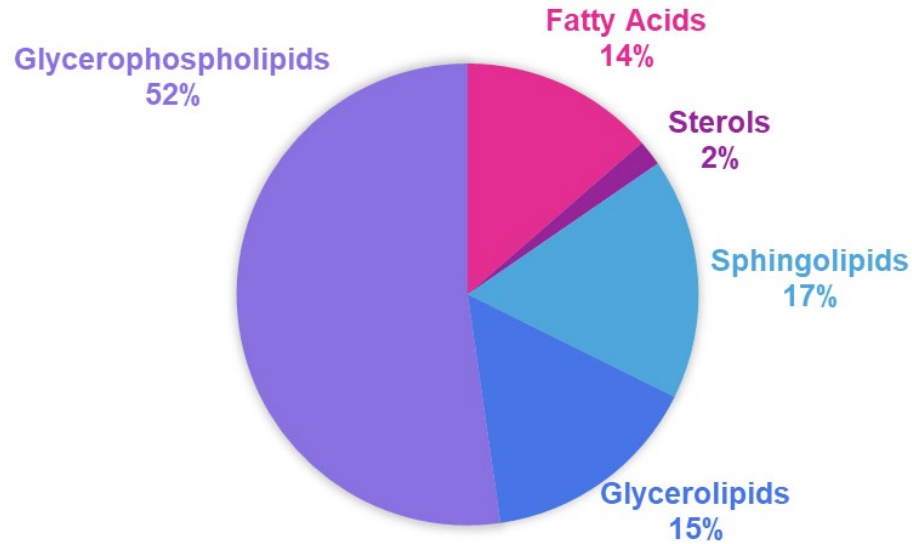
- 37 min LC-Drift Tube IMS-CID-MS
- m/z 50 to 1700

