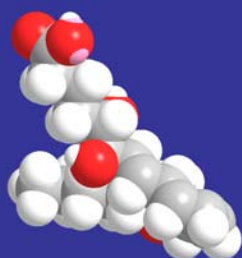
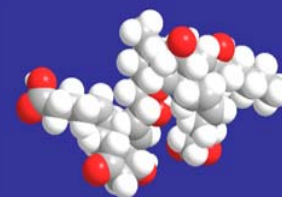


www.lipidmaps.org

LIPID MAPS Lipidomics Workshop

April 18, 2009



Eicosanoids

Richard Harkewicz



Departments of Pharmacology, Chemistry and Biochemistry



The University of California, San Diego



La Jolla, CA

Other LIPID MAPS Eicosanoid Core Members:

Aaron Armando

Matthew Buczynski

Edward A. Dennis
(Core Director)

Alexander Andreyev

Darren Dumlao

Joshua Brooks

Oswald Quehenberger

Outline

- A) Brief description of eicosanoids
- B) Sample preparation/extraction
- C) Analytical methodology: LC-MS
- D) Library of eicosanoid standards
- E) Chiral chromatography - *enzymatic or non-enzymatic?*
- F) GC-MS methodology for free fatty acids
- G) DIMPLES/MS: A stable isotope substrate labeling strategy enabling the search for novel eicosanoids
- H) Comparison of LIPID MAPS eicosanoid approach with others in literature
- I) Future plans

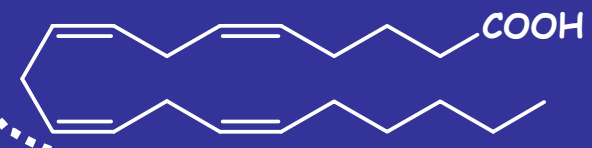
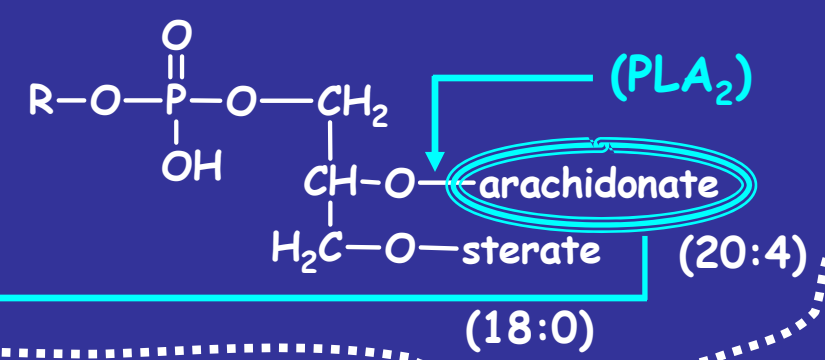
contact: rharkevicz@ucsd.edu

stimulation
(LPS or
Kdo₂-Lipid A)
(outside
the cell)

(inside
the cell)

release of
phospholipase A₂
(PLA₂)

membrane phospholipid
(R = choline or ethanolamine)

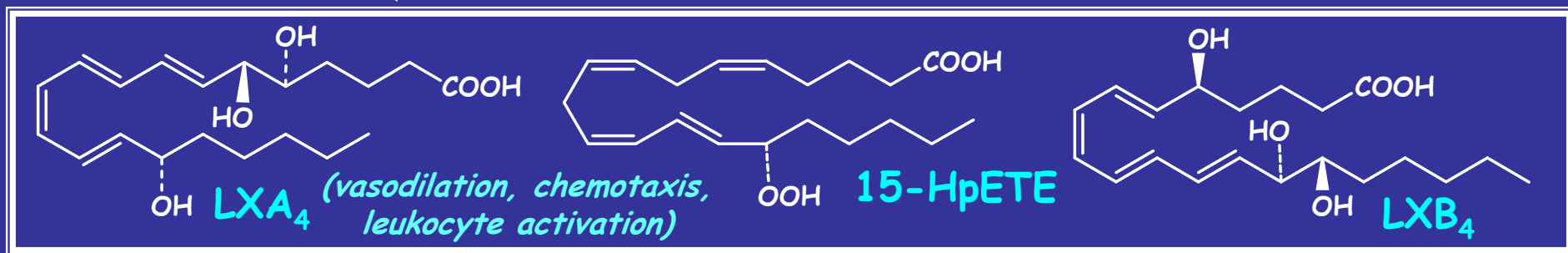
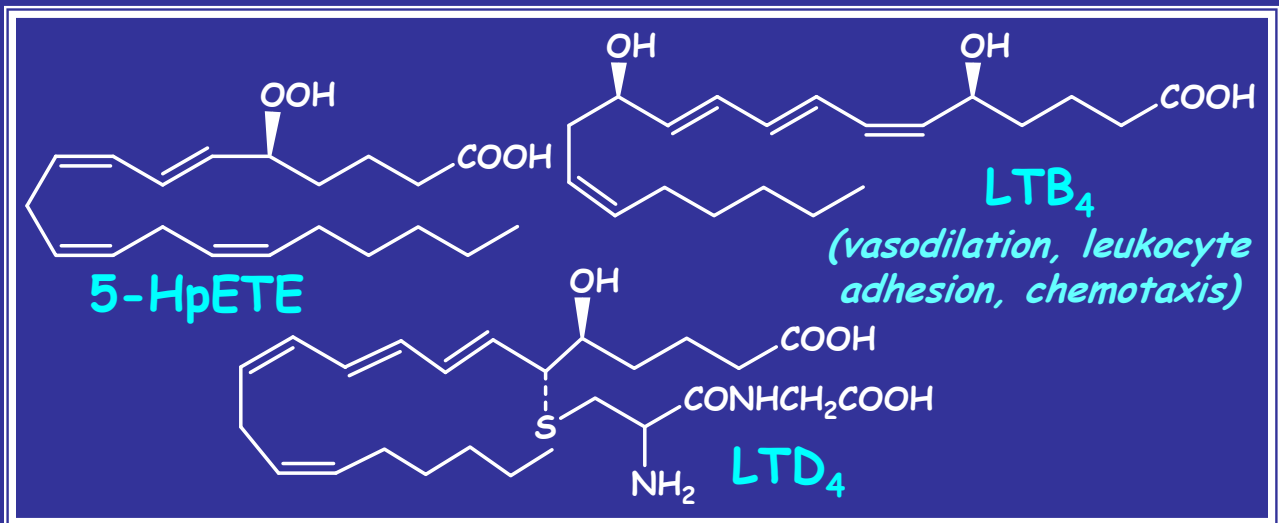
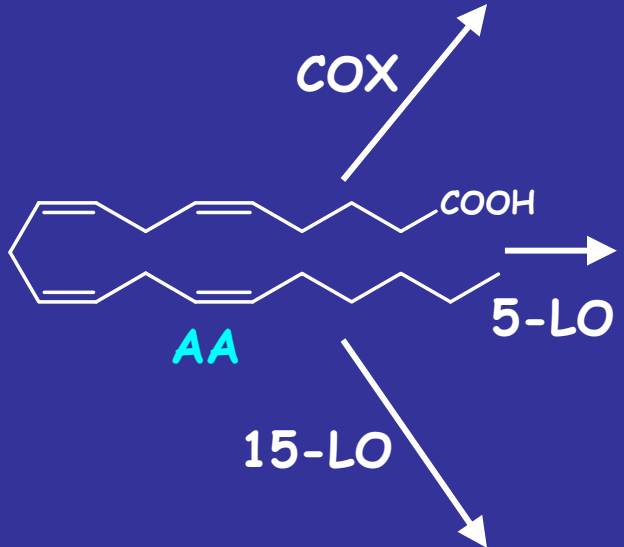
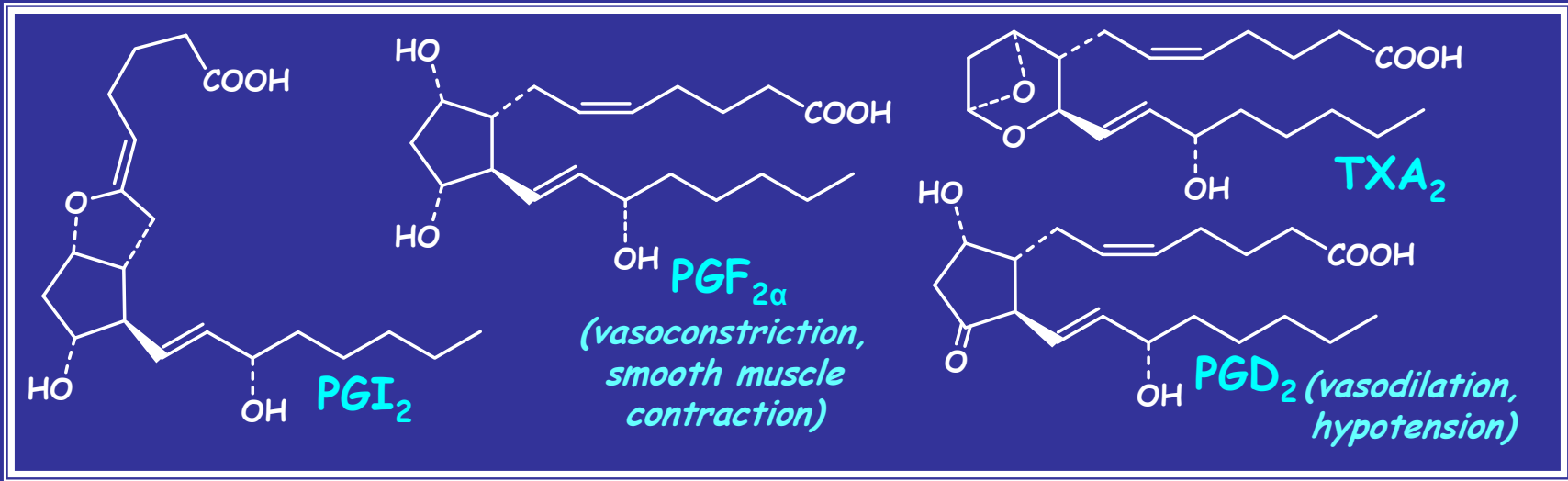


