



Washington University in St. Louis

SCHOOL OF MEDICINE

**Experimental Molecular
Therapeutic Strategies for
Treating Barth Syndrome:
Elucidation of the Functional Role of the
Mitochondrial Lipidome**

Barth Syndrome Conference 2012

Michael Kiebish

Disclaimer

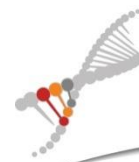
-All experiments and analysis were performed while at Washington University School of Medicine. The data and conclusions presented do not necessarily reflect the opinions of my current employer.



BERG
DIAGNOSTICS



BERG
PHARMA



BERG
BIOSYSTEMS

Overview

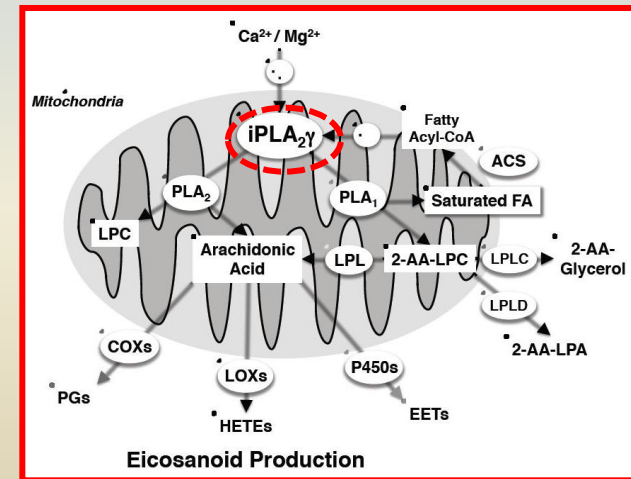
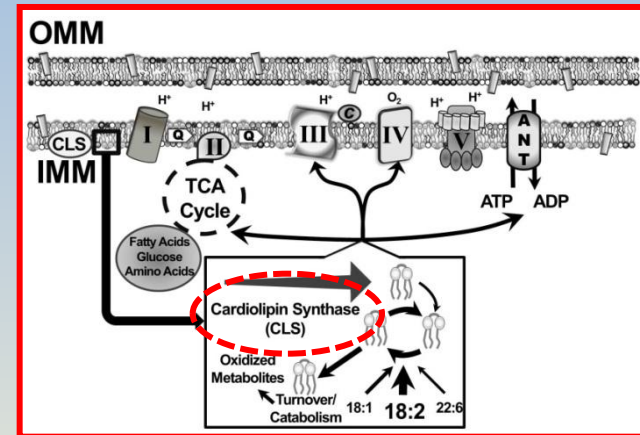
Barth Syndrome Mouse Model

Cardiac lipidomic, metabolic, and adaptive mechanisms

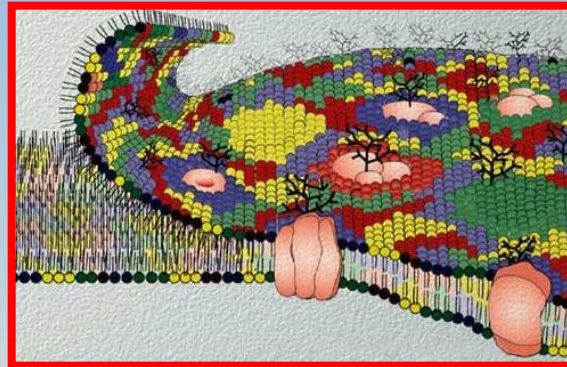
Effects of cardiac specific upregulation of cardiolipin synthase and the subsequent lipidomic and bioenergetic effects

Targeting phospholipases by transgenically expressing/ablating $iPLA_2\gamma$ and its subsequent lipidomic effects

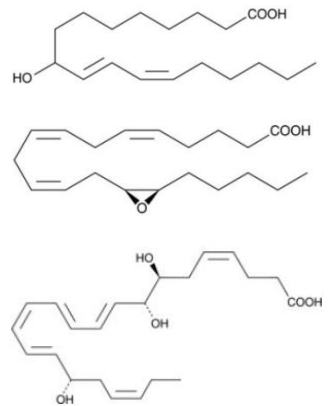
Novel interpretation of the role of the mitochondrial lipidome in health and disease as it relates to Barth syndrome



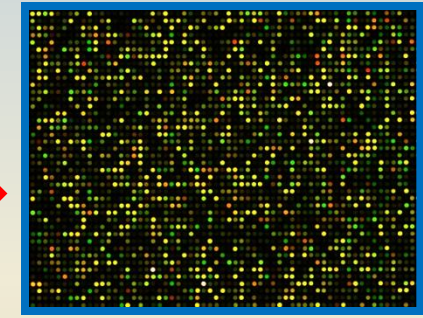
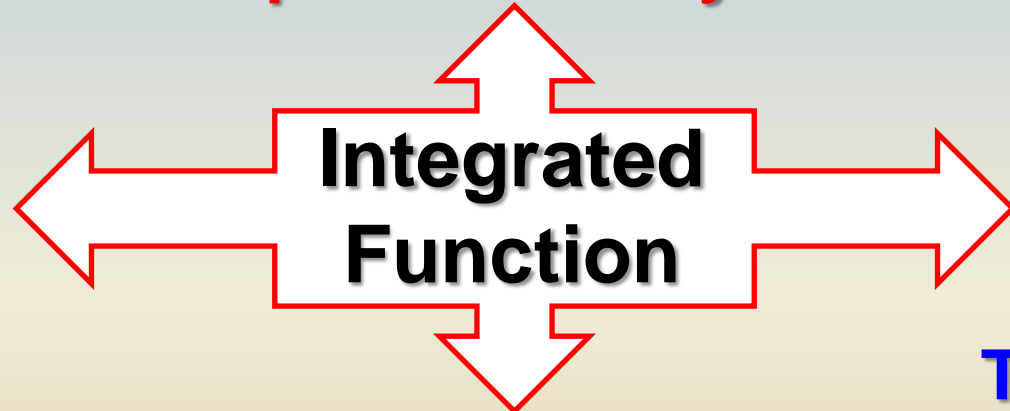
Multi-Omic Integrated Strategy to Elucidate Functional Changes in Pathophysiology



