

3rd Annual Workshop on Metabolomics

# Integration of Biology and the Metabolome: Breaking Barriers Across Disciplines

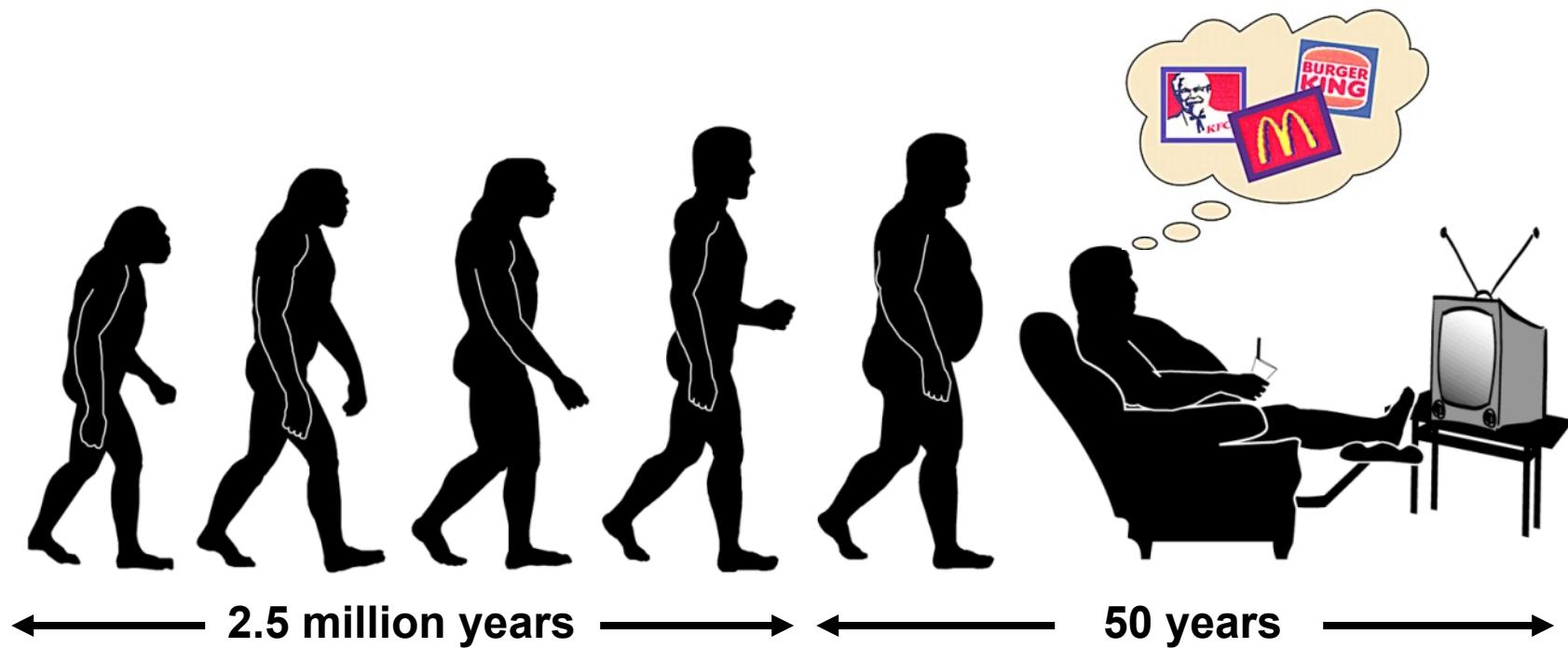
Monday, June 15, 2015

Adam R. Wende, Ph.D.

Assistant Professor  
Molecular and Cellular Pathology  
Department of Pathology

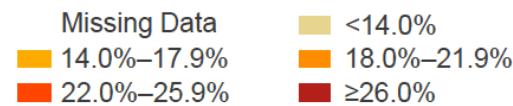
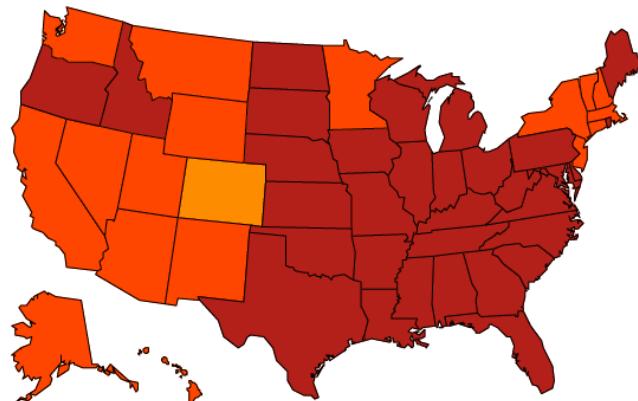


# Defining the Problem

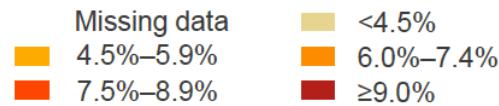
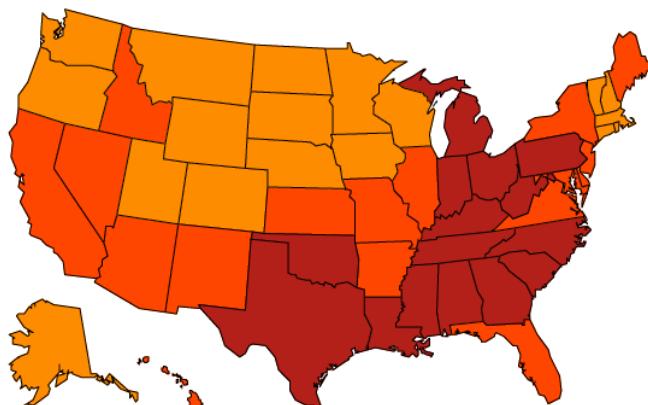