free eicosatetraenoic acid (from the Greek *eicosa* for 20)
AKA arachidonic acid (20:4, n=6)

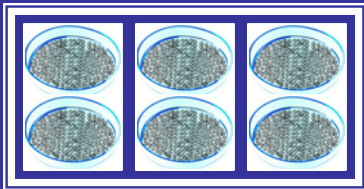
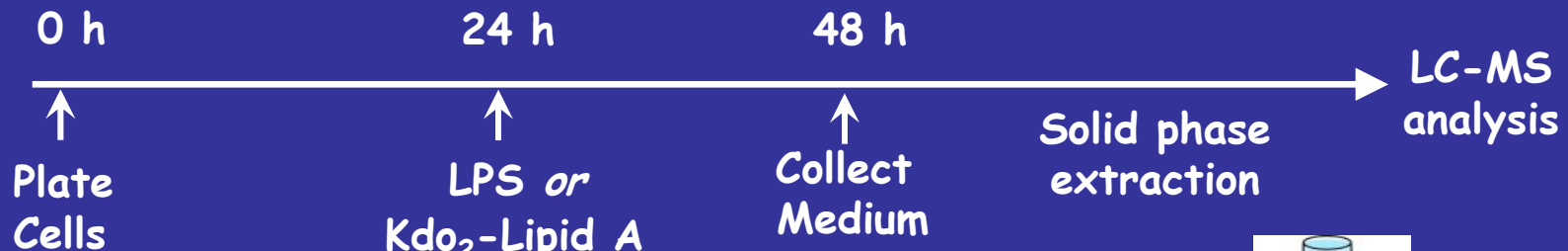
Eicosanoids - powerful mediators involved in inflammation that are derived from arachidonic acid and act in autocrine and paracrine fashion (signal at or immediately adjacent to their site of synthesis)

- Once they are made, they are quickly secreted from the cell
- Transcription factor: Can also enter cell's nucleus, binding and activating nuclear receptors

Sometimes referred to as *autocoids* or *local hormones*



sample preparation and extraction



1. Plate cells
- ↓ 24 h
2. Stimulate (LPS or Kdo)
- ↓ 24 h
3. Remove medium (≈2 ml per well) (≈2e6 cells per well)



1. Spike medium with deuterated internal standards (10 ng/100 µl)
2. Add EtOH to obtain 10% EtOH medium solution
3. Centrifuge 5 min @ 3000 rpm



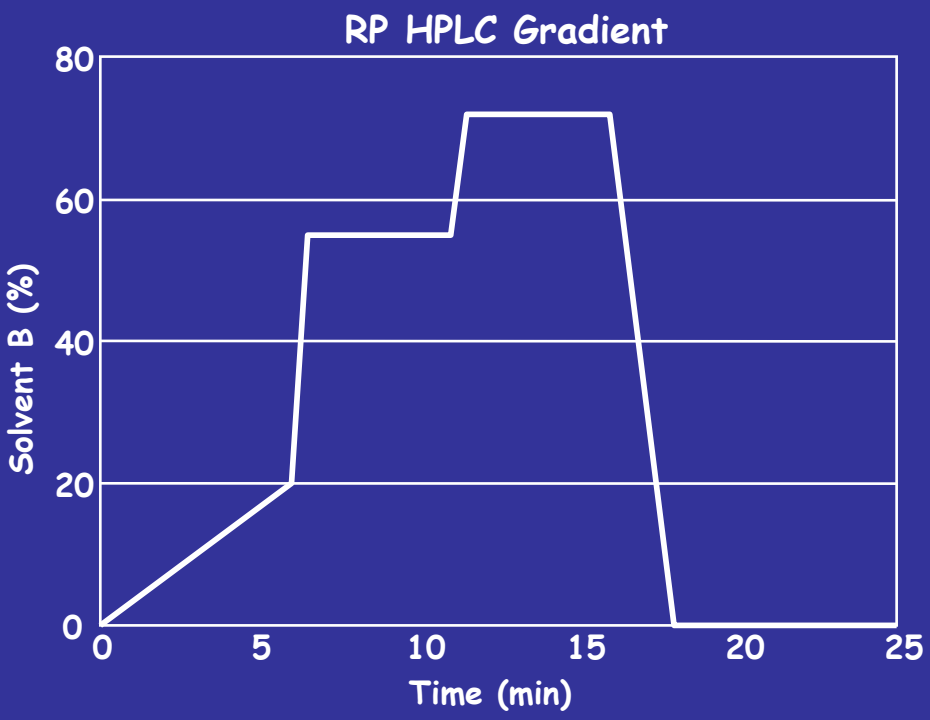
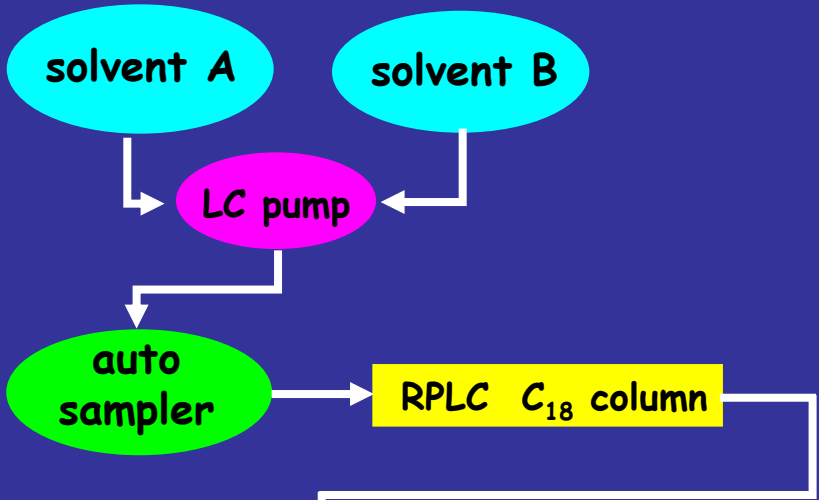
1. Solid phase extraction (Phenomenex® Strata-X 8B-S100-UBJ)
2. Prewash SPE column with 2 ml MeOH then 2 ml H₂O
3. Apply sample to column then wash with 2 ml 10% MeOH
4. Elute eicosanoids with 1 ml MeOH
5. Dry under vacuum and resuspend in 100 µl HPLC solvent A