### Data Analysis

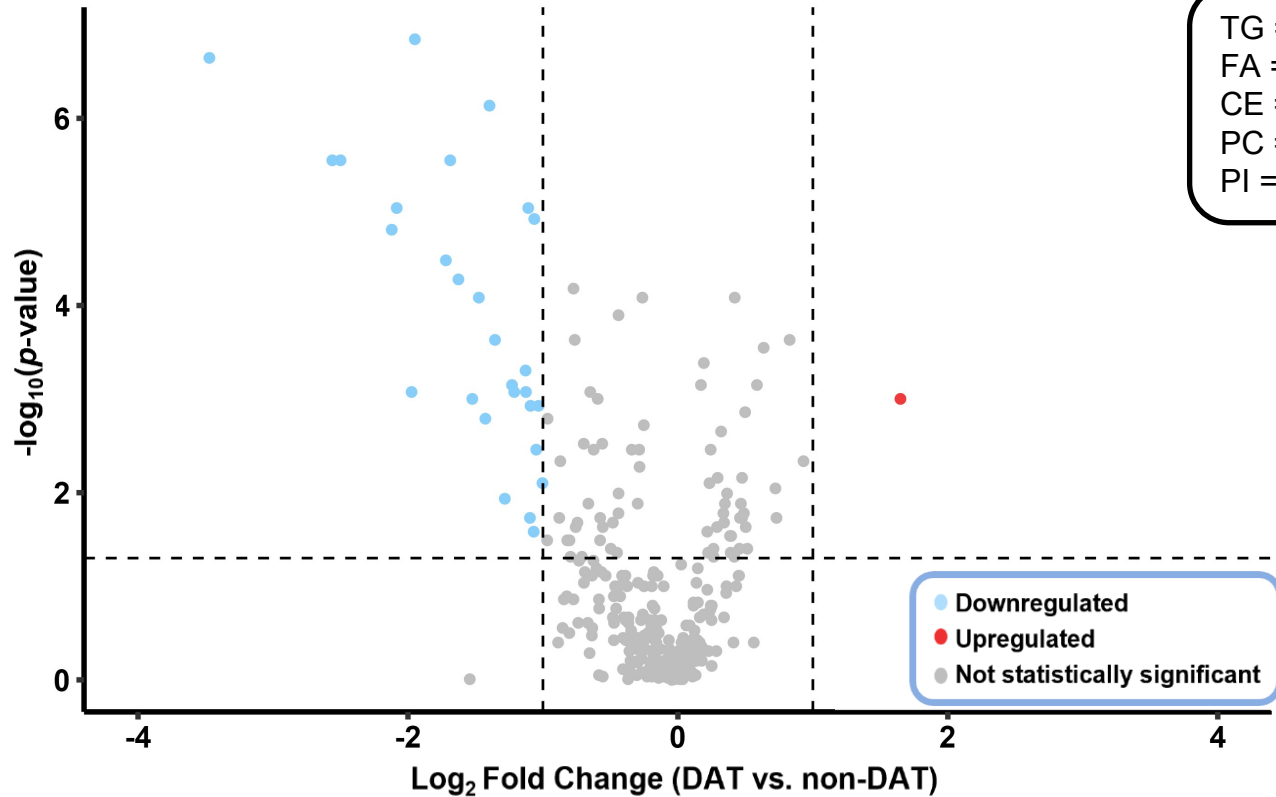


- Suspect Screening
- Statistics
- Data visualization

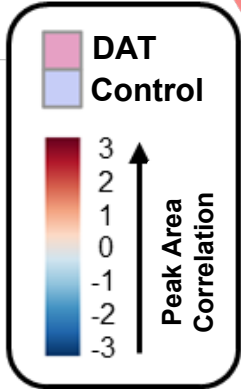
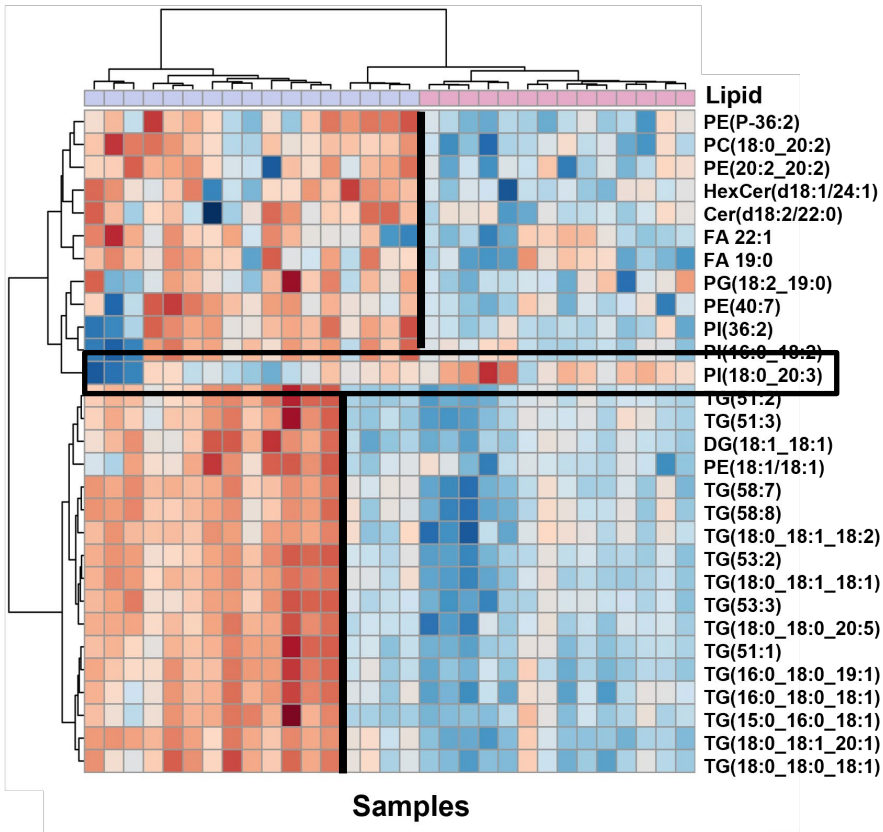
Detected Lipids: 331