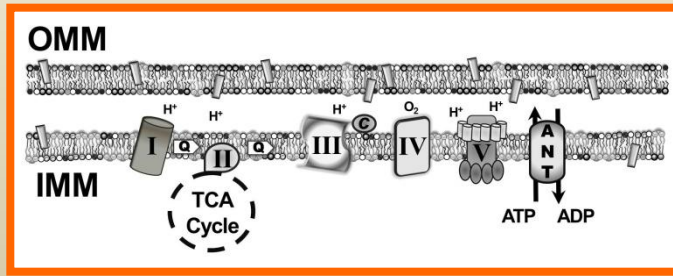
Lipidomic Analysis



Signaling Lipidomic Analysis



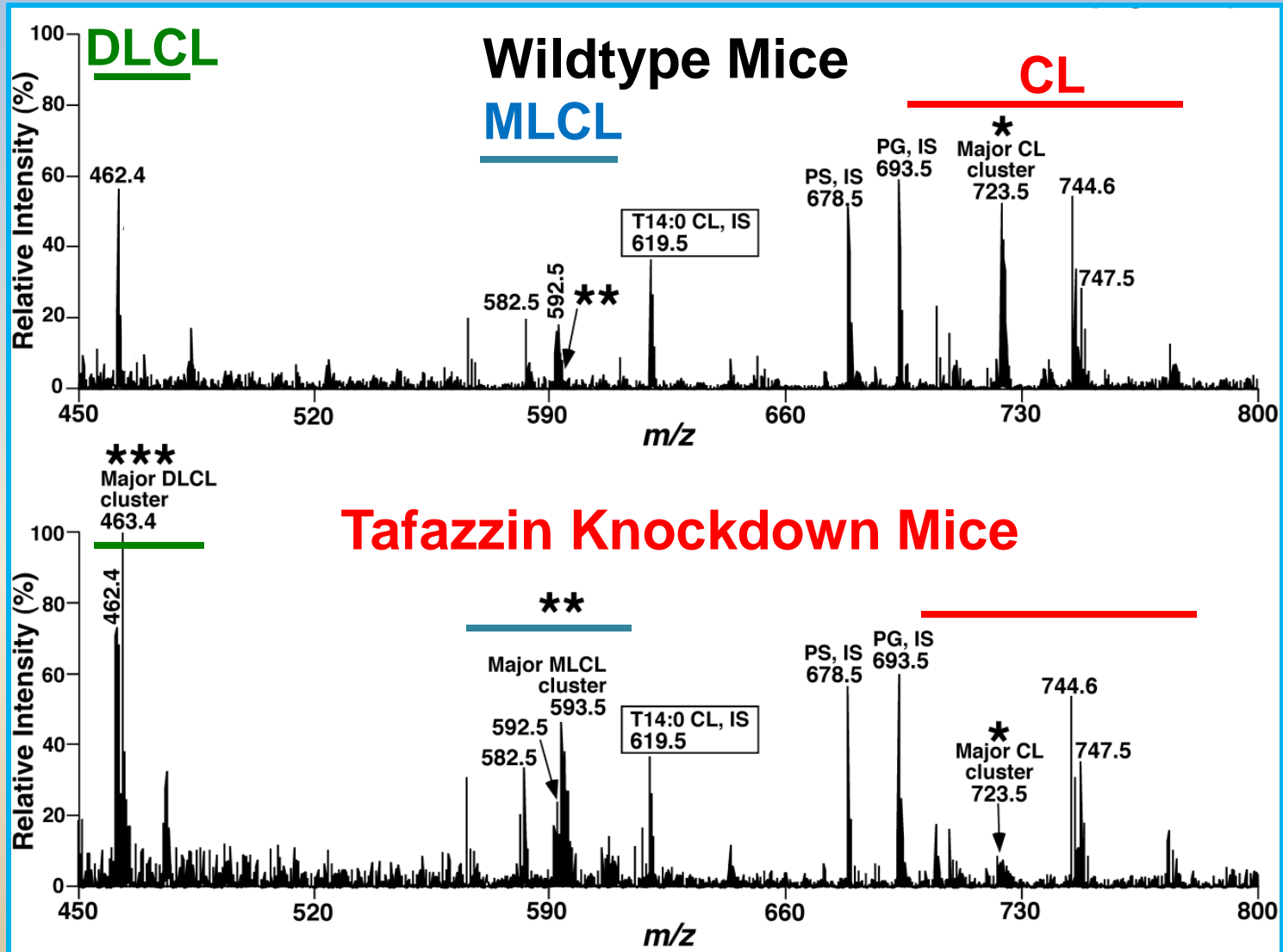
Transcriptomic Analysis



Bioenergetic Analysis

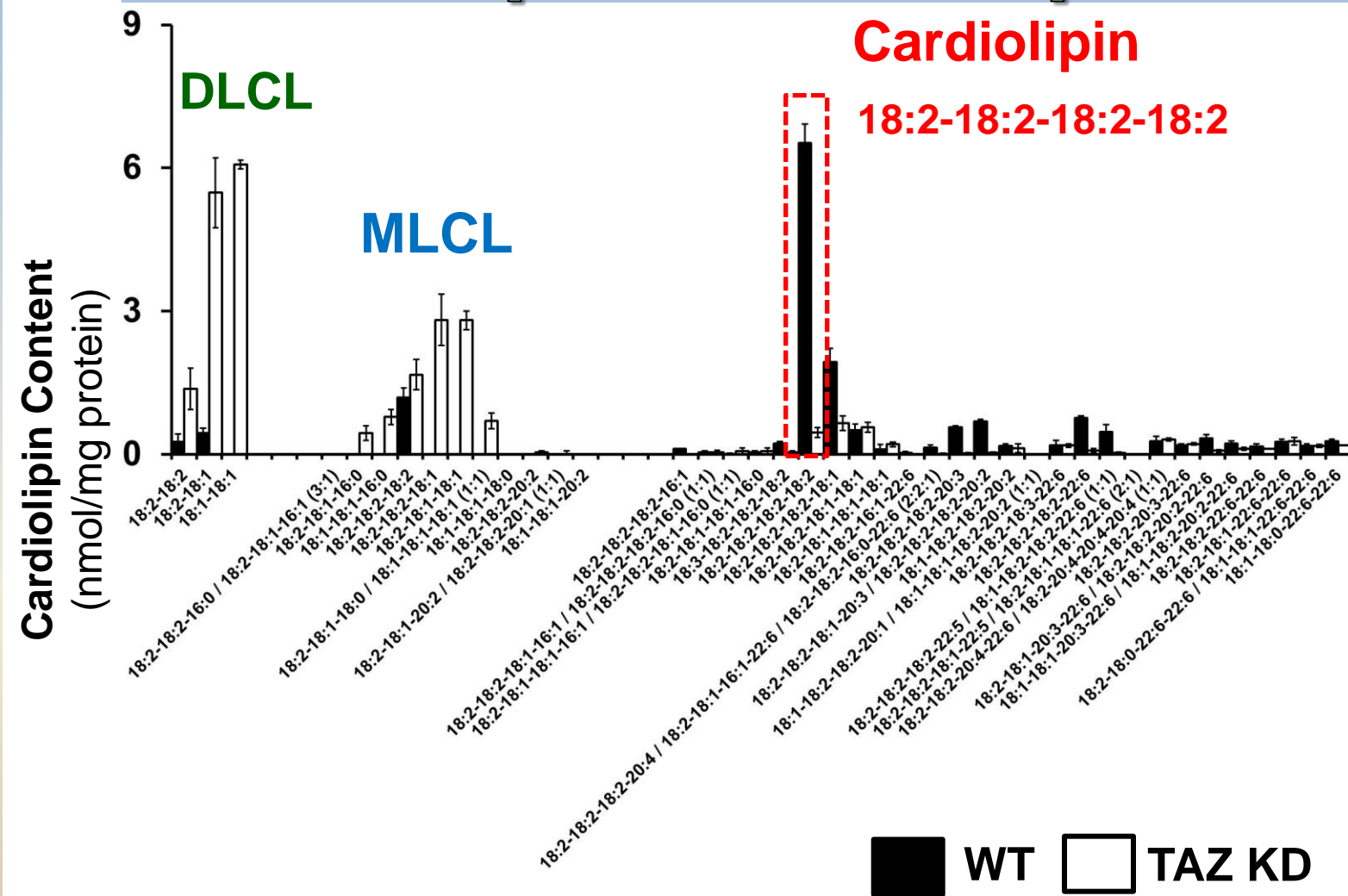
Barth Syndrome Mouse Model

Cardiac Anionic Lipidomic Spectrum



Barth Syndrome Mouse Model

Cardiac Cardiolipin Molecular Species

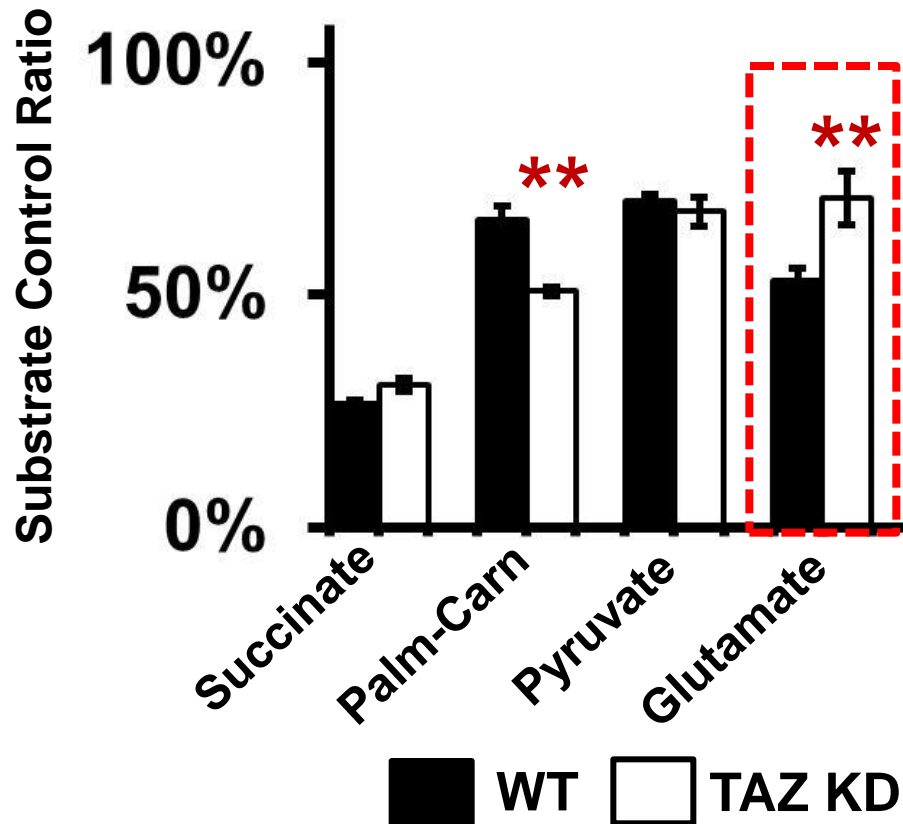


Values represent the mean \pm S.E. cardiolipin molecular species content (nmol/mg protein) in 2 month old WT and Tafazzin knockdown mice (N = 4)