column
Vydac® 201TP52
2.1 mm X 250 mm

Solvent A
H₂O/ACN/formic acid:
63/37/0.02

Flow rate
300 µl/min

Solvent B
ACN/IPA:
50/50



4000 QTRAP™
LC/MS/MS System

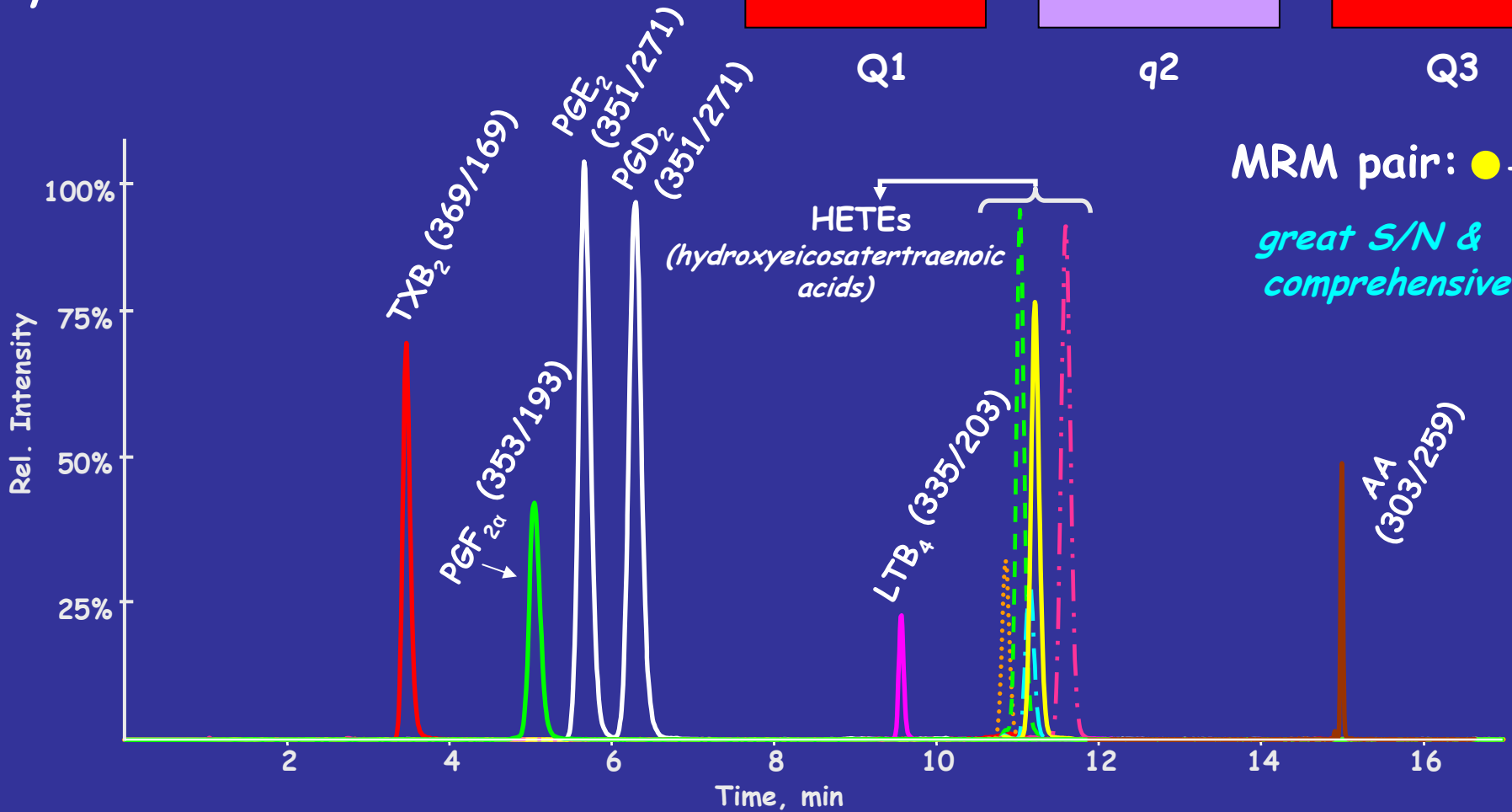
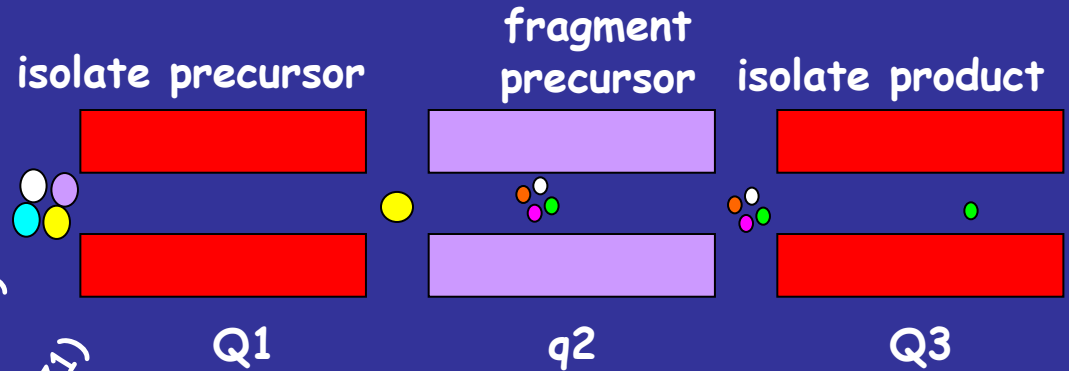
Applied Biosystems® 4000 QTrap
Quadrupole mass spectrometer

Eicosanoid Standard Mixture

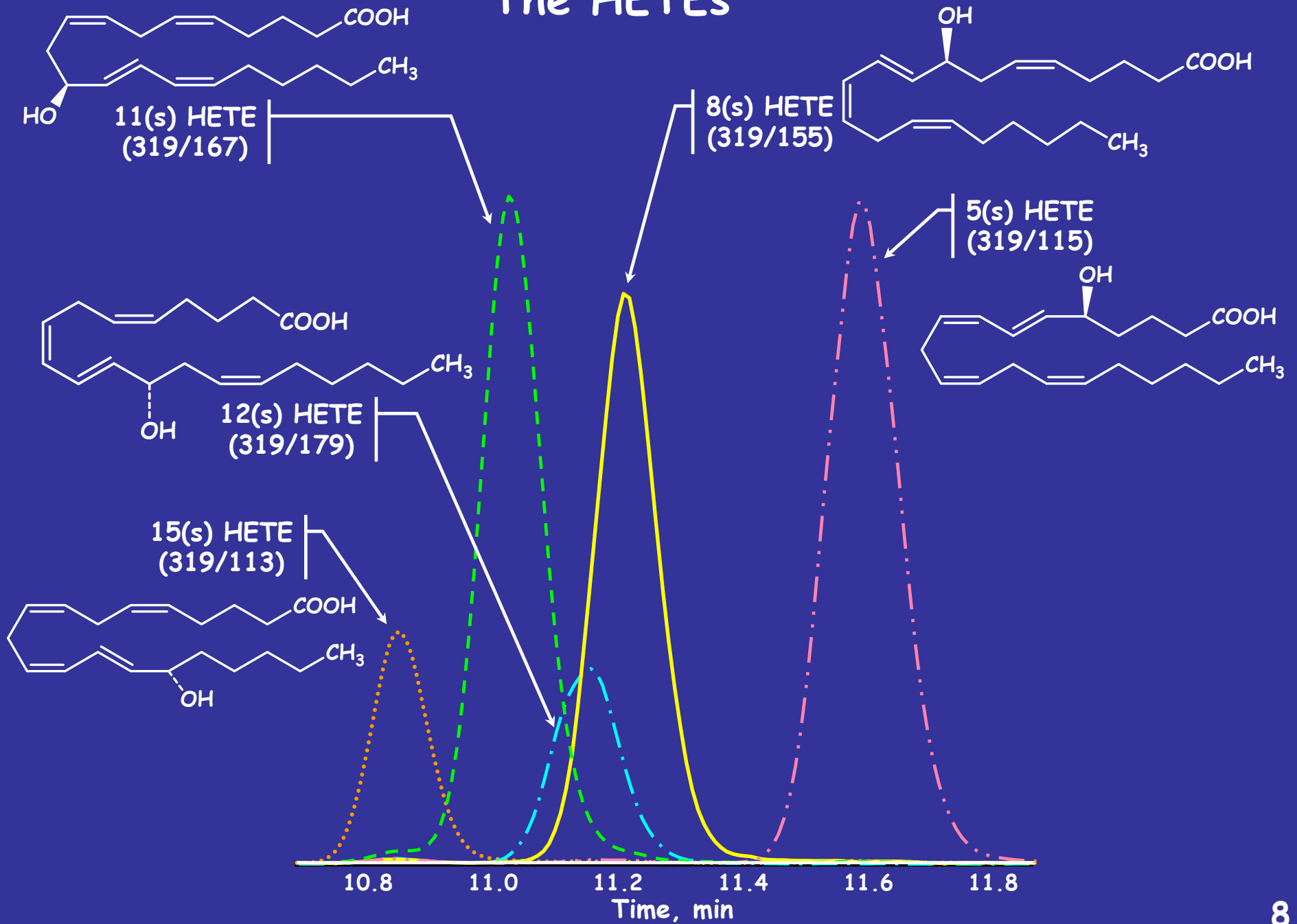
LC/MS - MRM

Multiple Reaction Monitoring
"specialized" form of MSMS

Applied Biosystems Analyst 1.5
"scheduled MRM" allowing >1000 MRM pairs



The HETEs



Library of Eicosanoid Standards (>200)

(<http://www.lipidmaps.org>)

Provides:

- ① Chemical structure in ChemDraw® format
- ① LC and MS protocols
- ① MSMS fragmentation spectra
- ① LC retention times for given set of conditions
- ① Web-link to Cayman Chemical for each eicosanoid

LIPID Metabolites And Pathways Strategy

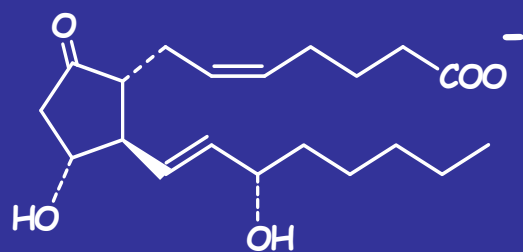
Lipid Standards: Eicosanoids

View LC/MS/MS protocols and retention-time data.
Clicking on a LM_ID displays the structure in GIF (or ChemDraw) format.
Clicking on a MS/MS value (nominal mass of precursor ion in negative-ion mode) displays the fragmentation spectrum, including structures of principal product ions.
Clicking on a CAYMAN_ID value displays the Cayman catalog website page.
Clicking on a Ref value displays a literature reference(s) pertaining to identification of fragment structures.