# Significance testing

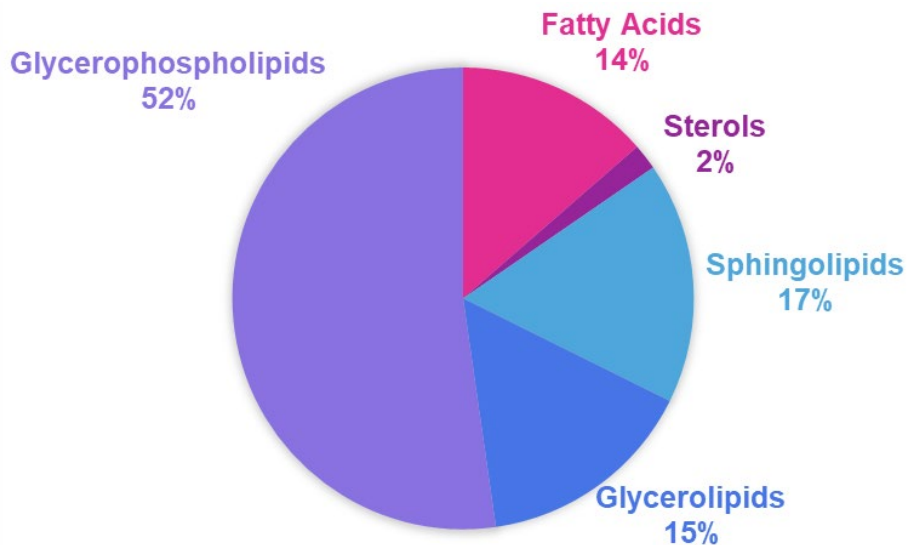


# Hierarchical clustering results

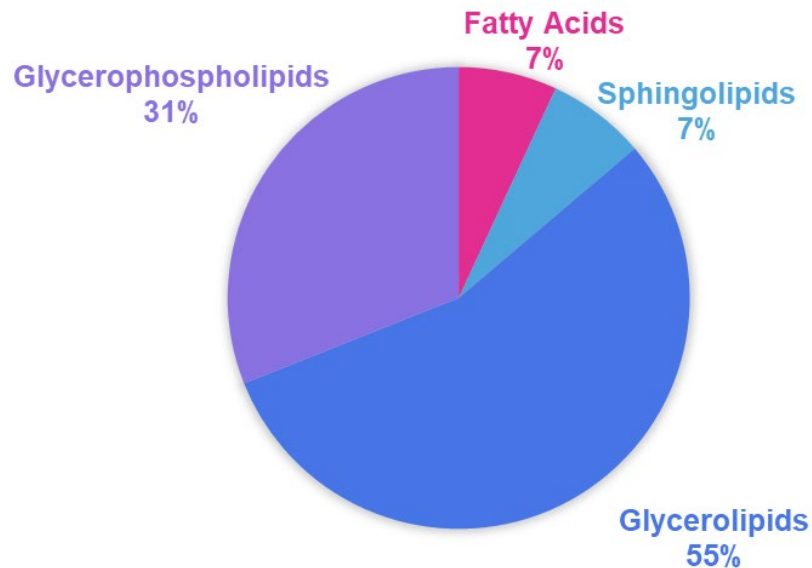


# Lipid category trends

**Detected Lipids: 331**



**Significant Lipids: 29**



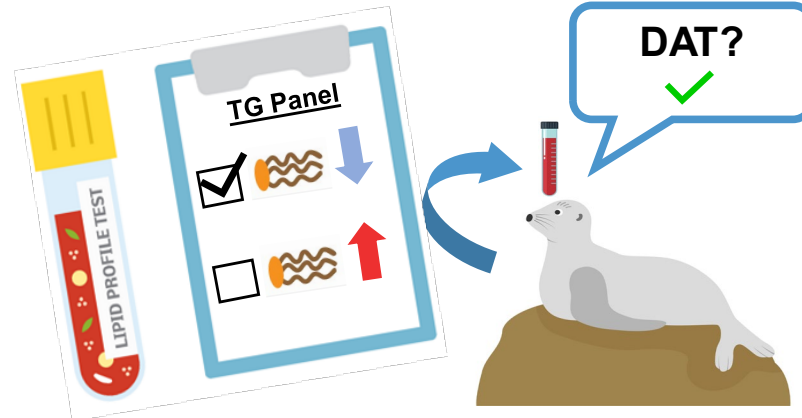


## Conclusions:

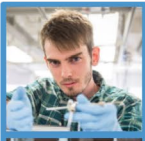
- 29 significant lipids & 50% are TGs
- **Triglycerides are dysregulated in DAT sea lions**
- This is a trend **unique to DAT** sea lions!
- Skyline was *essential* to these findings

## Future Directions:

- Develop a triglyceride blood panel to aid in diagnosis of DAT
- Test more samples to validate results



# Check out the Baker Lab at ASMS!



**James**

WP 475 - Cross Platform Assessment of Feature Detection as a Function of Measured Resolving Power in IMS



**Guozhi**

TP 366 - Lipids as Indicators of Successful Fecal Microbiota Transplantations



**Jessie**

TP 250 - Non-Targeted Analysis Feature Screening: Sample Classification and Feature Identification



**Jack**

WP - Utilizing High Resolution IMS-MS for Opioid Profiling  
Sunday – Agilent Users Meeting – Oligonucleotide Sequencing using IMS-MS/MS



**Anna**

ThOD - Comparing Target Screening, Suspect Screening, and Unknown Discovery Workflows in Non-Targeted Analyses of PFAS



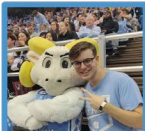
**Ashlee**

TP 101 - An Assessment of Serum Lipidomic Perturbations due to PFAS Exposures



**Amie**

TP 348 - Building a Multidimensional LC-IMS-MS Oxylipin Library  
Sunday - Skyline Users Meeting – Developing a Multidimensional Lipid Database