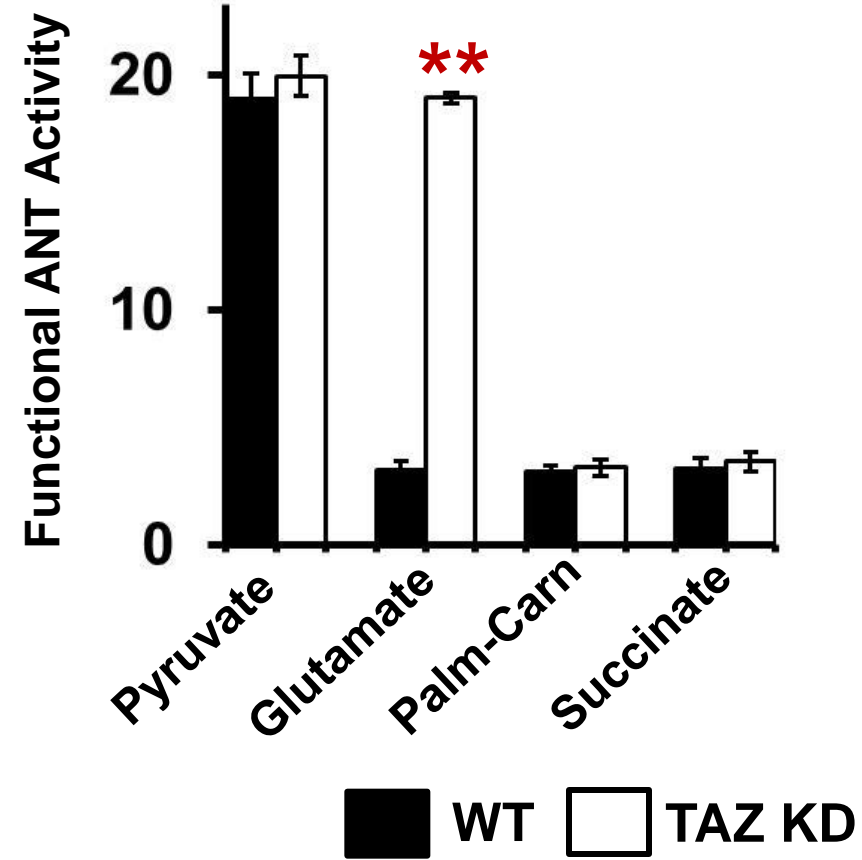
Barth Syndrome Mouse Model

Respiratory and Enzymatic Characterization

State 3 Respiration

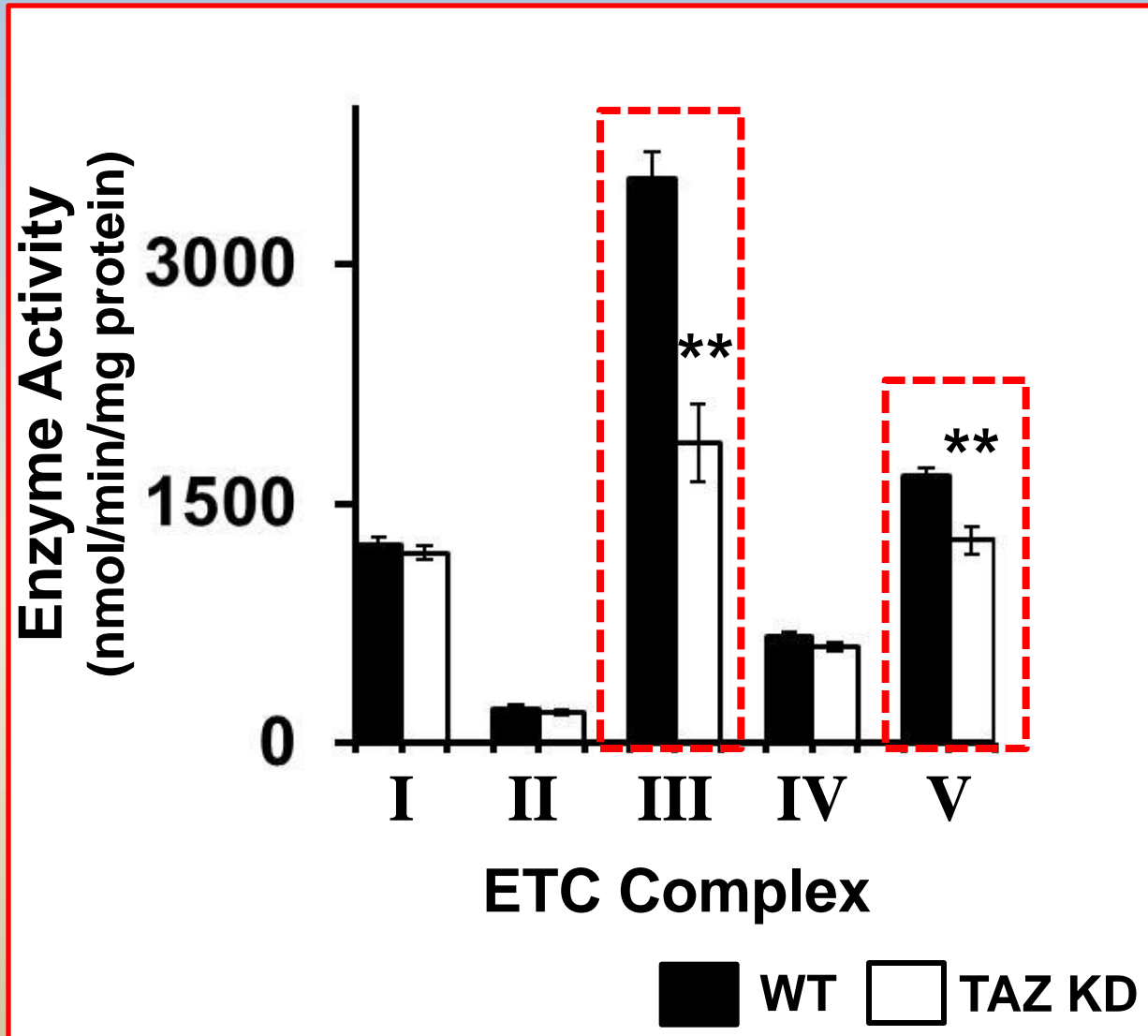


Adenine Nucleotide Translocase Activity



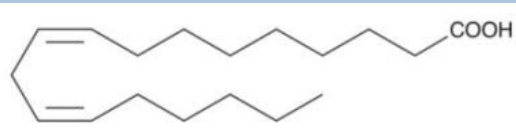
Barth Syndrome Mouse Model

Electron Transport Chain Activities

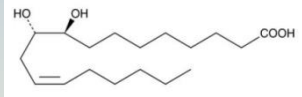


Oxidized Lipidomics

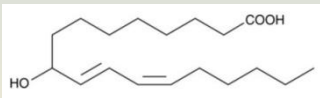
Linoleic



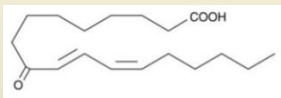
DiHOMEs



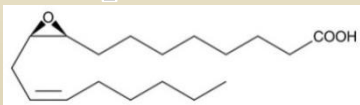
HODEs



oxoODE

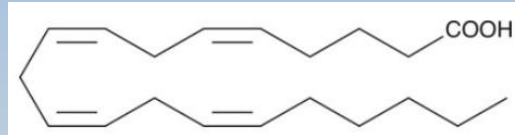


EpOME

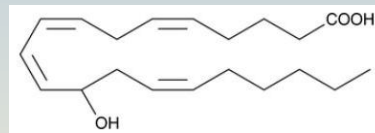


Calcium Influx
Immune Response

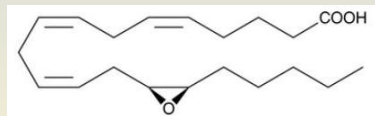
Arachidonic



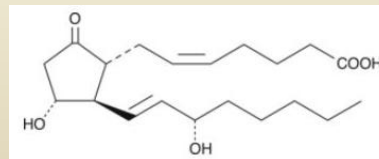
HETE s



EETs

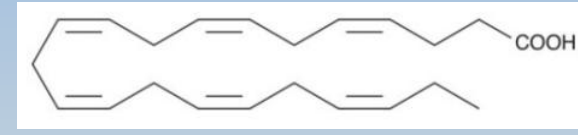


Prostanoids

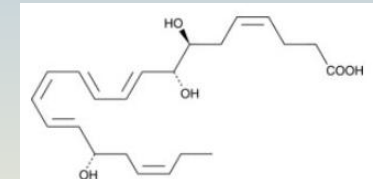


Vasodilation
Vasoconstriction
Inflammation

DHA

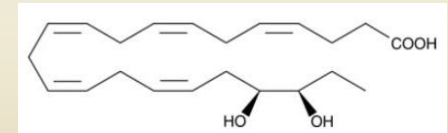


RVD



DiHDoHE

DiHDPA

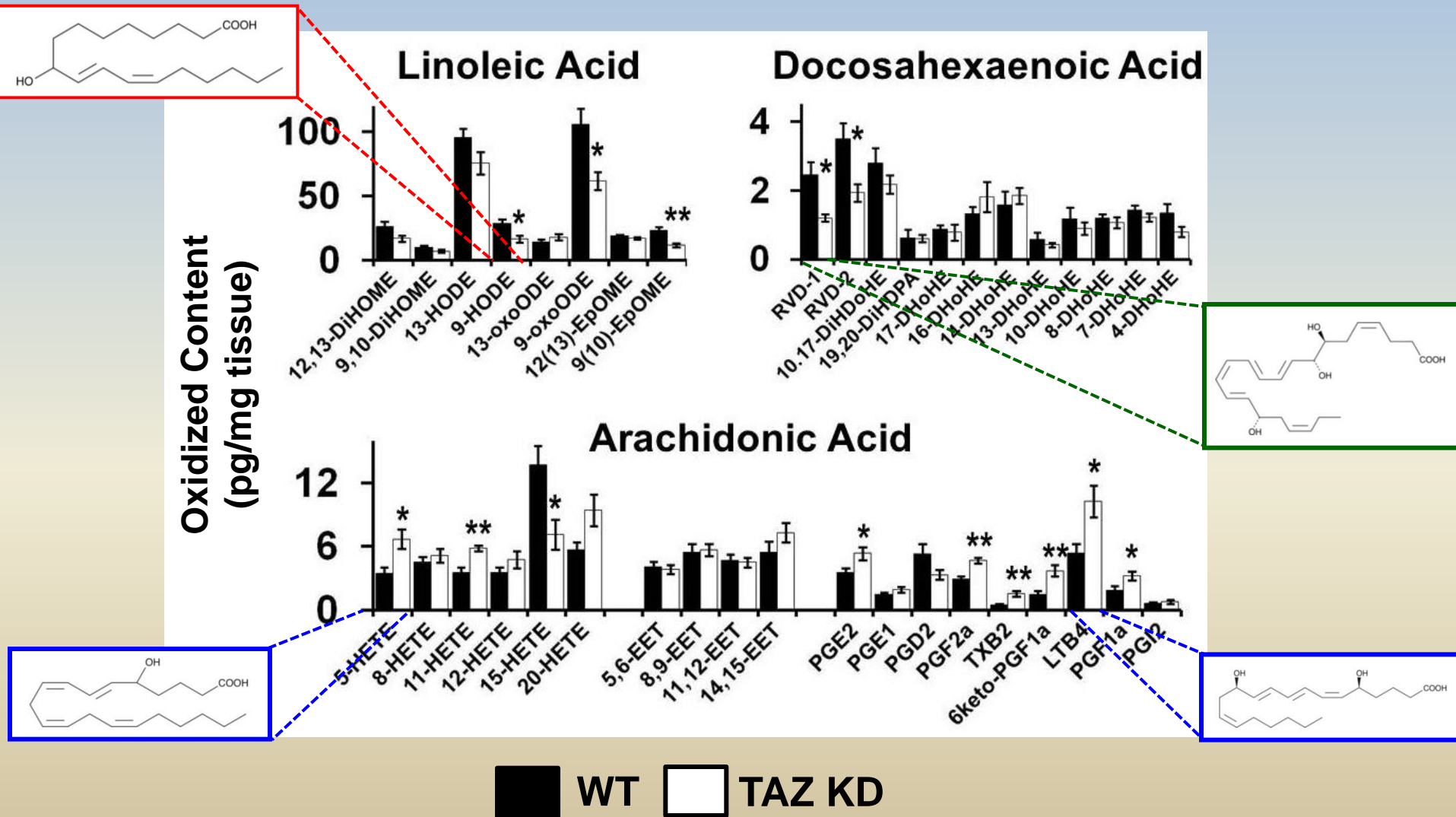


DHoHE

Anti-Inflammatory

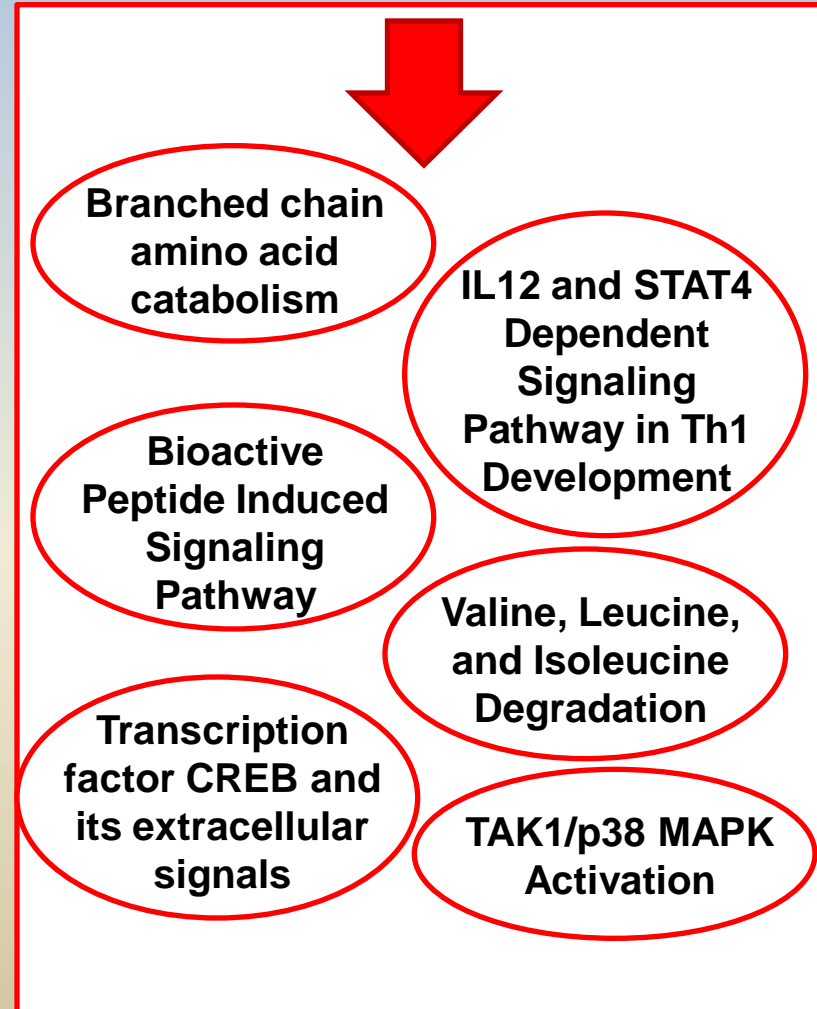
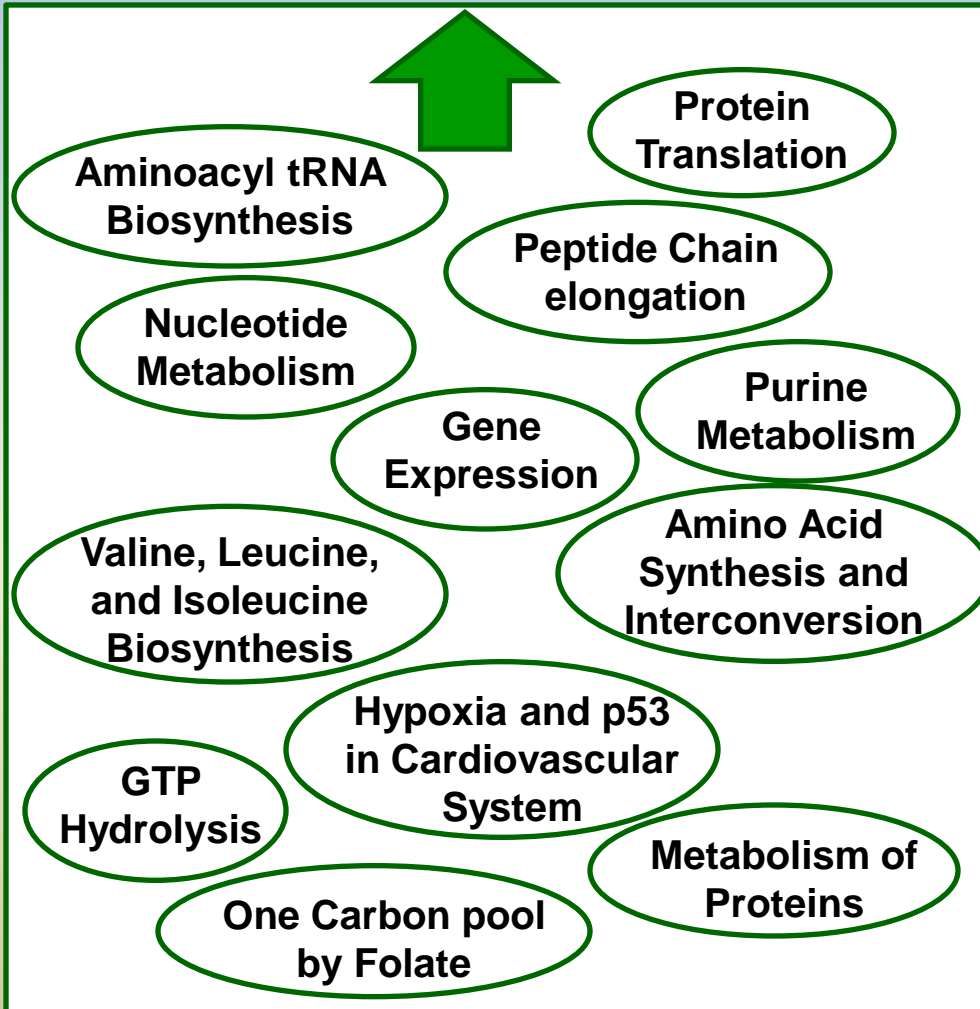
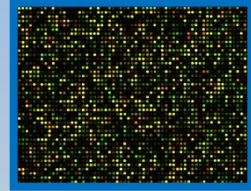
Barth Syndrome Mouse Model