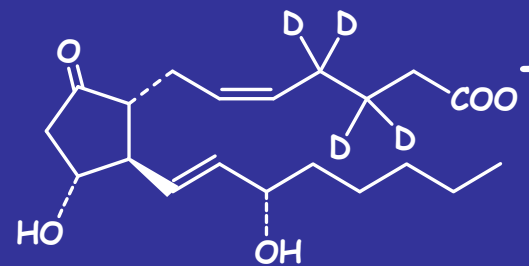
LM_ID	Name	Systematic Name	Cayman ID	MS/MS	Ref
LMFA01030001	AA	5Z,8Z,11Z,14Z - eicosatetraenoic acid	90010	303((M-H)-)	-
LMFA01030003	AA - d8	5Z,8Z,11Z,14Z - eicosatetraenoic acid (5,6,8,9,11,12,14,15 - d8)	390010	311((M-H)-)	-
LMFA03010001	6k - PGF1α	9S,11R,15S - trihydroxy - 6 - oxo - 13E - prostenoic acid	15210	369((M-H)-)	6
LMFA03010002	PGF2α	9S,11R,15S - trihydroxy - 5Z,13E - prostadienoic acid	16010	353((M-H)-)	1,2,3
LMFA03010003	PGE2	11R,15S - dihydroxy - 9 - oxo - 5Z,13E - prostadienoic acid	14010	351((M-H)-)	4
LMFA03010004	PGD2	9S,15S - dihydroxy - 11 - oxo - 5Z,13E - prostadienoic acid	12010	351((M-H)-)	-
LMFA03010006	PGF2α - d4	9S,11R,15S - trihydroxy - 5Z,13E - prostadienoic acid (3,3,4,4 - d4)	316010	357((M-H)-)	-
LMFA03010007	PGD2 - d4	9S,15S - dihydroxy - 11 - oxo - 5Z,13E - prostadienoic acid (3,3,4,4 - d4)	312010	355((M-H)-)	-
LMFA03010008	PGE2 - d4	11R,15S - dihydroxy - 9 - oxo - 5Z,13E - prostadienoic acid (3,3,4,4 - d4)	314010	355((M-H)-)	-
LMFA03010009	PGG2	9S,11R - epidioxy - 15S - hydroperoxy - 5Z,13E - prostadienoic acid	17010	367((M-H)-)	-
LMFA03010010	PGH2	9S,11R - epidioxy - 15S - hydroxy - 5Z,13E - prostadienoic acid	17020	351((M-H)-)	-
LMFA03010011	2,3 - Dinor - 11β - PGF2a	9S,11S,13S - trihydroxy - 2,3 - dinor - 5Z,13E - prostadienoic acid	16530	325((M-H)-)	-
LMFA03010012	6keto - PGE1	11R,15S - dihydroxy - 6,9 - dioxo - 13E - prostenoic acid	13260	367((M-H)-)	-
LMFA03010013	6,15 - diketo - 13,14 - dihydro - PGF1α	9S,11R - dihydroxy - 6,15 - dioxo - 13E - prostenoic acid	15270	369((M-H)-)	-
LMFA03010014	20 - hydroxy - PGE2	11R,15S,20 - trihydroxy - 9 - oxo - 5Z,13E - prostadienoic acid	14950	367((M-H)-)	-
LMFA03010015	PGF2α - EA	N - (9S,11R,15S - trihydroxy - 5Z,13E - prostadienoyl) - ethanalamine	16013	396((M-H)-)	-
LMFA03010016	PGE2 - EA	N - (11R,15S - dihydroxy - 9 - oxo - 5Z,13E - prostadienoyl) - ethanalamine	14012	394((M-H)-)	-
LMFA03010017	PGD2 - EA	N - (9S,15S - dihydroxy - 11 - oxo - 5Z,13E - prostadienoyl) - ethanalamine	12012	394((M-H)-)	-
LMFA03010018	PGB2	15S - hydroxy - 9 - oxo - 5Z,8(12),13E - prostatrienoic acid	11210	333((M-H)-)	-
LMFA03010019	PGJ2	15S - hydroxy - 11 - oxo - 5Z,8Z,13E - prostatrienoic acid	18500	333((M-H)-)	-
LMFA03010020	δ - 12 - PGJ2	15S - hydroxy - 11 - oxo - 5Z,9,12E - prostatrienoic acid	18550	333((M-H)-)	-
LMFA03010021	15 - deoxy - δ - 12,14 - PGJ2	11 - oxo - 5Z,9,12E,14Z - prostatetraenoic acid	18570	315((M-H)-)	-
LMFA03010022	13,14 - dihydro - 15 - keto - PGD2	11,15 - dioxo - 9S - hydroxy - 5Z - prostenoic acid	12610	351((M-H)-)	-
LMFA03010023	PGK2	9,11 - dioxo - 15S - hydroxy - 5Z,13E - prostadienoic acid	18900	349((M-H)-)	-
LMFA03010024	19R - hydroxy - PGE2	9 - oxo - 11R,15S,19R - trihydroxy - 5Z,13E - prostadienoic acid	14910	367((M-H)-)	-
LMFA03010025	PGF28	9R,11R,15S - trihydroxy - 5Z,13E - prostadienoic acid	16410	353((M-H)-)	-
LMFA03010026	15 - keto - PGE2	9S,11R - dihydroxy - 15 - oxo - 5Z,13E - prostadienoic acid	16700	351((M-H)-)	-

The eicosanoid library provides information from which comprehensive methods can be created and used to survey eicosanoid release from stimulated cells

use of deuterium labeled internal standards allows absolute quantitation for a number of eicosanoids