**Greg**

TP 103 - Evaluating Environmental Factors in Systemic Autoimmune Rheumatic Diseases



**Emily**

MP 196 - Assessing Antidepressant Pharmaceuticals in the Environment



**Sarah**

TP 055 - Evaluating Bone Marrow Lipidomic Changes and Their Potential Association with Alzheimer's Disease



**Kara**

TP 079 - Extraction Optimization and Non-Targeted Analyses of (PFAS) in Mammalian Milk



**Allison**

TP 116 - A Lipidomic Exploration of the Impact of PFOA Exposure on SARS-CoV-2 Infection



**Haley**

TP 115 - Evaluating Lipidomic Changes in Mice Exposed to Wildfire-Relevant Smoke from Different Fuel and Burn Conditions

Skyline used in the project

# Acknowledgements

## Baker Lab Members

### Principal Investigator

- Dr. Erin S. Baker

### Research Assistant Professor

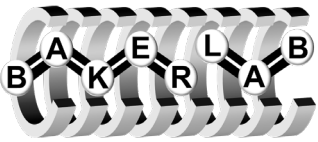
- Prof. James Dodds

### Postdoctoral Researchers

- Dr. Guozhi Zhang
- Dr. Jessie Chappel

### Graduate Students

- Jack Ryan
- Anna Boatman
- Ashlee Falls
- Greg Kudzin
- Allison Fry
- Emily Crawford
- Haley Jostes
- Kara Joseph
- Sarah Clark



### Past Graduates with Significant Contributions

- Dr. Karen Butler
- Dr. Melanie Odenkirk
- Dr. Kaylie Kirkwood Donelson
- Dr. Nancy Lee Alexander

### Contributing Undergraduate Students

- Iliana Claudio (alum)
- Quentin DuVal

## Sources of Funding

- NIEHS P42 ES027704 & P42 ES031009
- NIGMS RM1 GM145416 & R01 GM141277
- US EPA



@BakerLabMS   
 @AmieSolosky 

## Collaborators

### U Washington

- Mike MacCoss
- Brendan MacLean

### NIST

- Ben Neely

### College of Charleston

- Alison Bland
- Michael Janech

### University of California Davis

- Frances M.D. Gulland