Cardiac Oxidized Lipid Analysis



Barth Syndrome Mouse Model

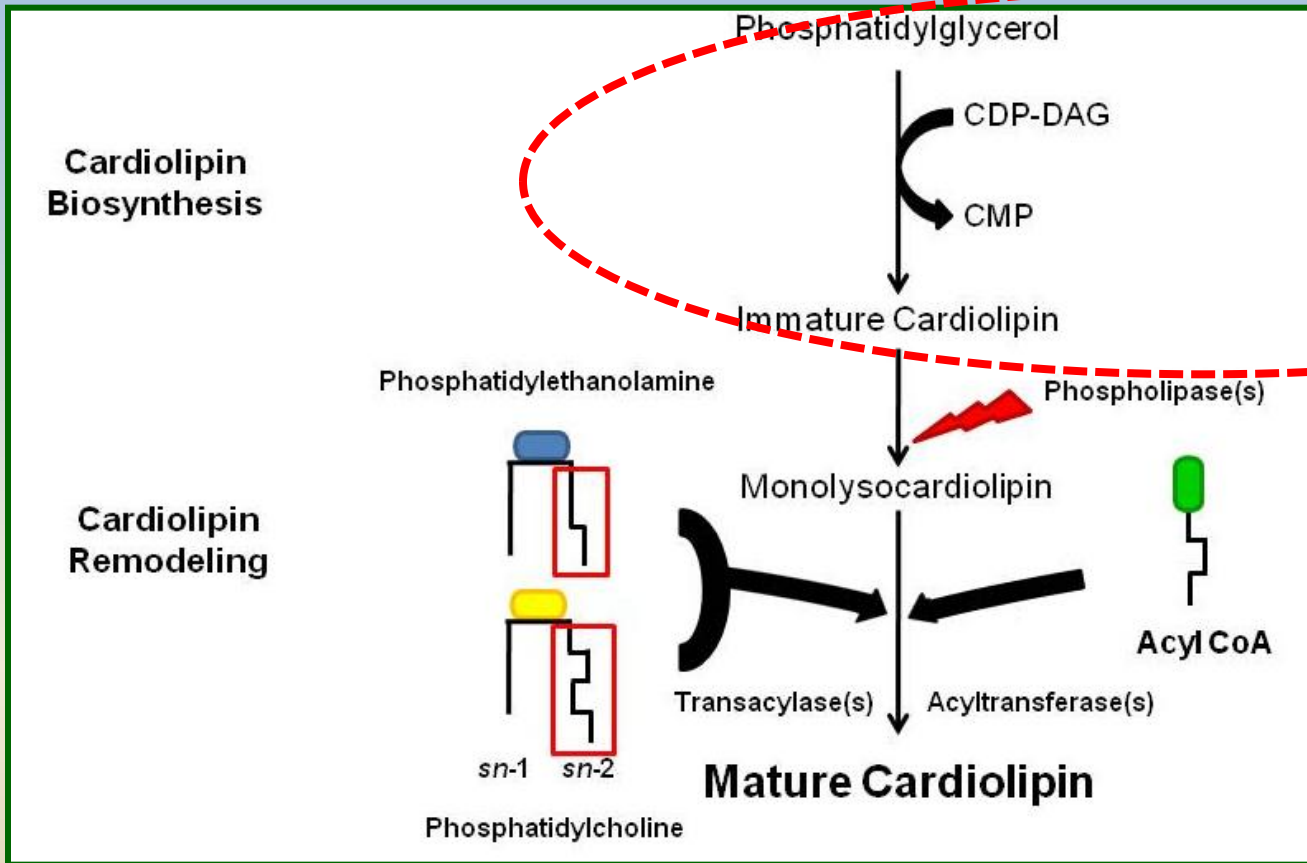
Transcriptomics/GSEA Analysis



Barth Syndrome Mouse Model

- **Demonstrates the lipidomic abnormalities discovered in boys with Barth syndrome**
- **Displays altered substrate utilization (decreased fatty acid and increased glutamate stimulated respiration)**
- **Demonstrates altered mediator lipidomic signature**
- **Compensatory enzyme kinetics (Adenine nucleotide translocase and Electron transport chain) and gene expression (microarray analysis)**

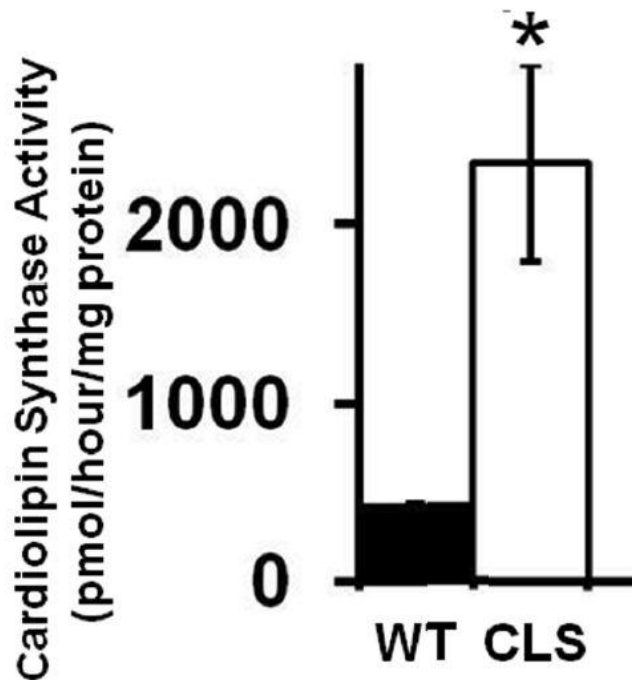
Cardiolipin Synthase (CLS) Mouse Model



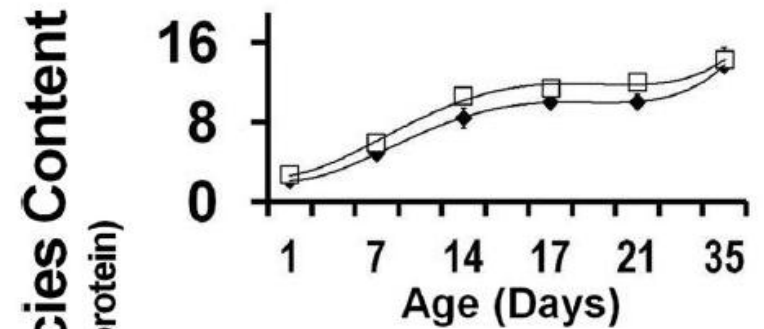
The CLS Model Provides a Molecular Therapeutic Tool to Investigate the Role of Cardiolipin in Attenuating Mitochondrial Dysfunction in Disease