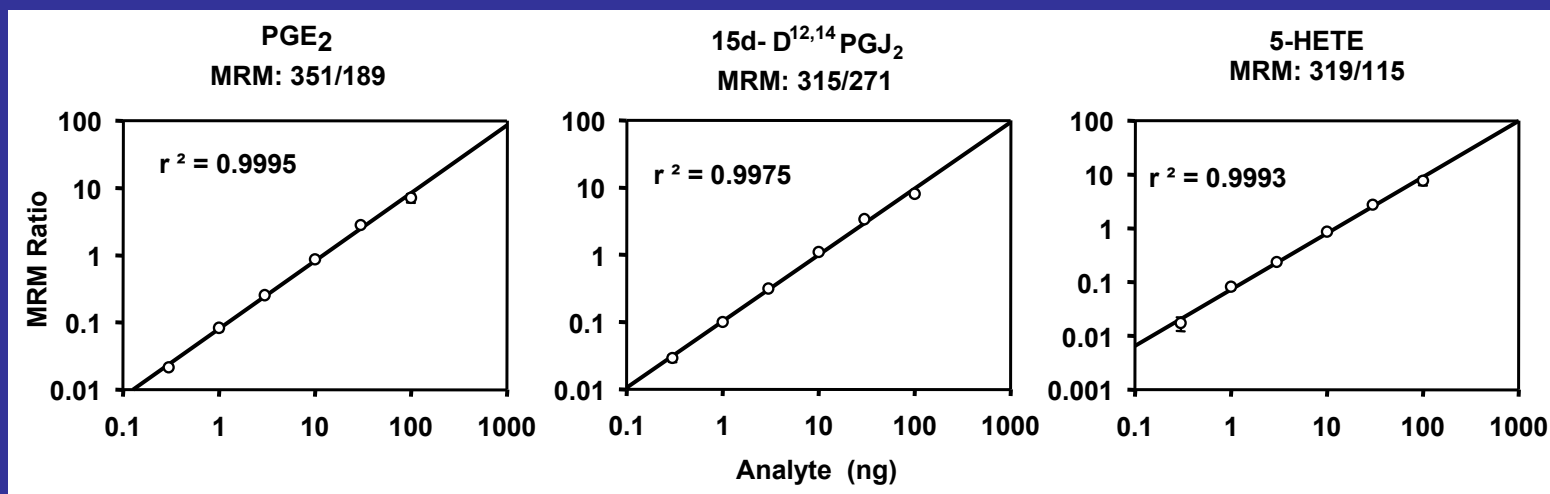


PGE₂ [M-H]⁻ = 351



PGE₂ - d4 [M-H]⁻ = 355

example of internal standard calibration curves used for quantitation



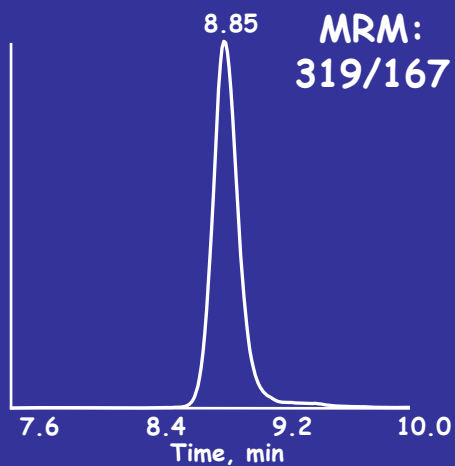
chiral chromatography aids in determining enzymatic vs. nonenzymatic origin

atmospheric pressure chemical ionization (APCI) used

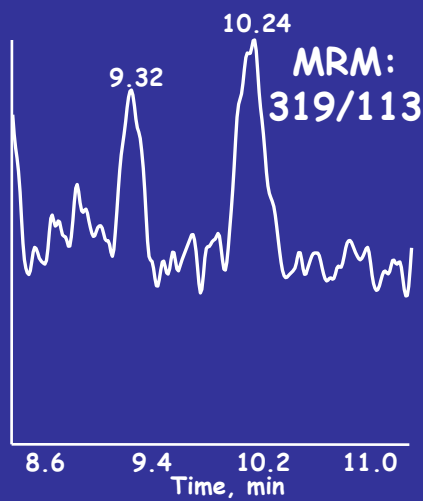
column
Chiralpak® AD-H
4.6 mm X 250 mm

Flow rate
500 µl/min

Kdo₂-Lipid A stimulated RAW cells

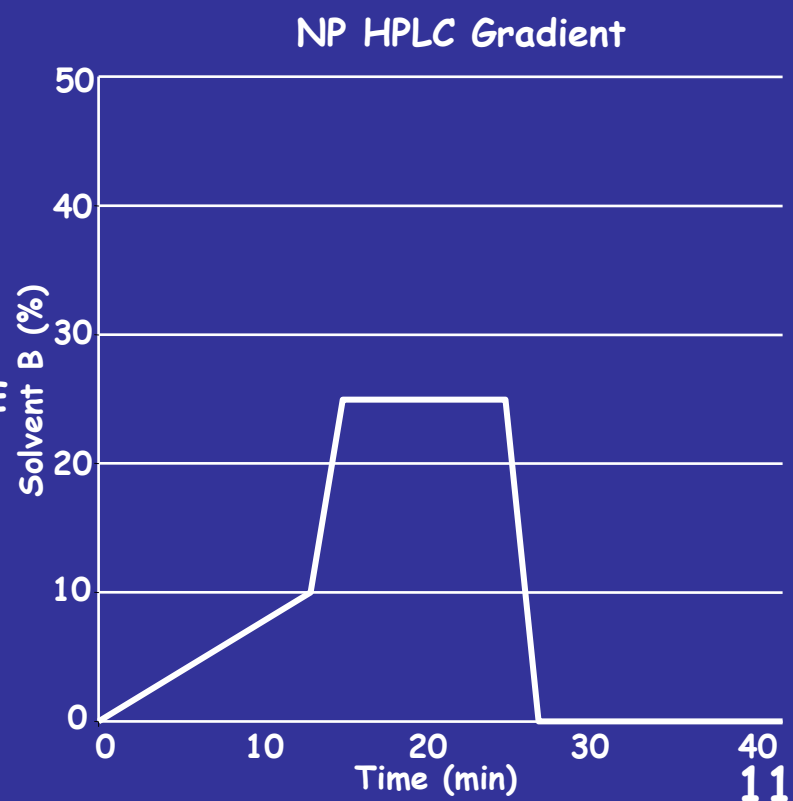
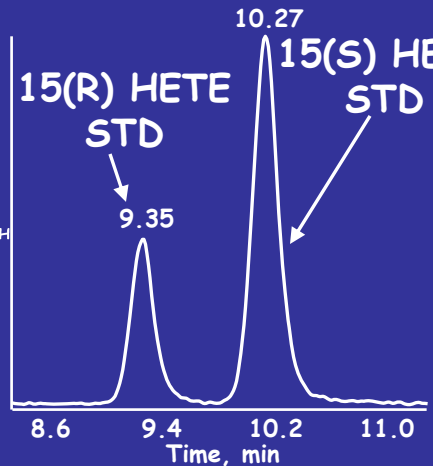
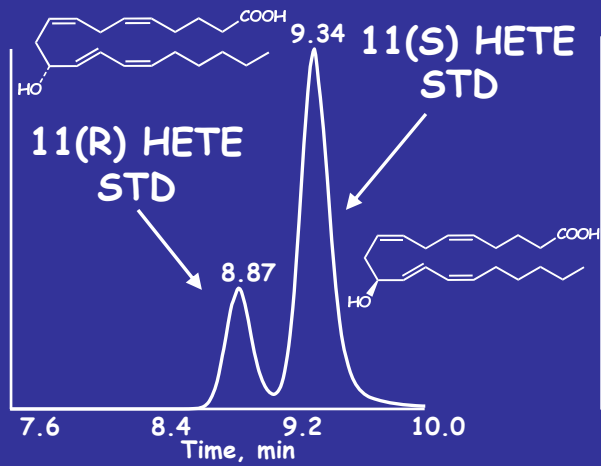


Kdo₂-Lipid A stimulated RAW cells



Solvent A
hexane/EtOH/H₂O/formic acid:
96/4/0.08/0.02

Solvent B
EtOH : 100



GC-MS analysis of free fatty acids

- Pentafluorobenzyl (PFB) derivatives of the free fatty acids are formed and negative chemical ionization employed. *Produces intact molecular anions - optimal for quantitation (using deuterium labeled standards).*
- Agilent 6890N gas chromatograph
- Agilent 5973 single quadrupole mass analyzer
- Selected ion monitoring (SIM) mode monitoring the $[M-H]^-$ anions.

analysis of essential fatty acids in macrophages

Quehenberger et al. (2008) Prostaglandins Leukot. Essent. Fatty Acids 79 pp. 123-129

Lauric acid (12:0)
Myristic acid (14:0)
Pentadecanoic acid (15:0)
Palmitic acid (16:0)
Palmitoleic acid (16:1)
Heptadecanoic acid (17:0)
Heptadecaenoic acid (17:1)
Stearic acid (18:0)
Oleic acid (18:1)
Linoleic acid (18:2)
 α -Linolenic acid (18:3)
 γ -Linolenic acid (18:3)
Stearidonic acid (18:4)
Arachidic acid (20:0)
Eicosadienoic acid (20:2)
11,14,17-Eicosatrienoic acid (20:3)
Bishomo- γ -linolenic acid (20:3)
5,8,11-Eicosatrienoic acid (20:3)
Arachidonic acid (20:4)
Eicosapentaenoic acid (20:5)
Behenic acid (22:0)
cis-Erucic acid (22:1)
Docosadienoic acid (22:2)
Docosatrienoic acid (22:3)
Docosatetraenoic acid (22:4)
Docosapentaenoic acid (22:5)
Docosahexaenoic acid (22:6)
Tricosanoic acid (23:0)
Lignoceric acid (24:0)
cis-Selacholeic acid (24:1)
Cerotic acid (26:0)

Are biologically significant eicosanoids being overlooked?

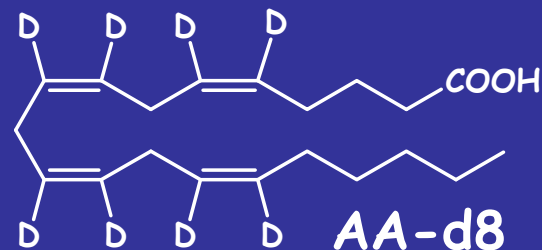
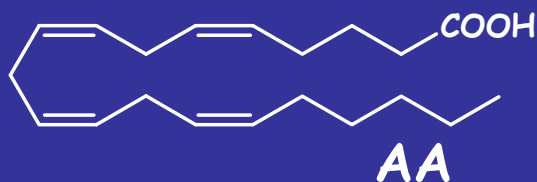
Are there species for which we have no prior knowledge of or expectation of their presence and, hence, no available MRM pairs required for their detection?