From: Roger Unger - UTSW

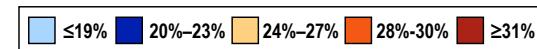
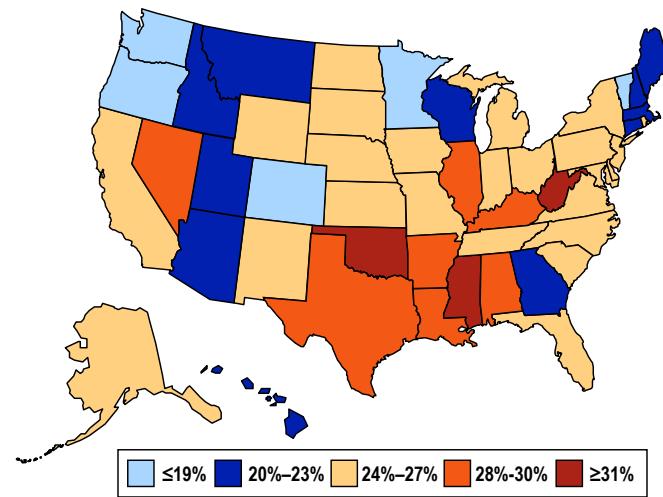
## 2010 – Obesity



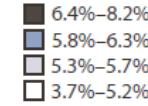
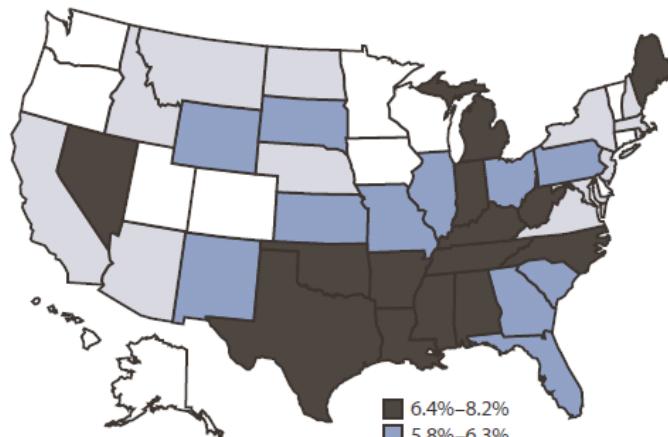
## 2010 – Diabetes



## 2010 – Physical Inactivity



## 2010 – Heart Disease



[www.cdc.gov/diabetes/statistics](http://www.cdc.gov/diabetes/statistics) and [www.cdc.gov/mmwr](http://www.cdc.gov/mmwr)

# Cardiac Metabolic Substrate Utilization

**Table 2. Brief Overview of Myocardial Metabolism in Physiological and Pathophysiological Conditions**

	MVo <sub>2</sub>	Glucose Metabolism	Fatty Acid Metabolism
Aging	↑	↑	↓
Female sex	↑	↓	↑
Obesity	↑	—	↑
Diabetes, types 1 and 2	—↑	↓	↑
Hypertension: LV hypertrophy	—	↑	↓
Dilated cardiomyopathy	—	↑	↓
Ischemia	↓	↑	↓

Peterson and Gropler 2010 *Circ Cardiovasc Imaging* 3:211

# Cardiac Metabolic Substrate Utilization

## Studies on Myocardial Metabolism\*

### *IV. Myocardial Metabolism in Diabetes*

I. UNGAR, M.D., M. GILBERT, M.D., A. SIEGEL, M.S., J. M. BLAIN, M.D. and R. J. BING, M.D.

\* From the Department of Medicine and Physiology, University of Alabama Medical Center, Birmingham, Ala.  
Work supported by the U. S. Public Health Service Grant No. H-1129(CS), The Life Insurance Medical Research Fund and the American Heart Association.

UAB founded in 1969

amino acids by the heart in both species.

Myocardial glucose consumption is reduced in dog and man relative to the elevation in blood glucose concentration. The myocardial

usage of ketones is slightly increased in diabetic hearts of patients and significantly elevated in the dog. The main difference concerns the utilization of fatty acids; this is significantly increased in the human heart but is unchanged in the dog. Whether this is due to a species difference or to differences in type and severity