Cardiolipin Synthase Mouse Model

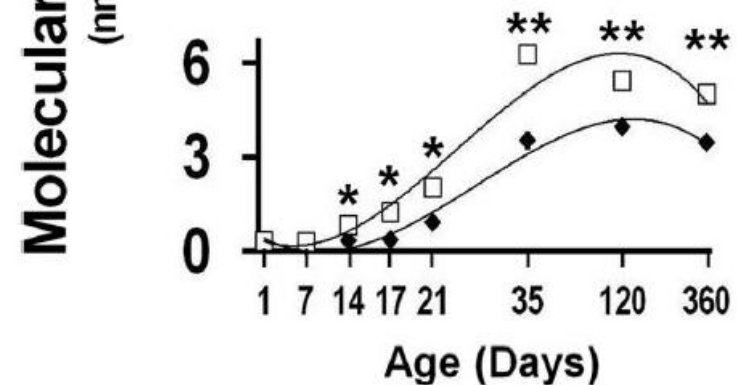
Cardiolipin Synthase Activity



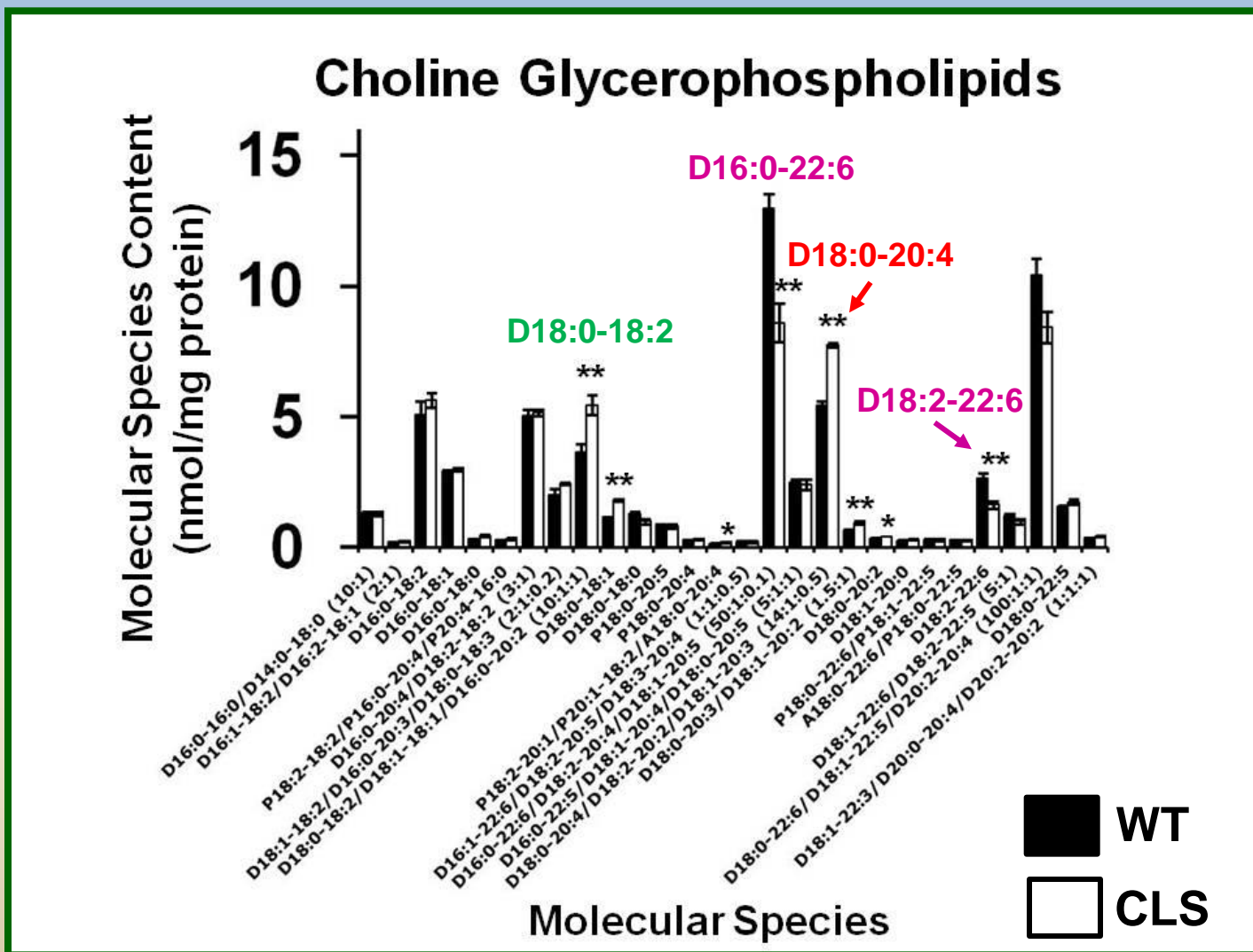
Cardiolipin



Tetra18:2 Cardiolipin

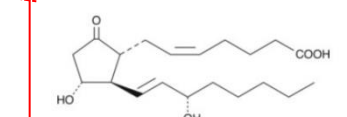
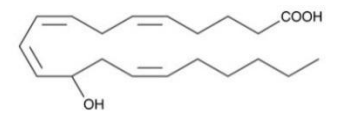
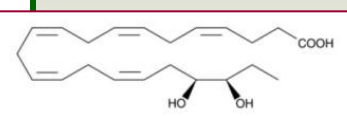
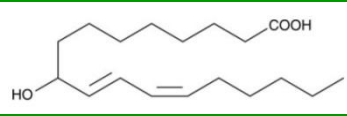
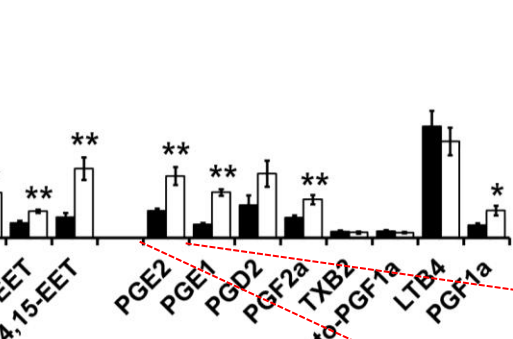
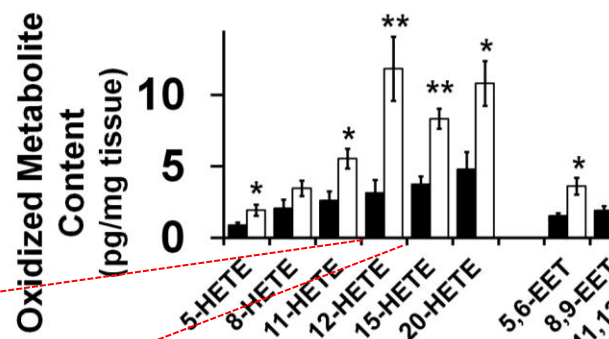
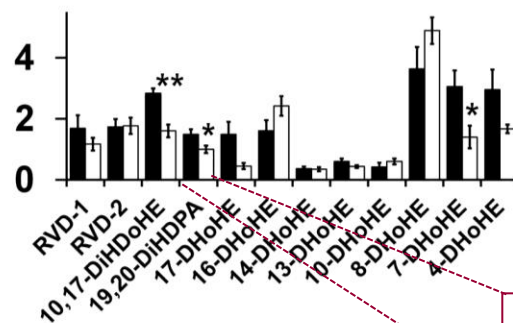
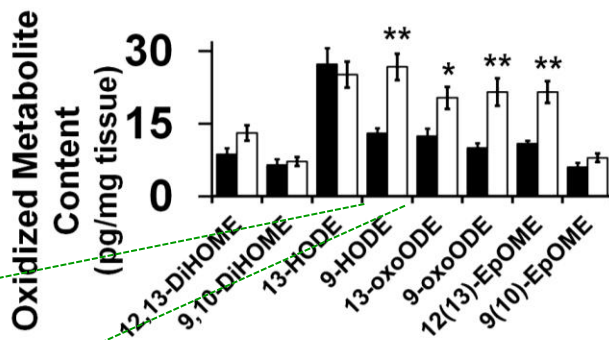
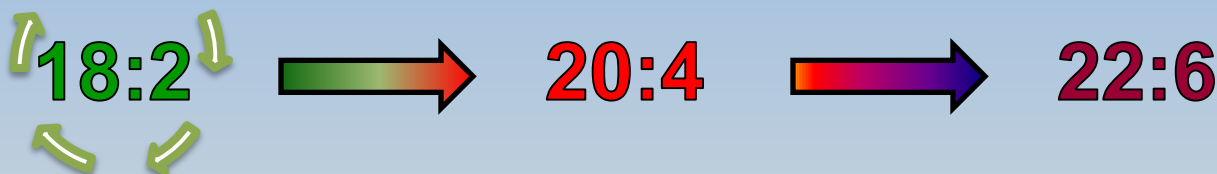


Cardiolipin Synthase Mouse Model



Cardiolipin Synthase Mouse Model

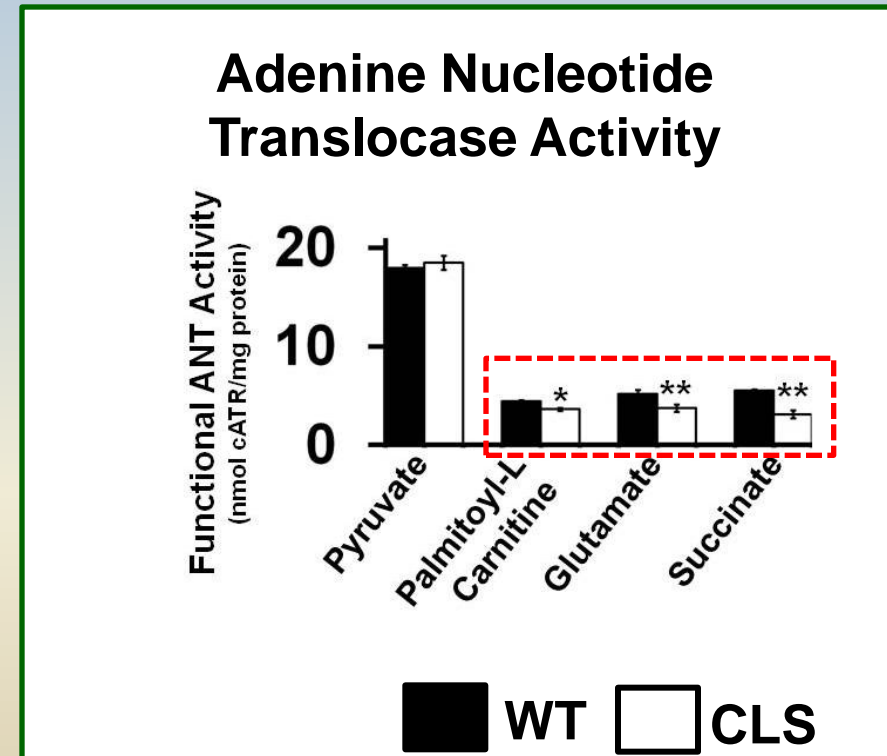
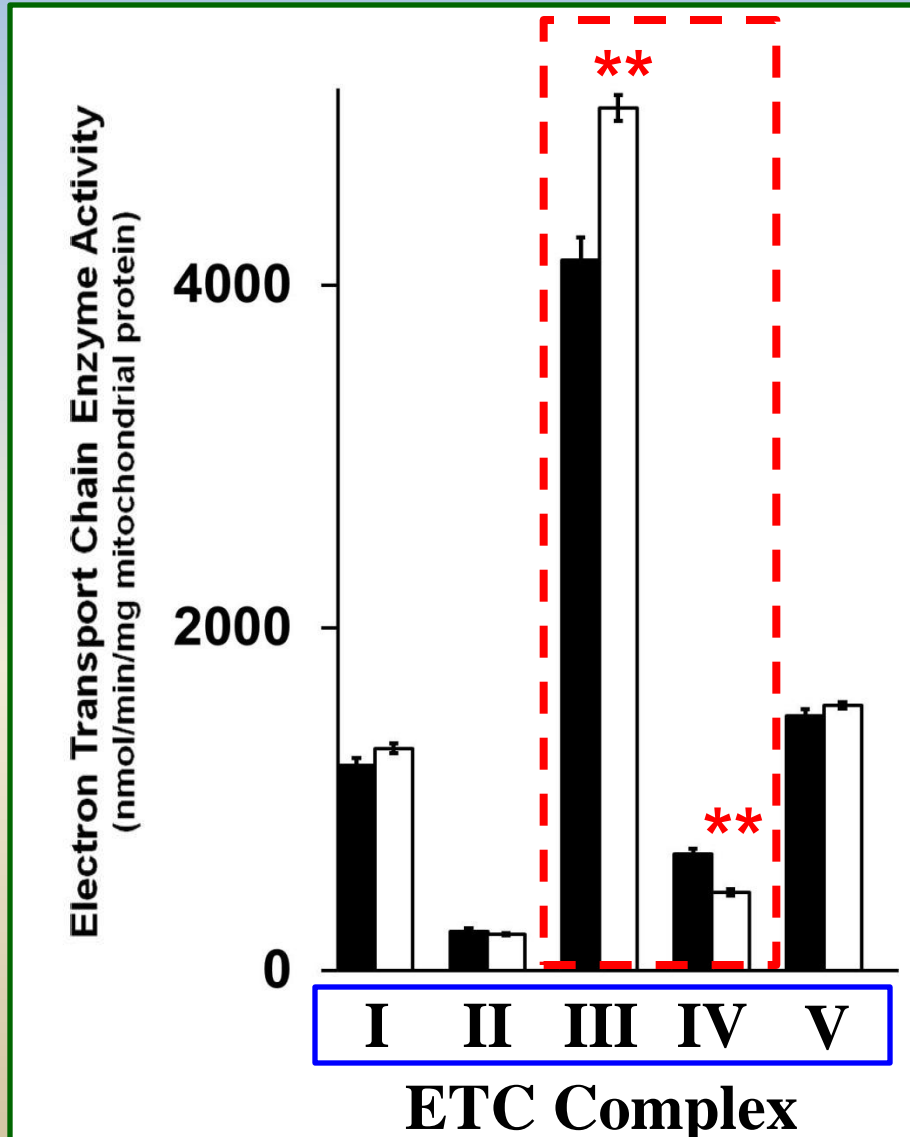
Oxidized Lipid Species



■ WT □ CLS

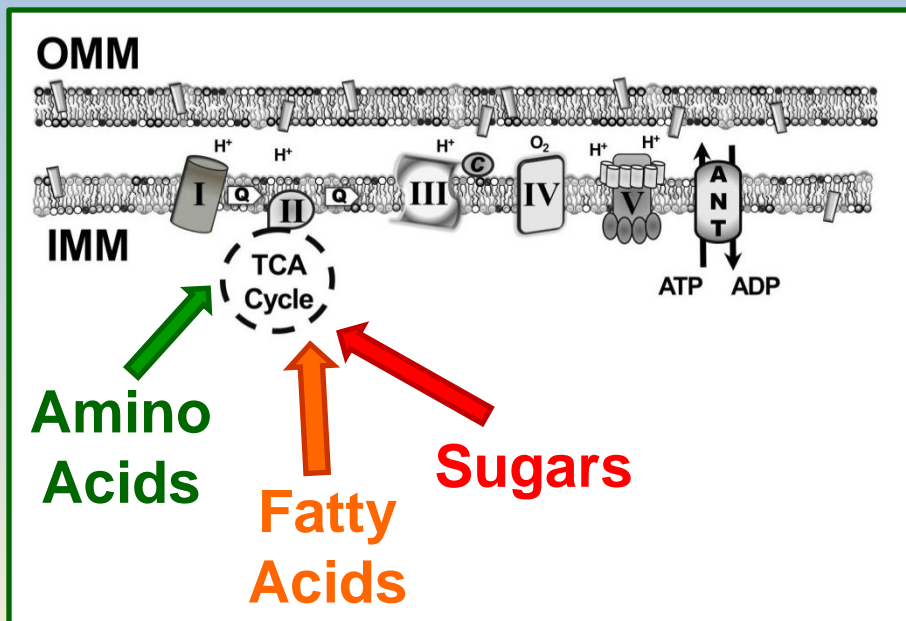
Cardiolipin Synthase Mouse Model

Mitochondrial Enzyme Activities



Cardiolipin Synthase Mouse Model

Mitochondrial Respiration

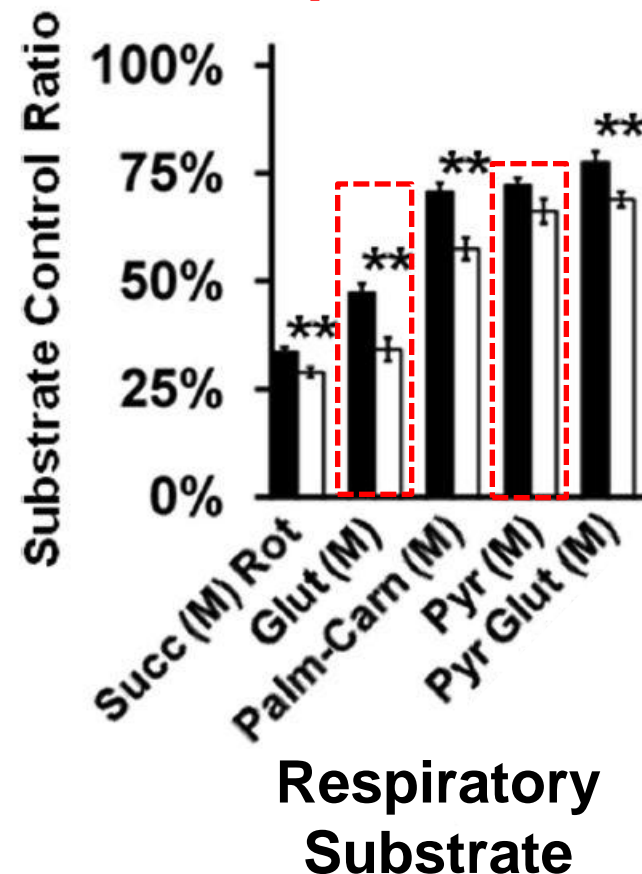


Pyruvate = Normal

Glutamate = ↓

Palm-Carnitine = ↓

State 3 Respiration



Cardiolipin Synthase Mouse Model

Transcriptomics/GSEA Analysis

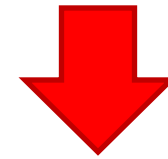


Aminoacyl
tRNA
Biosynthesis

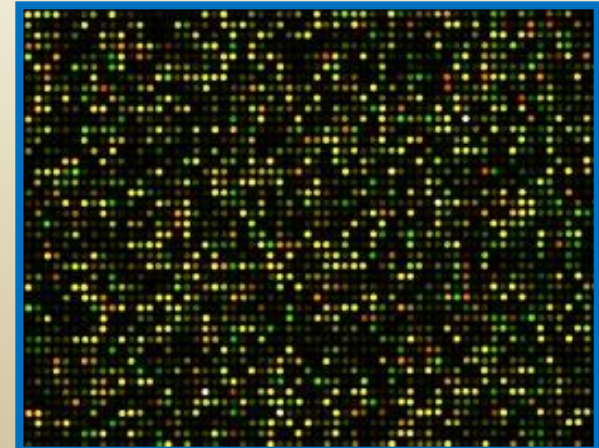
P53
Signaling
Pathway

Glycine,
Serine,
Threonine
Metabolism

Amino Acid
Synthesis and
Interconversion



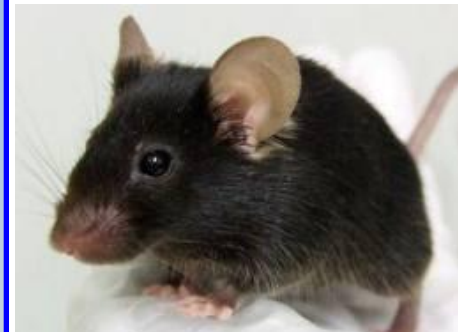
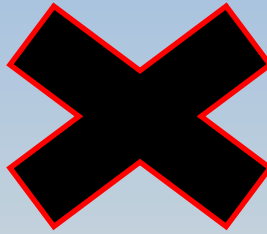
AHSP
Pathway