To address these concerns a mass spectral based stable isotope labeling strategy has been developed

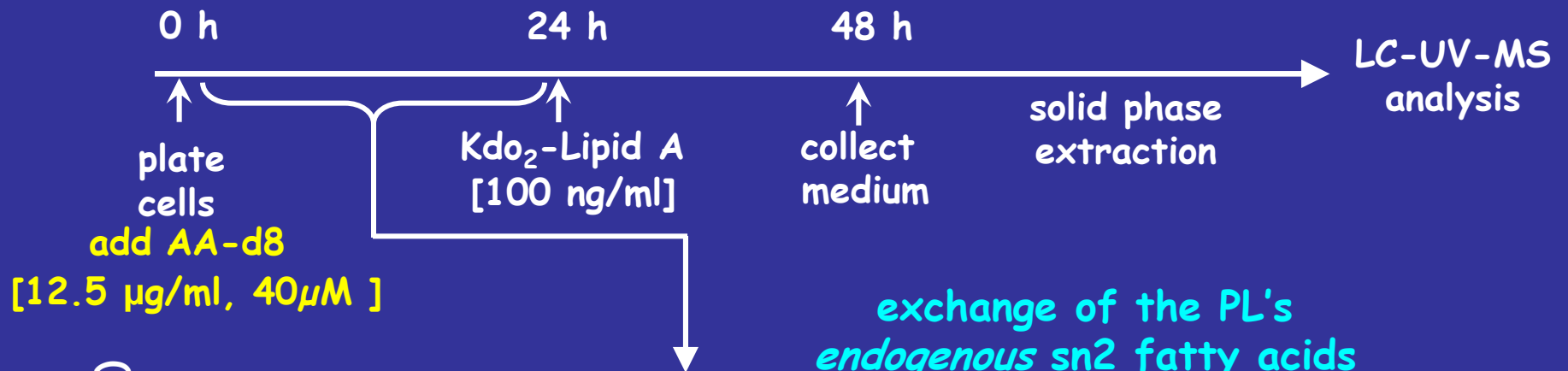
DIMPLES/MS: Diverse Isotope Metabolic Profiling of Labeled Exogenous Substrates using Mass Spectrometry

Harkewicz et al. (2007) J. Biol. Chem. 282 pp. 2899-2910

Incubation of cells in medium supplemented with deuterium-labeled arachidonic acid (AA-d8)

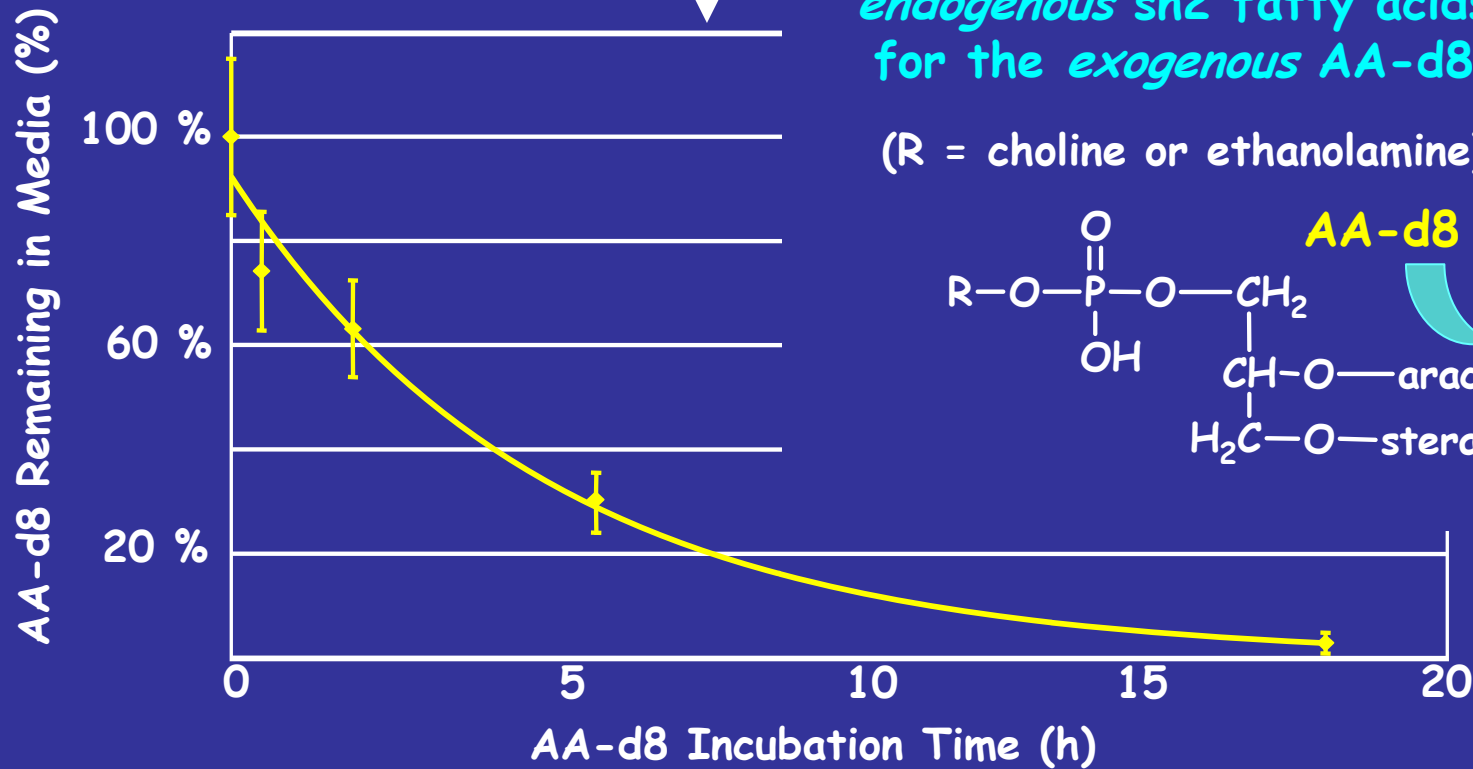
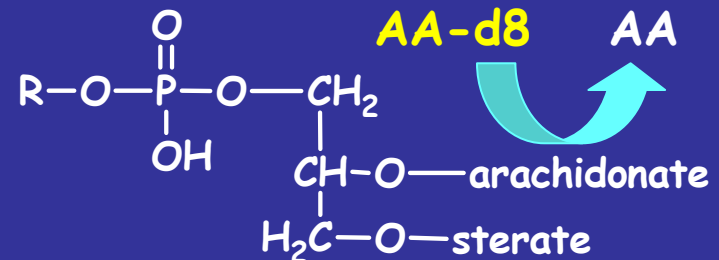