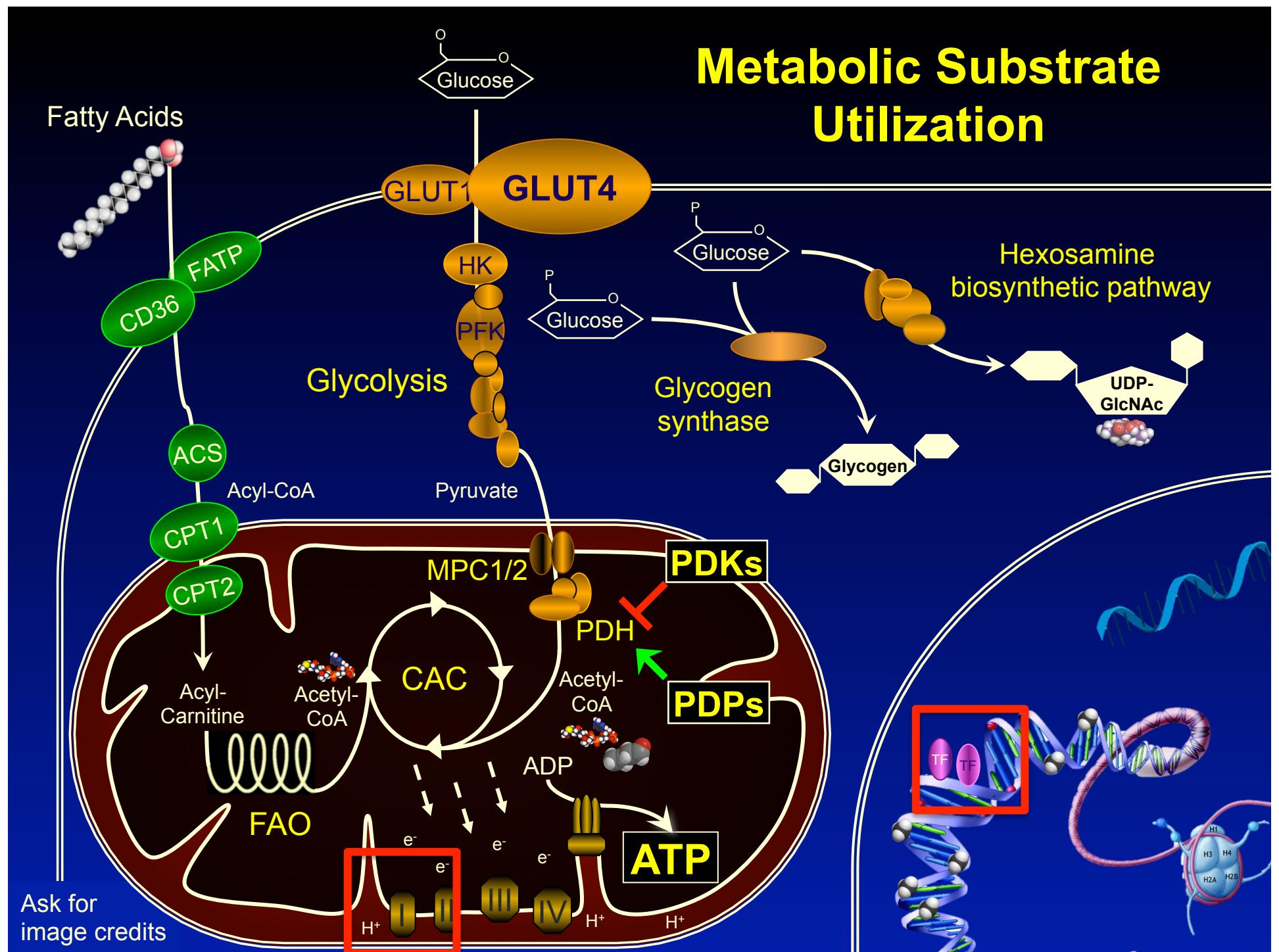
Ungar ... Bing 1955 Am J Med 18(3):385



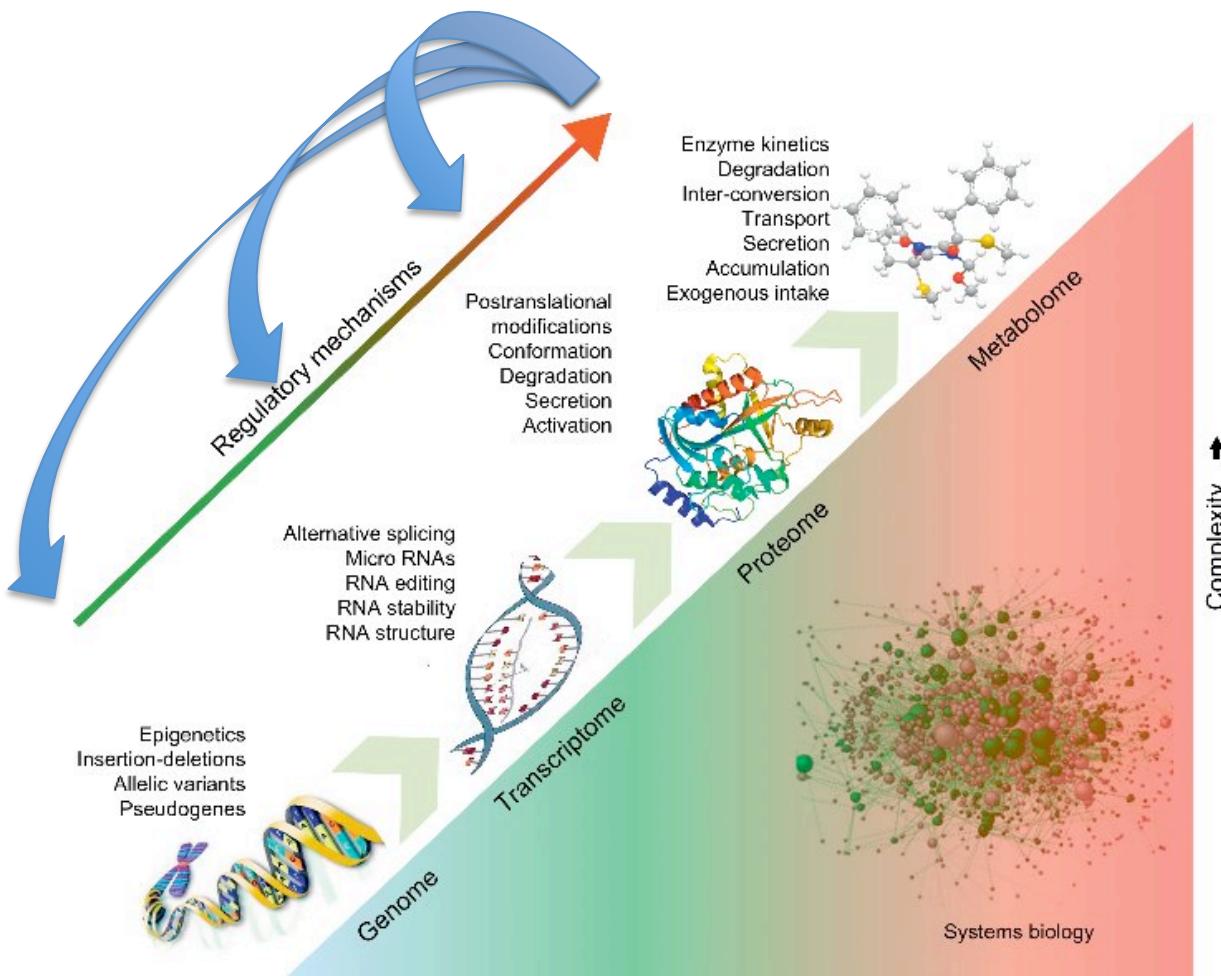
Metabolomics

**UAB** THE UNIVERSITY OF  
ALABAMA AT BIRMINGHAM  
Knowledge that will change your world