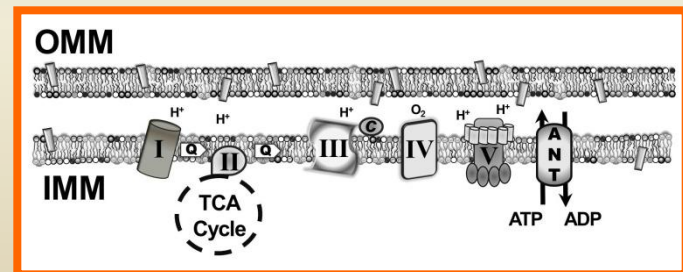
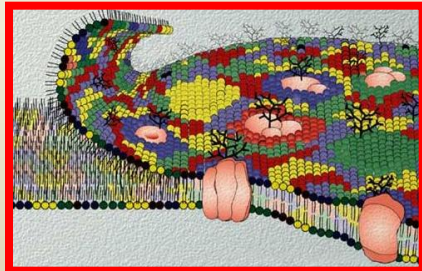
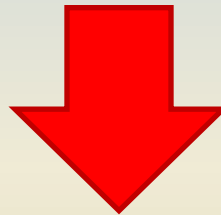
CLS-TG/Tafazzin KD Mouse Model



**Cardiac specific
Tg-Cardiolipin
Synthase Mouse
Model**



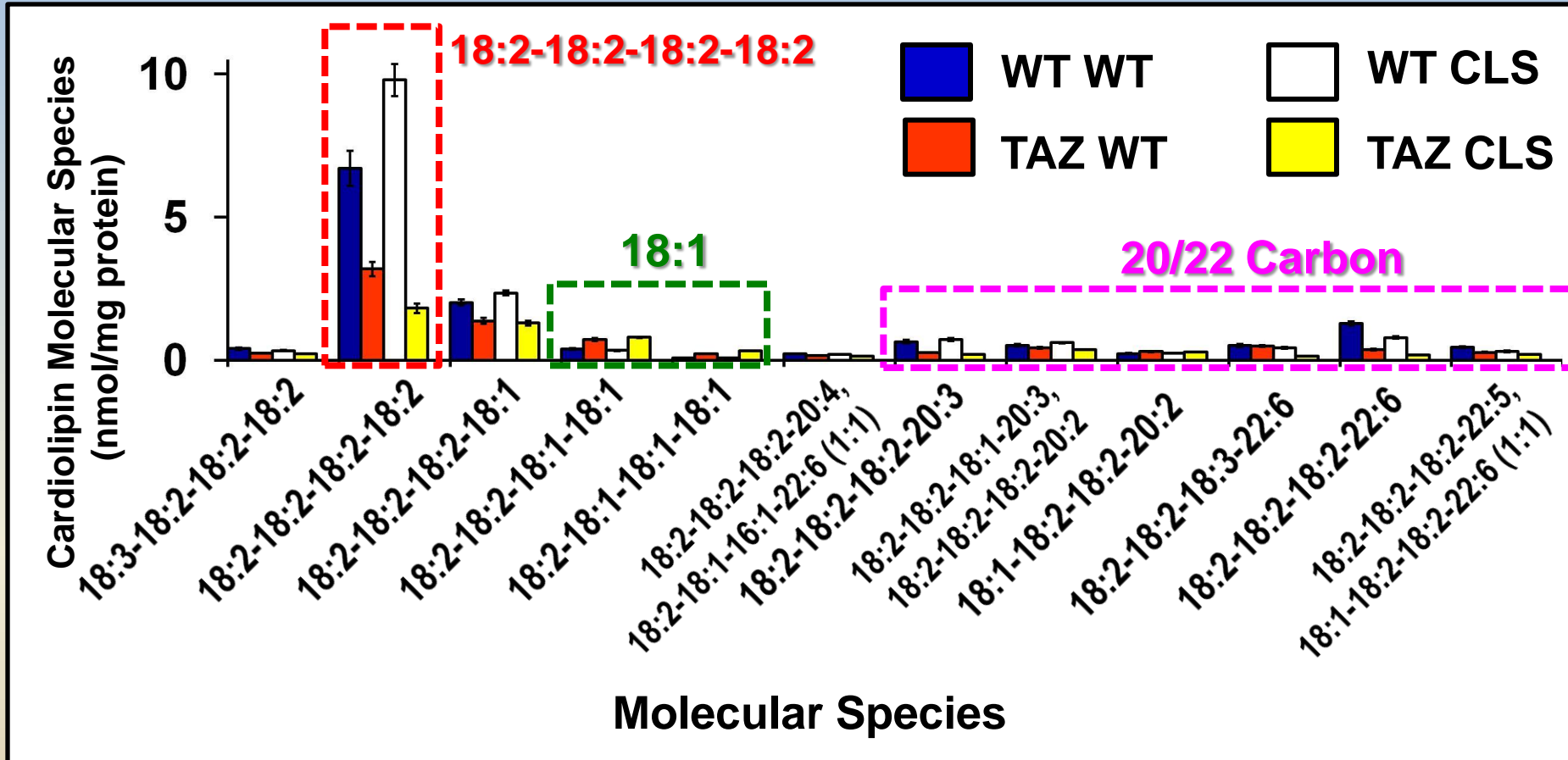
**Tafazzin inducible
shRNA
Knockdown Mouse
Model**



Goal was to use cardiolipin synthase over expression in the heart to attenuate altered cardiolipin molecular species in Barth Syndrome, thus restoring the homeostatic balance of mitochondrial substrate utilization

CLS-TG/Tafazzin KD Mouse Model

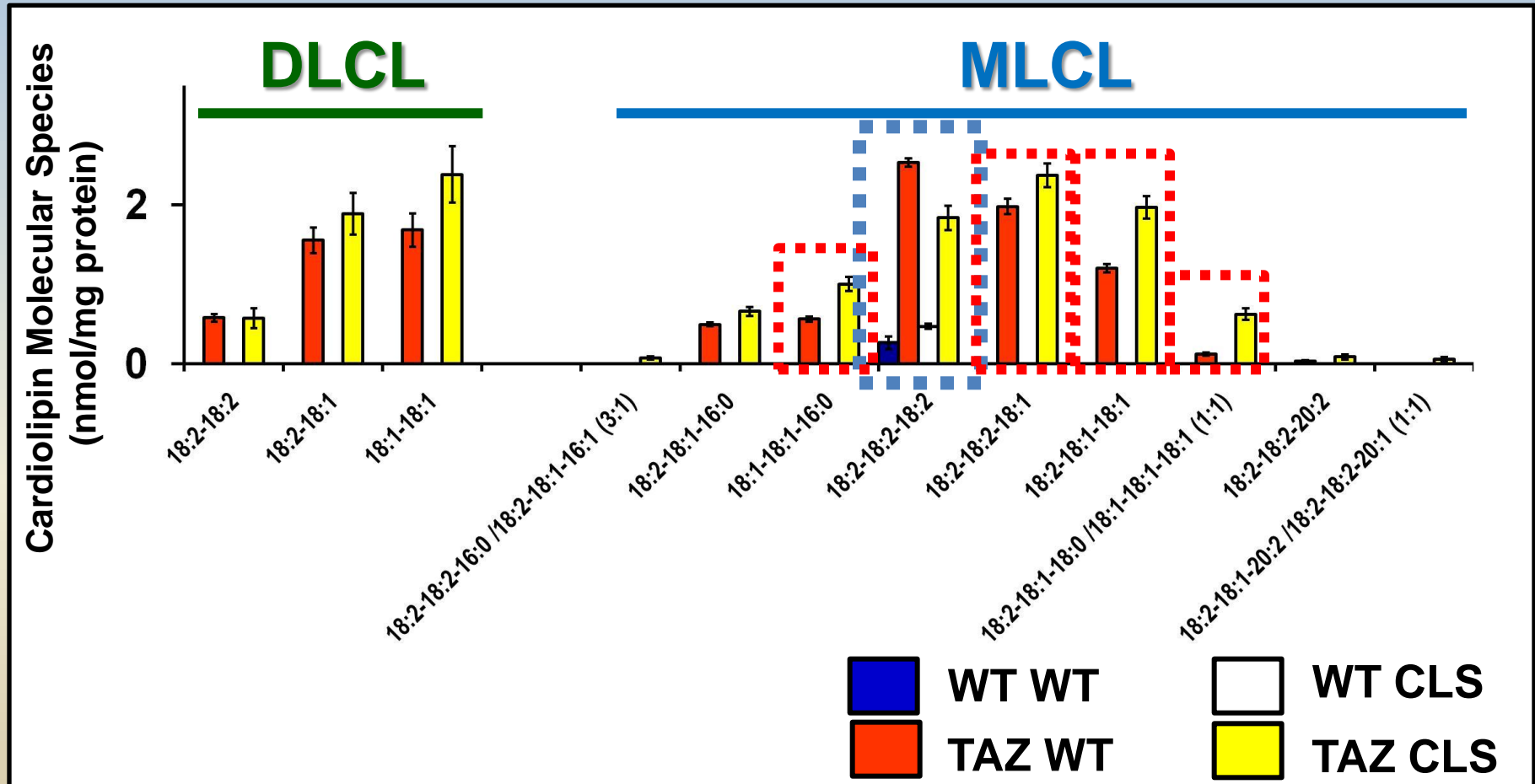
Cardiolipin Molecular Species



Values represent the mean \pm S.E. cardiolipin molecular species content (nmol/mg protein) in WT, Taz, CLS, and Taz x CLS treated with doxycycline for 2 months (N = 4)