uptake and incorporation of AA-d8 by cells



exchange of the PL's endogenous sn2 fatty acids for the exogenous AA-d8

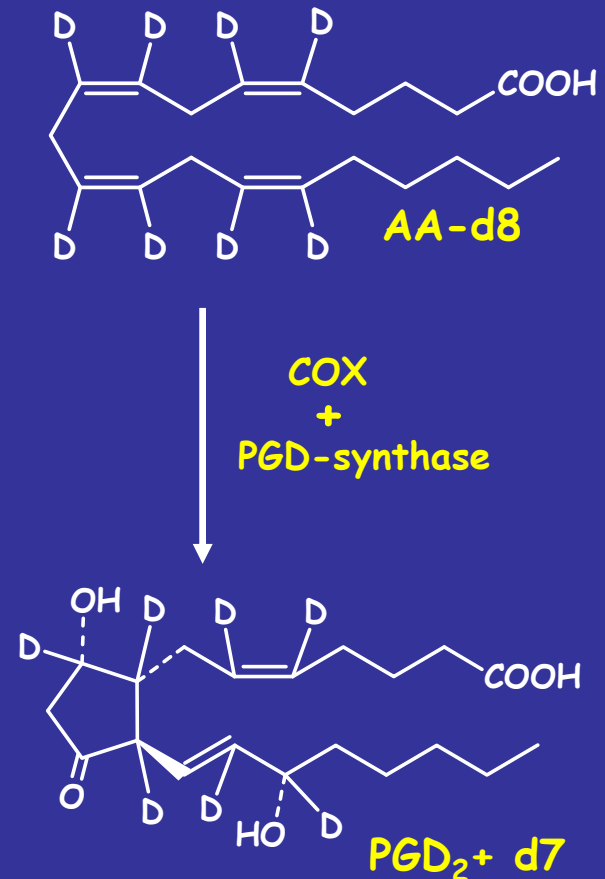
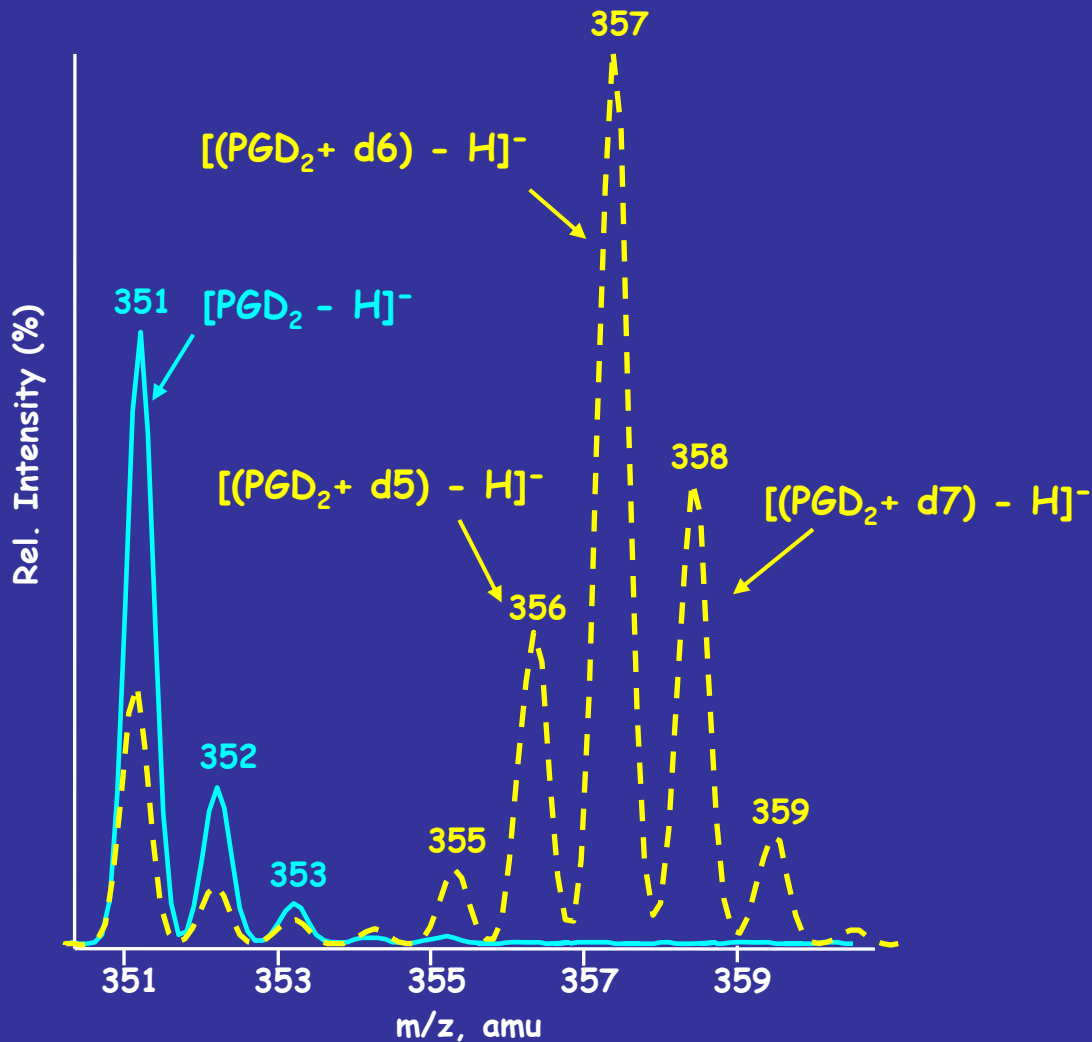
(R = choline or ethanolamine)



*so as not to be a
targeted analysis*
full-scan MS

*AA/AA-d8 mass spectral doublet
indicative of eicosanoid:
(1) AA origin
2) upregulated by Kdo*

———— 10% serum (fetal calf)
----- AA-d8 supplemented



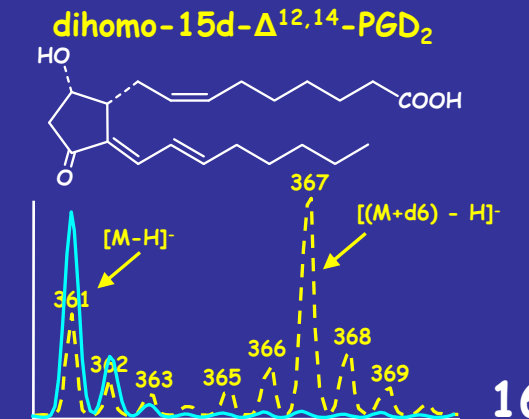
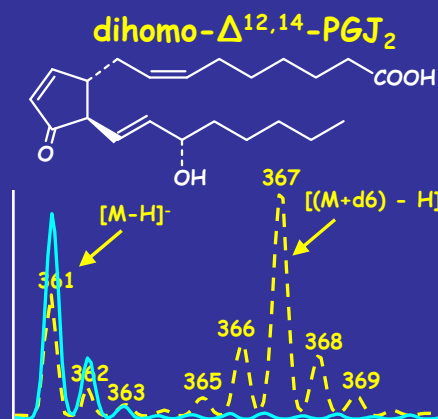
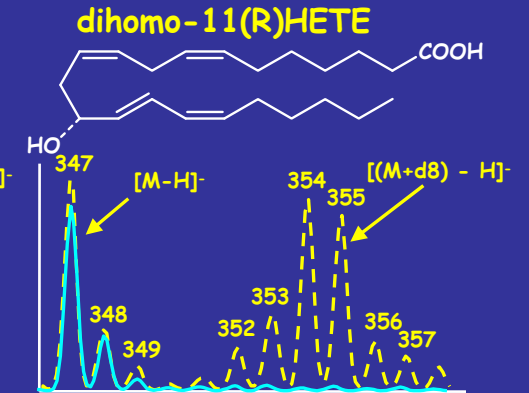
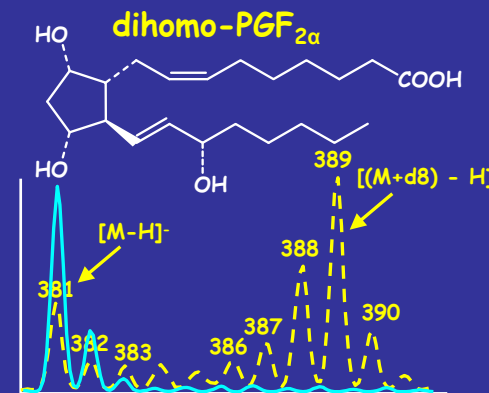
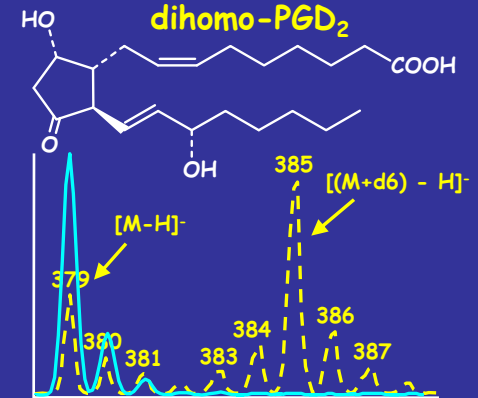
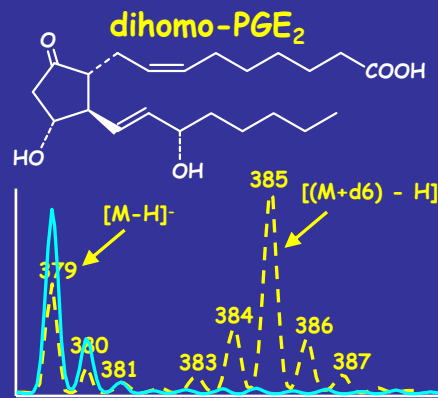
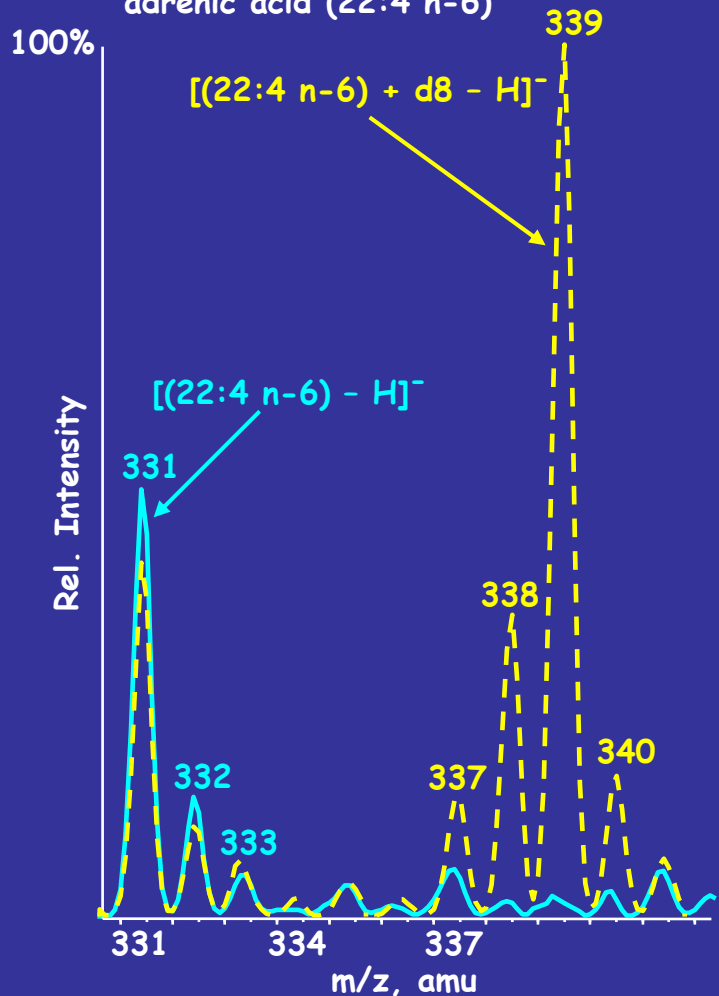
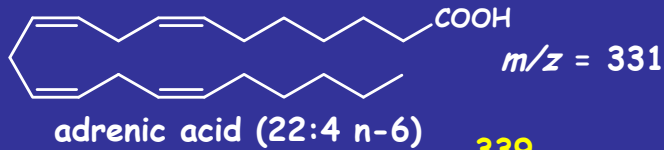
observed production of a class of dihomoprostaglandins