# Metabolic Substrate Utilization



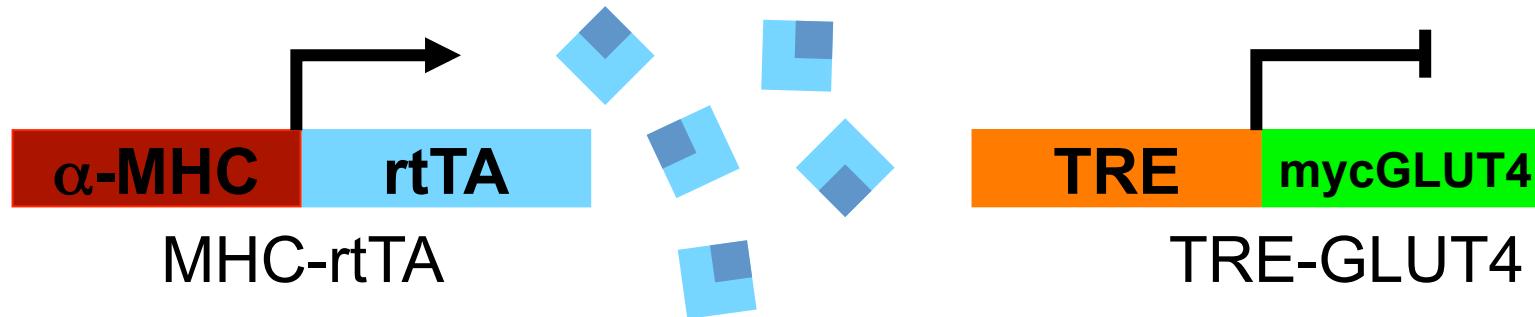
# Defining the Mechanism



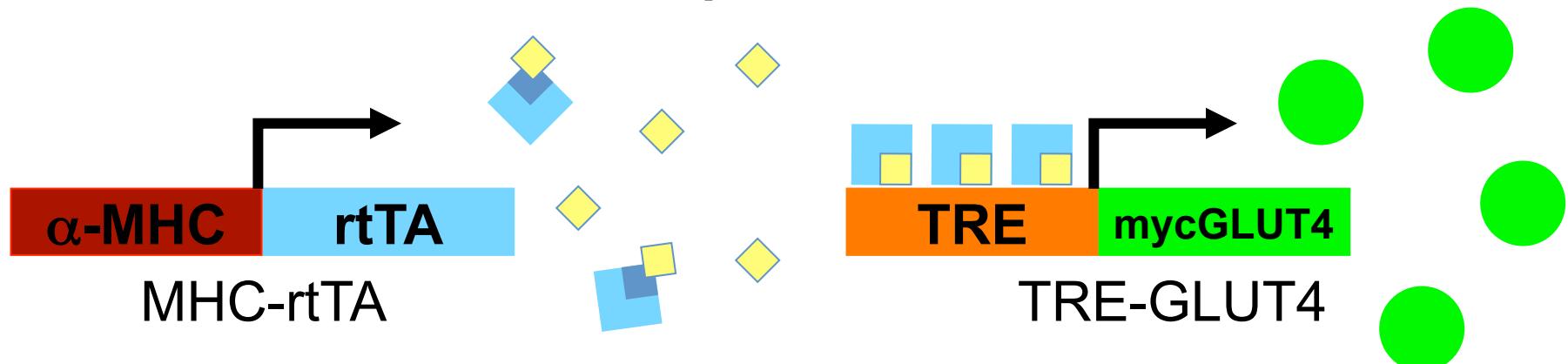
Barallobre-Barreiro ... Mayr 2013 Rev Esp Cardiol 66:657