CLS-TG/Tafazzin KD Mouse Model

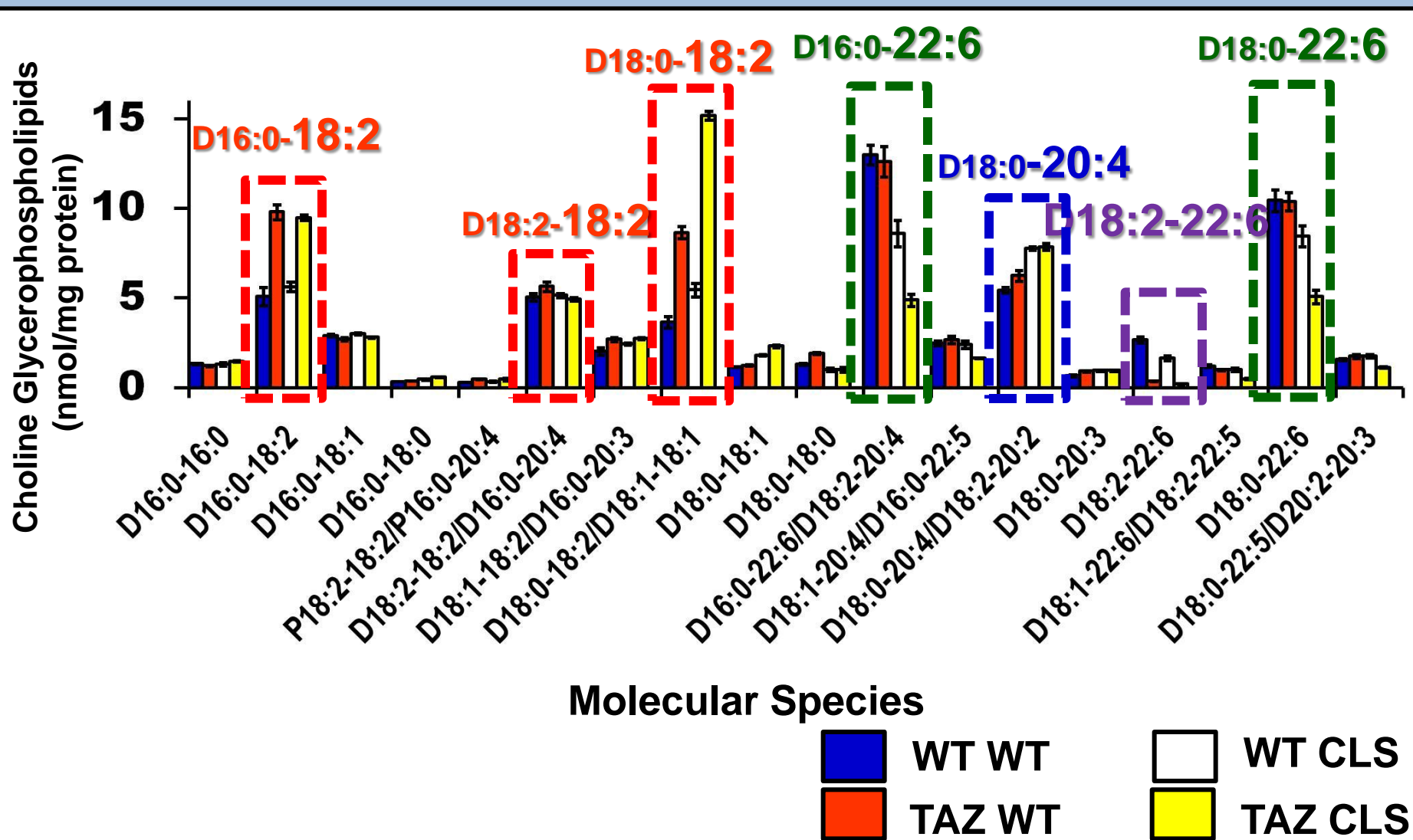
Di/Monolyso - Cardiolipin Molecular Species



Values represent the mean \pm S.E. cardiolipin molecular species content (nmol/mg protein) in WT, Taz, CLS, and Taz x CLS treated with doxycycline for 2 months (N = 4)

CLS-TG/Tafazzin KD Mouse Model

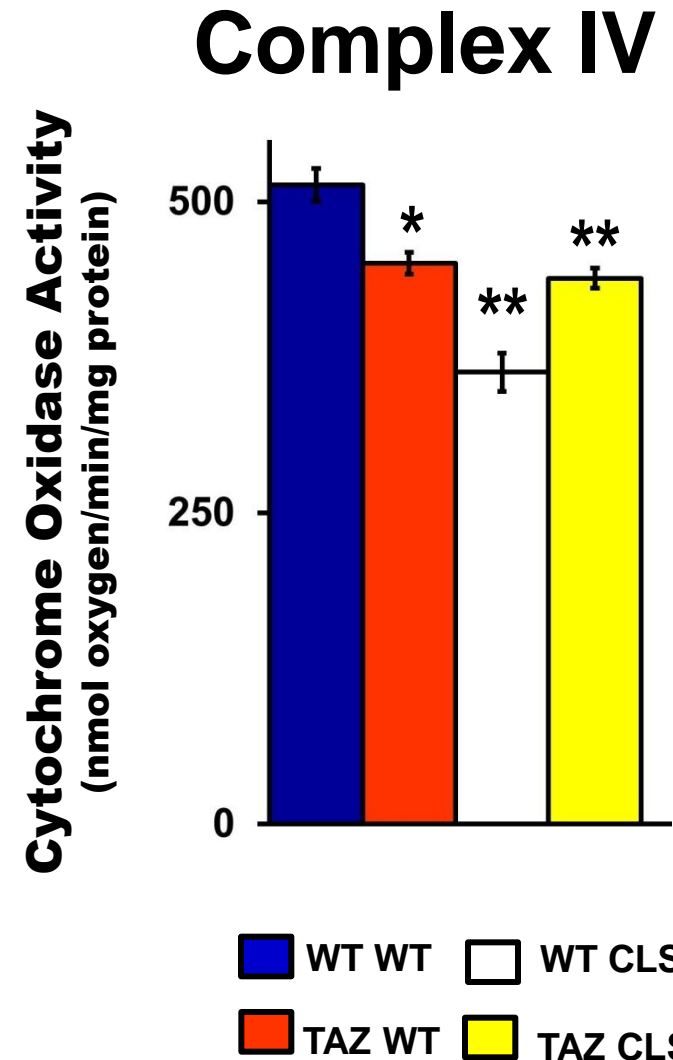
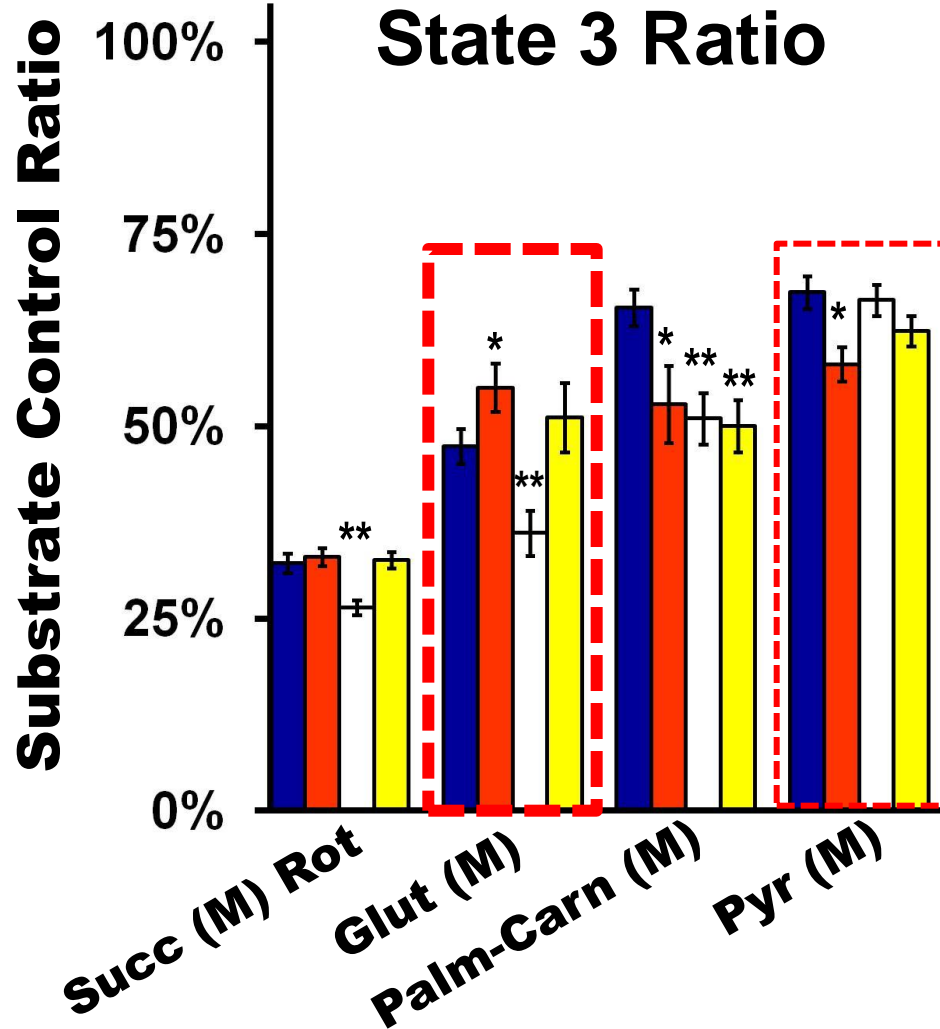
Choline Glycerophospholipid Molecular Species



Values represent the mean \pm S.E. choline glycerophospholipid molecular species content (nmol/mg protein) in WT, Taz, CLS, and Taz x CLS treated on doxycycline for 2 months (N = 4)

CLS-TG/Tafazzin KD Mouse Model

Bioenergetic Analysis



CLS-TG/Tafazzin KD Mouse Model



**Cardiac specific
Tg-Cardiolipin
Synthase Mouse
Model**



**Tafazzin inducible
shRNA
Knockdown Mouse
Model**

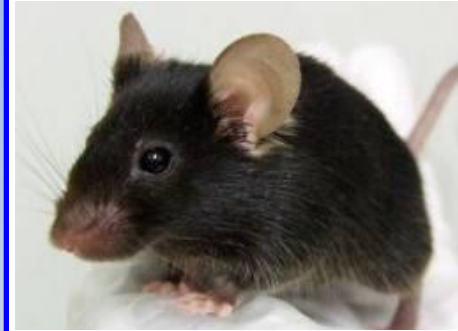
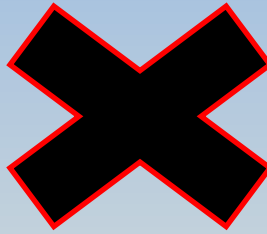
- **Although cardiolipin molecular species were less remodeled in the Taz X CLS mouse model, the overall bioenergetic phenotype was attenuated, possibly suggesting that alternative factors may be influencing bioenergetics**

iPLA₂ γ -TG and KO Mouse Model

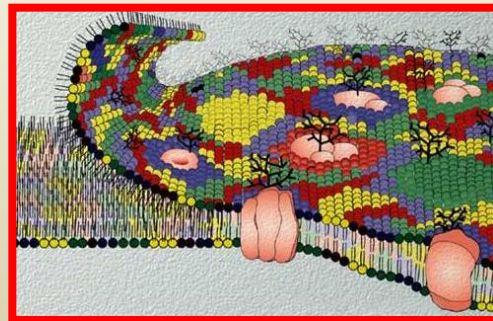
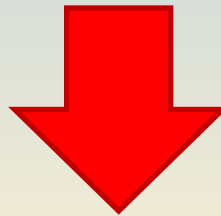
iPLA₂ γ KO