Harkewicz et al. (2007) *J. Biol. Chem.* **282** pp. 2899-2910

Kdo₂ - Lipid A stimulated RAW cells

———— Control (NO AA-d8)

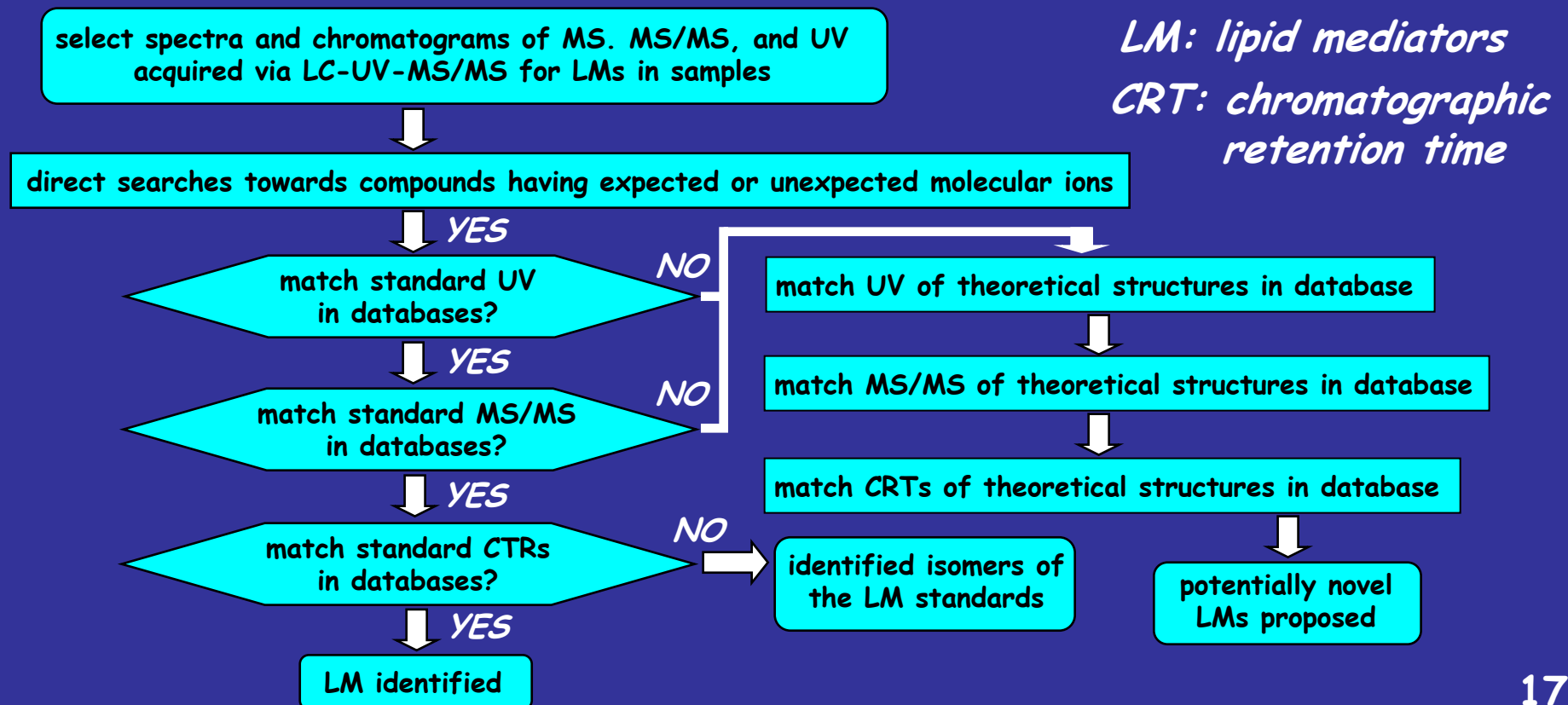
- - - - - AA-d8 supplemented



another approach to eicosanoid lipidomics and the search for novel eicosanoids

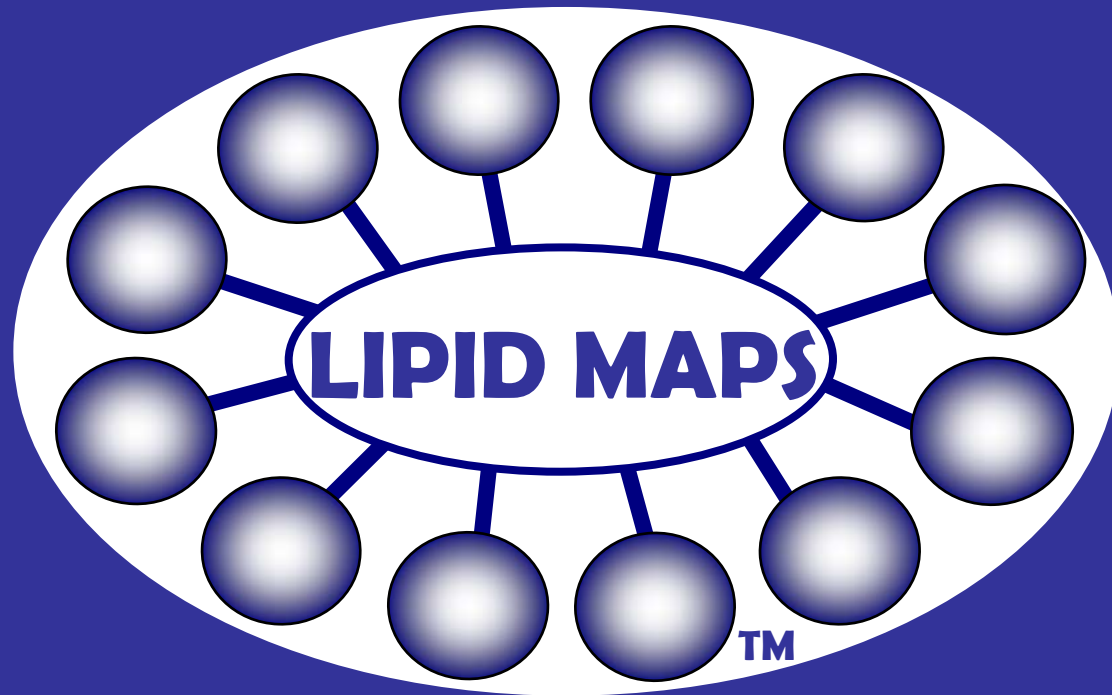
Serhan Lab developing theoretical databases and algorithms based on virtual UV absorption, tandem mass spectrometry and LC retention times for identifying potential eicosanoids .

Y. Lu et al. (2005) J. Lipid Res. 46 pp. 790-802



Future Plans

- Continue to expand eicosanoid library
- Incorporate UV detection and analyses into eicosanoid surveys
- Expand search for novel eicosanoids
- Similar studies with Ω -3 fatty acid supplementation
EPA (20:5 n-3) and DHA (22:6 n-3)



www.lipidmaps.org

Thank You