# Model Development

**DOX absent = OFF**

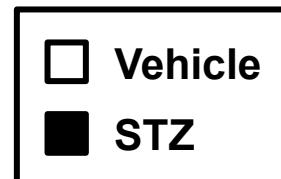


**DOX present = ON**

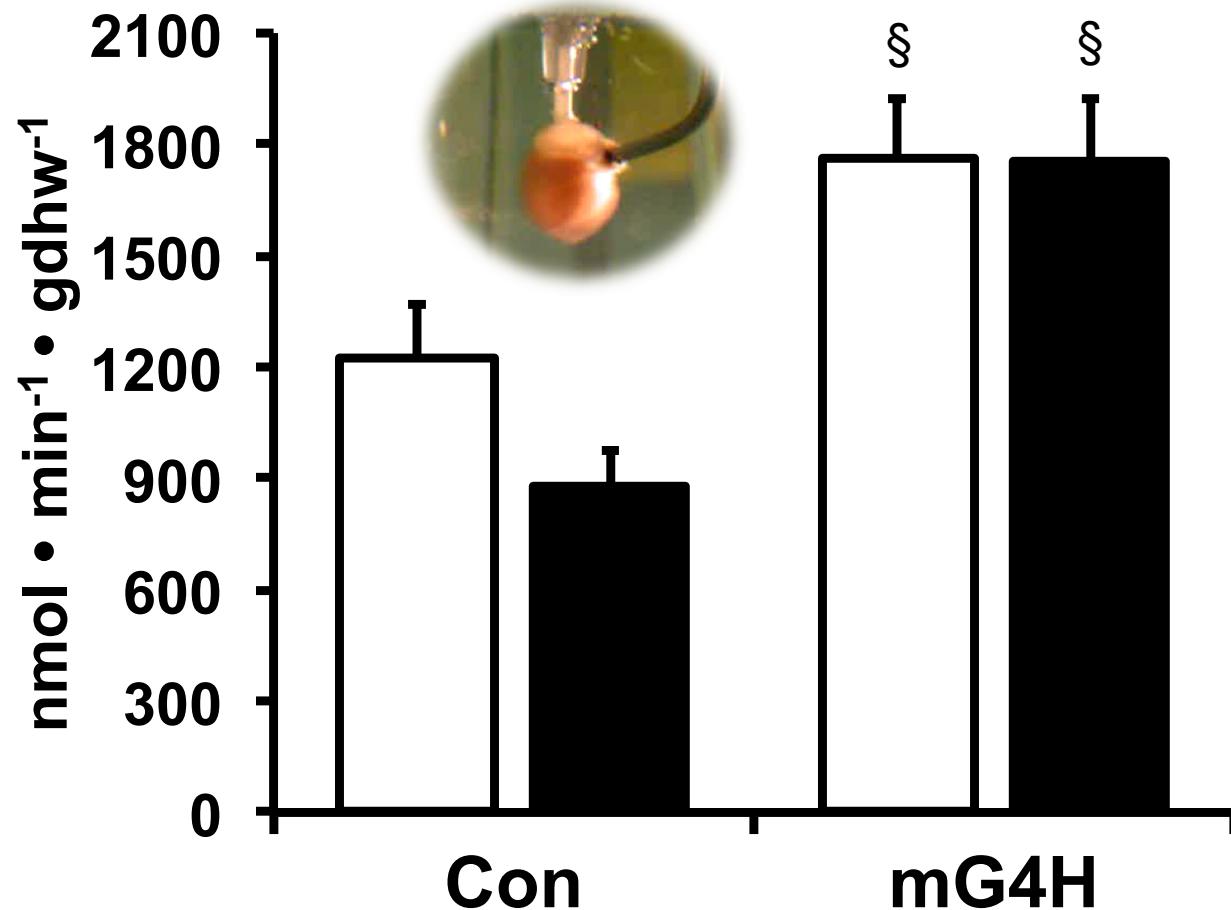


# GLUT4 Induction Increases Glycolysis and Rescues Diabetic Cardiac Glycolytic Defects

Isolated  
Working  
Hearts  
Glycolysis

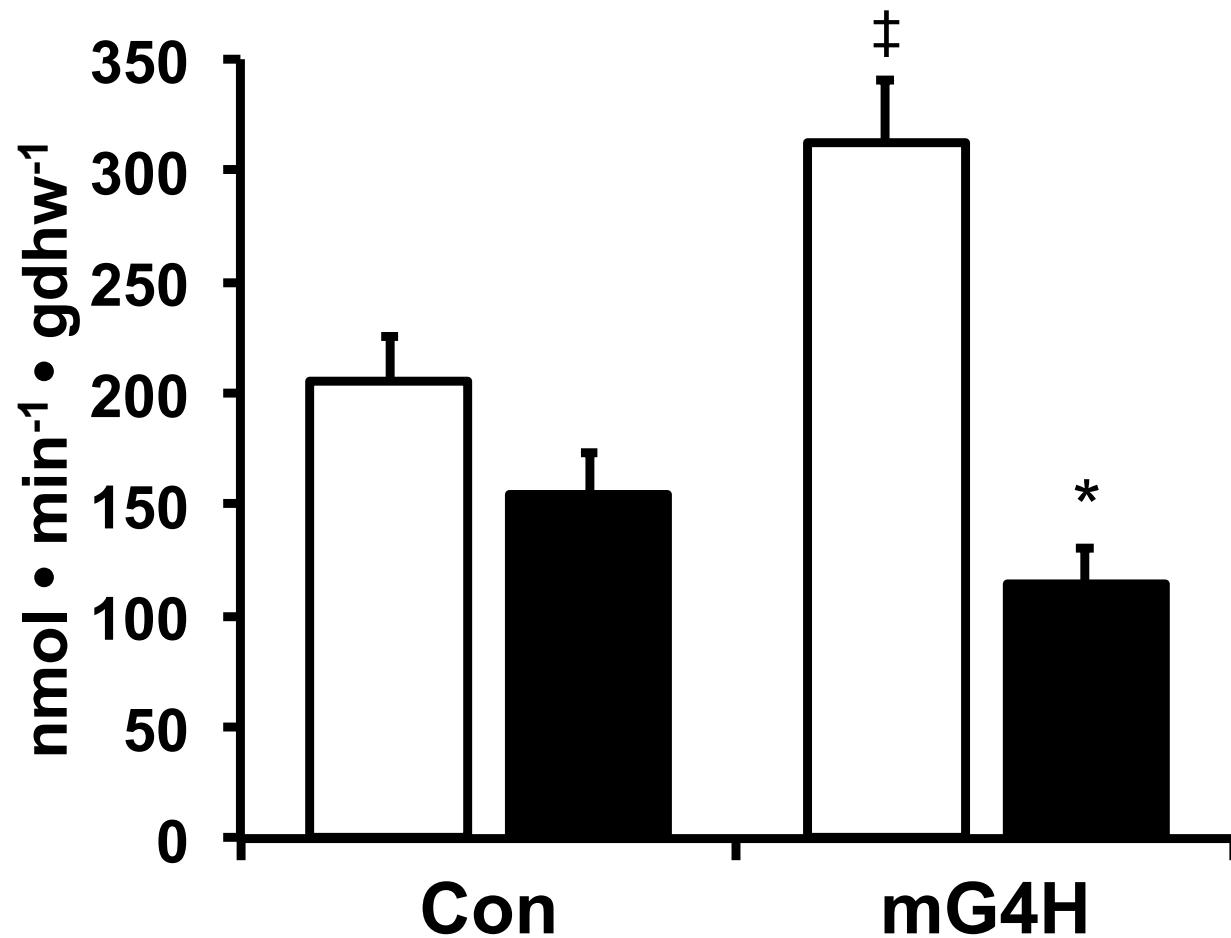
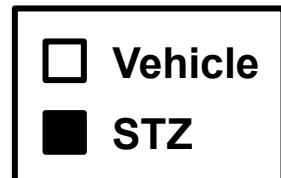