iPLA₂ γ -TG



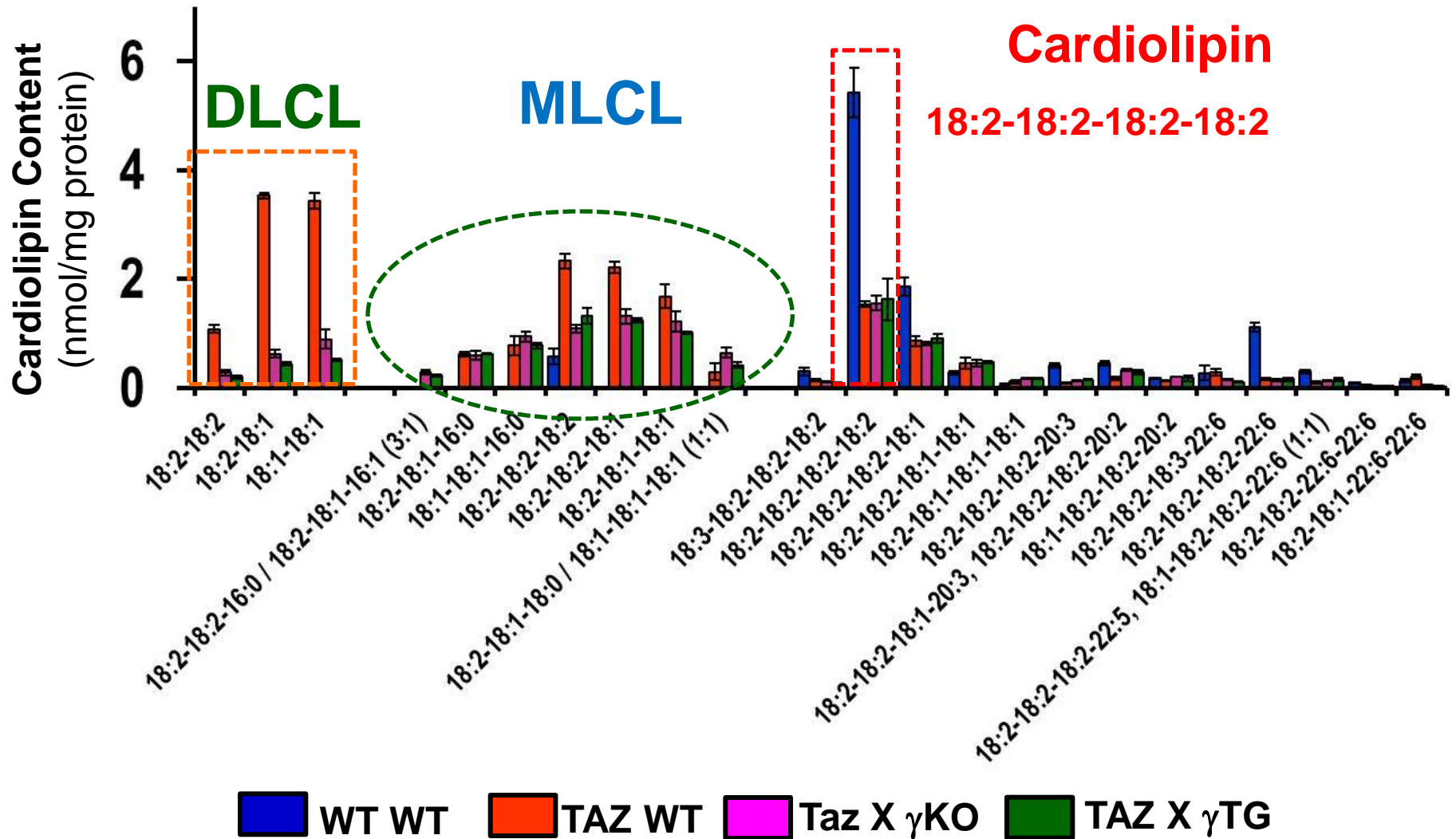
**Tafazzin inducible
shRNA
Knockdown Mouse
Model**



Does iPLA₂ γ expression attenuate dysfunctional cardiolipin remodeling as well as effect survival of the cross?

(iPLA γ -TG or KO) X Tafazzin KD Mouse Model

Cardiolipin Molecular Species



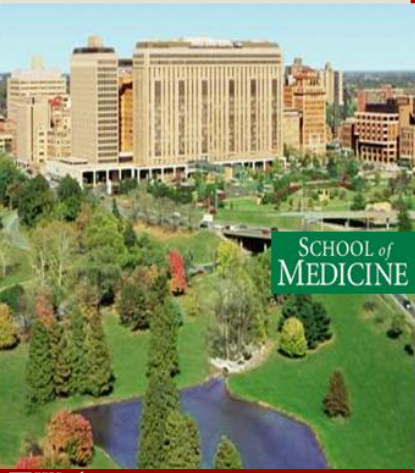
Values represent the mean \pm S.E. cardiolipin molecular species content (nmol/mg protein) in WT, Taz, Taz x iPLA γ KO, and Taz x iPLA γ -TG treated on doxycycline for 2 months (N = 4)

Conclusion

- Through the utilization of various transgenic models as well as investigating dynamic flux of the lipidome, we can elucidate the lipidomic/bioenergetic connection that initiates pathological sequelae or that can be identified for therapeutic efficacy
- Regulation of the mitochondrial lipidome may hold unknown mechanism that go well beyond just structure and enzyme kinetics
- The inducible Tafazzin shRNA knockdown mouse model of Barth Syndrome is an invaluable tools to discover unknown mechanism that embody the Barth Syndrome phenotype which can be therapeutically targeted



Barth Syndrome
Foundation



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CENTER FOR
Cardiovascular Research

Acknowledgements

Barth Syndrome Foundation

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University School of Medicine**

Washington University School of Medicine

Dr. Todd Cade

Division of Bioorganic Chemistry and Molecular Pharmacology

Dr. Xianlin Han

Dr. Richard W. Gross

Dr. Kui Yang

Dr. Ari Cedars

Dr. David Mancuso

Dr. Chris Jenkins

Dr. Xinping Luo

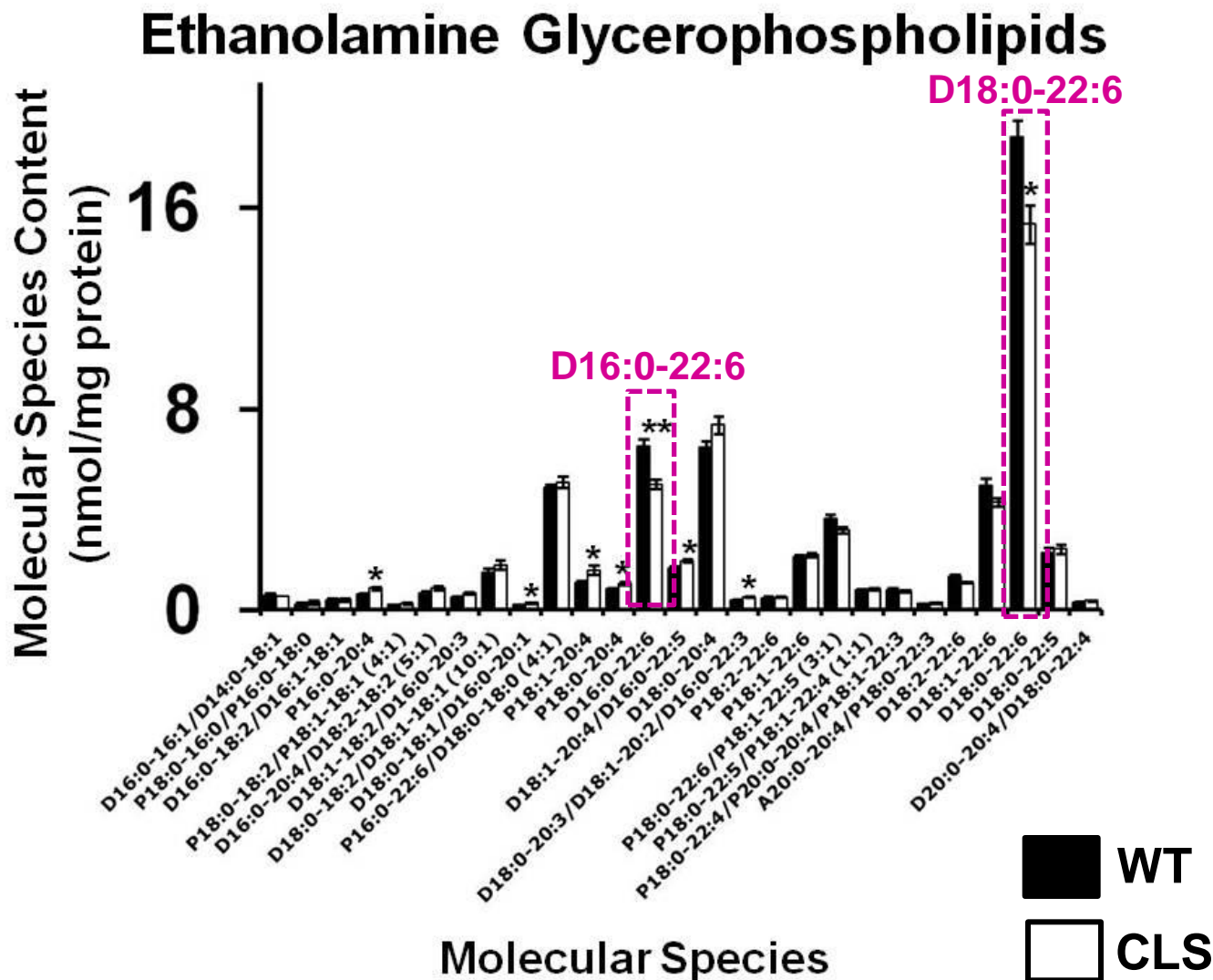
Mr. Harold Sims

Mrs. Zhongdan Zhao

Mrs. Hua Cheng

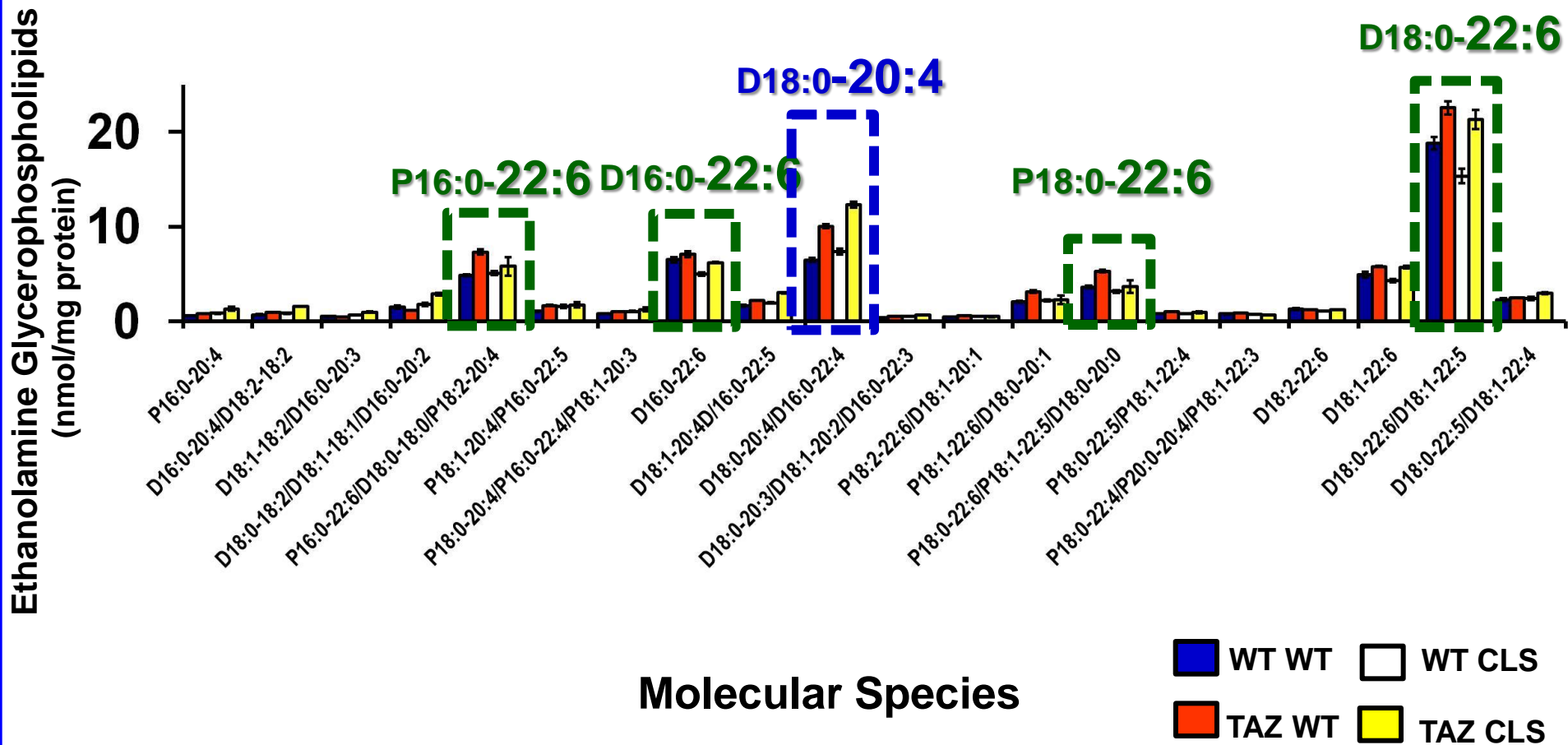
Mrs. Shaoping Guan

Cardiolipin Synthase Mouse Model



CLS-TG/Tafazzin KD Mouse Model

Ethanolamine Glycerophospholipid Molecular Species



Values represent the mean \pm S.E. ethanolamine glycerophospholipid molecular species content (nmol/mg protein) in WT, Taz, CLS, and Taz x CLS treated on doxycycline for 2 months (N = 4)

Cardiolipin Synthase Mouse Model

Cardiolipin Molecular Species