$n = 6 - 10$   
§  $P < 0.01$  vs. Con



# GLUT4 Induction Increases GLOX but Accelerates Diabetic Cardiac GLOX Defects

Isolated  
Working  
Hearts  
Glucose  
Oxidation  
(GLOX)



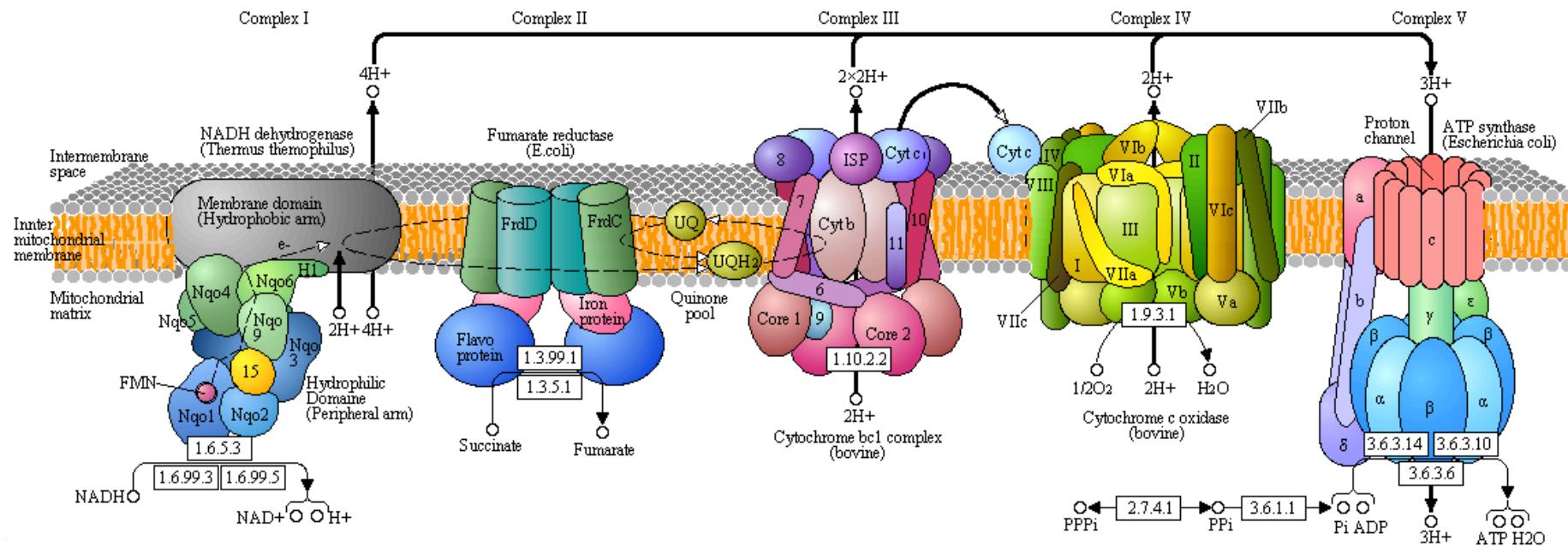
$n = 6 - 10$

§  $P < 0.01$  vs. Con



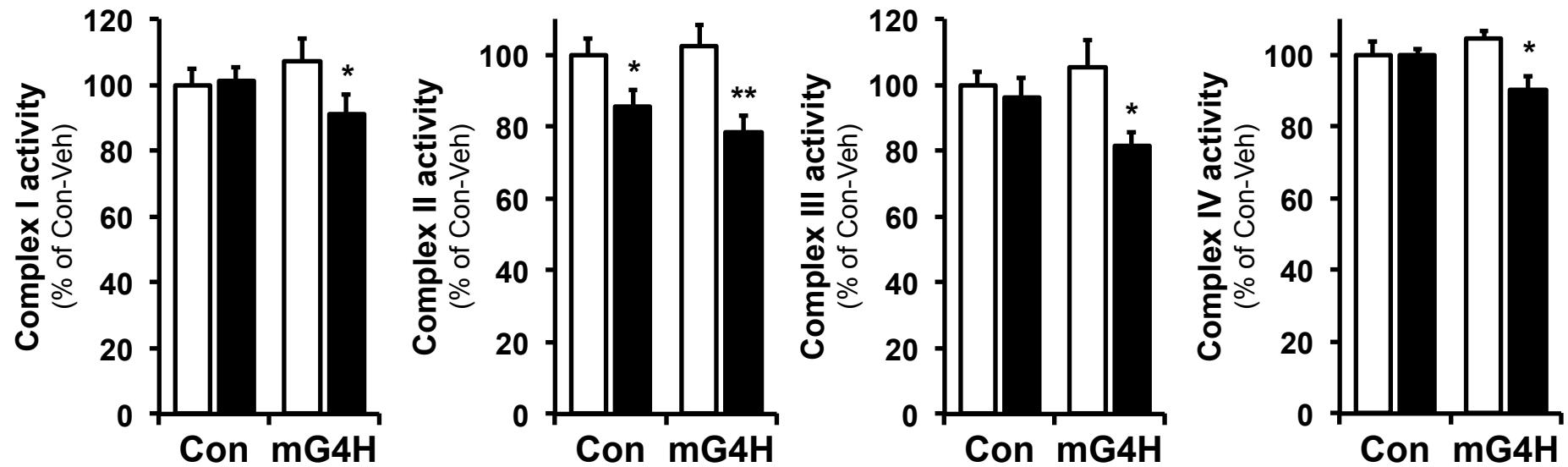
Metabolomics

# Oxidative Phosphorylation



[www.genome.jp/kegg/pathway.html](http://www.genome.jp/kegg/pathway.html)

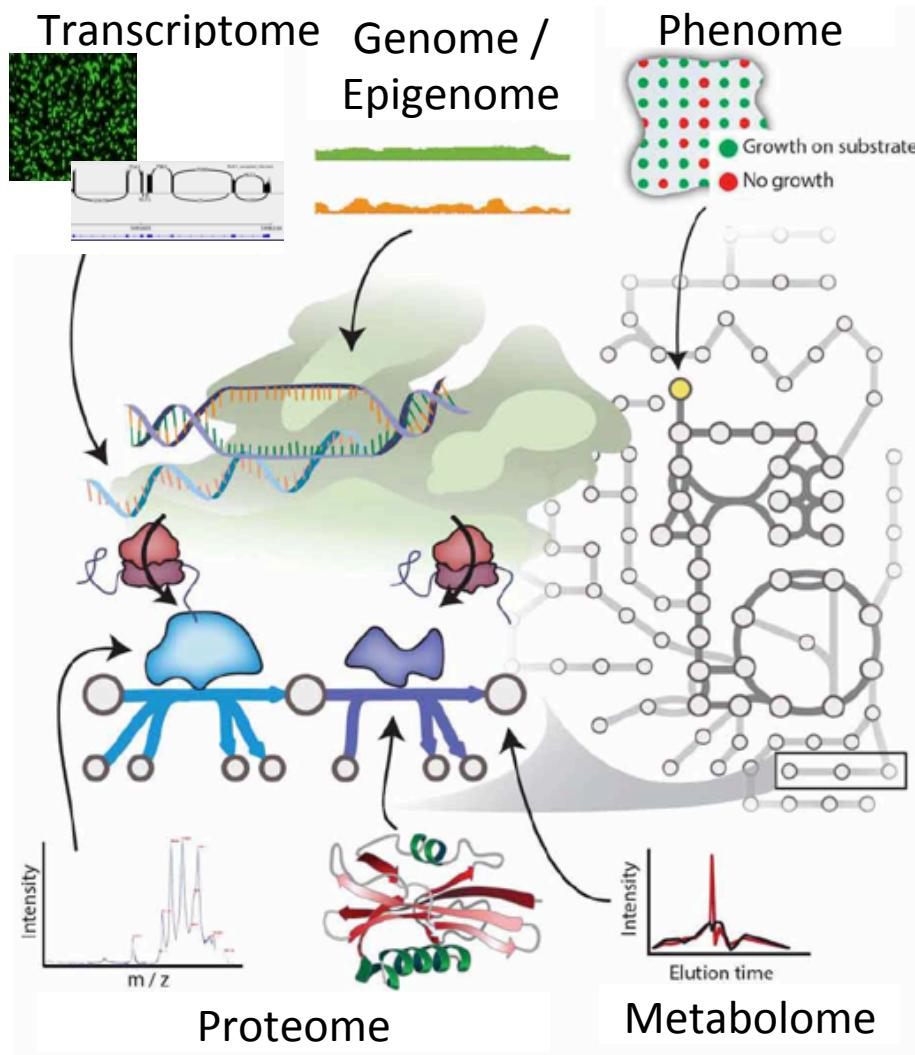
# GLUT4 Induction Accelerates Development of Mitochondrial Dysfunction



$n = 3 - 4$   
\*  $P < 0.05$

Oleh Khalimonchuk  
Wende ... Abel *in prep*

# Systems Biology of the Diabetic Heart



## Phenome

Obesity, diabetes, heart failure, BHI, etc.

## Transcriptome

Northerns, qPCR, microarray  
RNA-seq, miR, lncRNA, etc.

## Proteome

Mass spec, western blot, Co-IP,  
IHC, PTMs, etc.

## Metabolome

Glucometer, ELISA, GC-MS,  
HPLC, NMR, fluxomics, etc.

## Genome / Epigenome

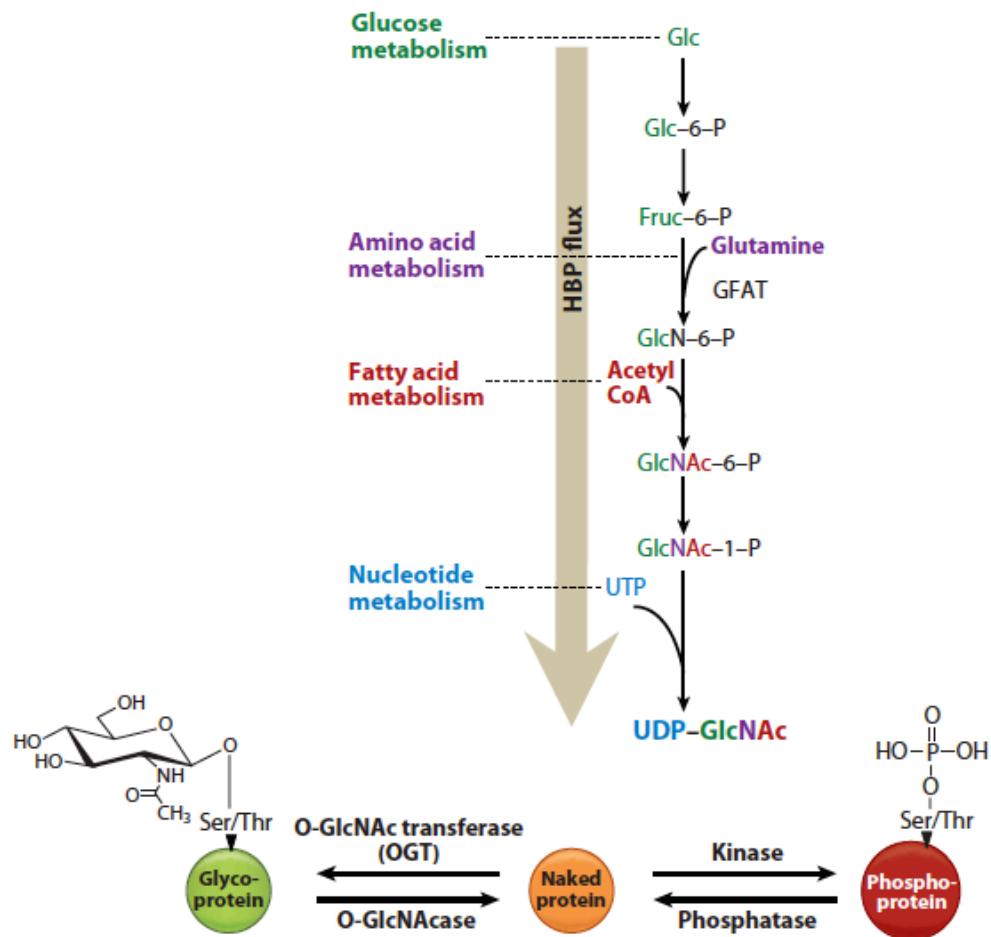
Southerns, sequencing,  
GenBank, ENCODE,  
ChIP-seq, bsDNA-seq, etc.

Adapted from Lewis and Abdel-Haleem **2013** *Front Physiol* 4:237



Metabolomics

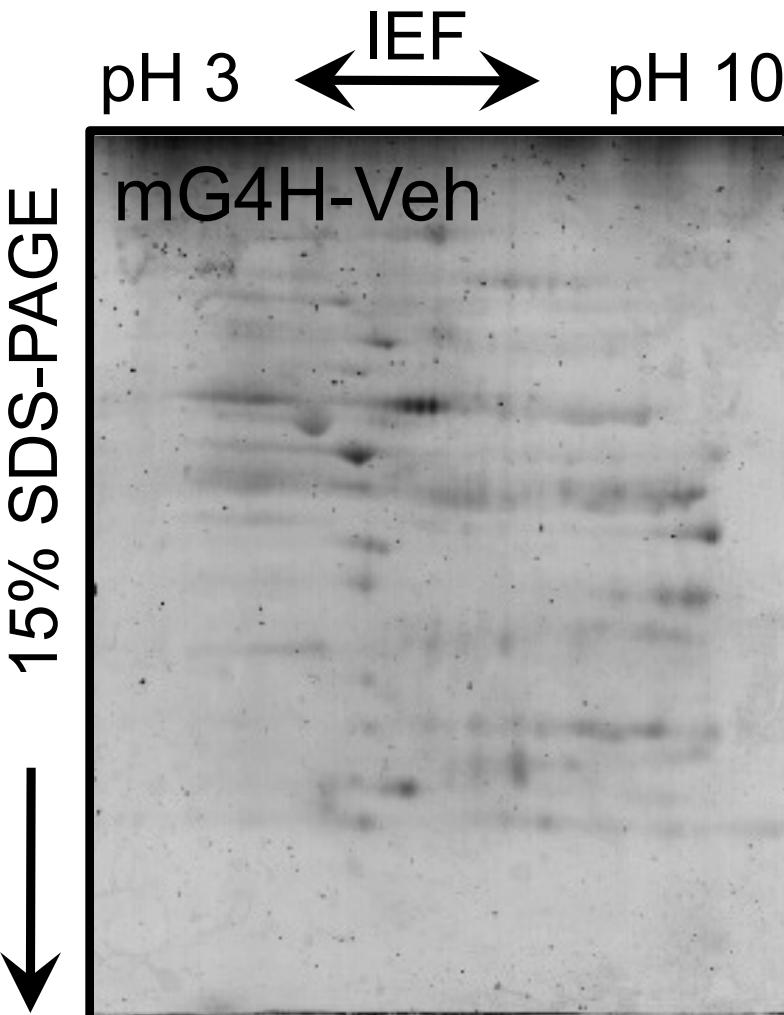
# Metabolism, Bringing the System Together



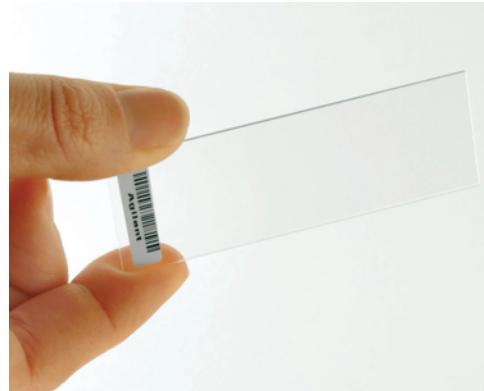
Hart ... Lagerlof 2011 *Annu Rev Biochem* 80:825

# Metabolite Modification of the Proteome

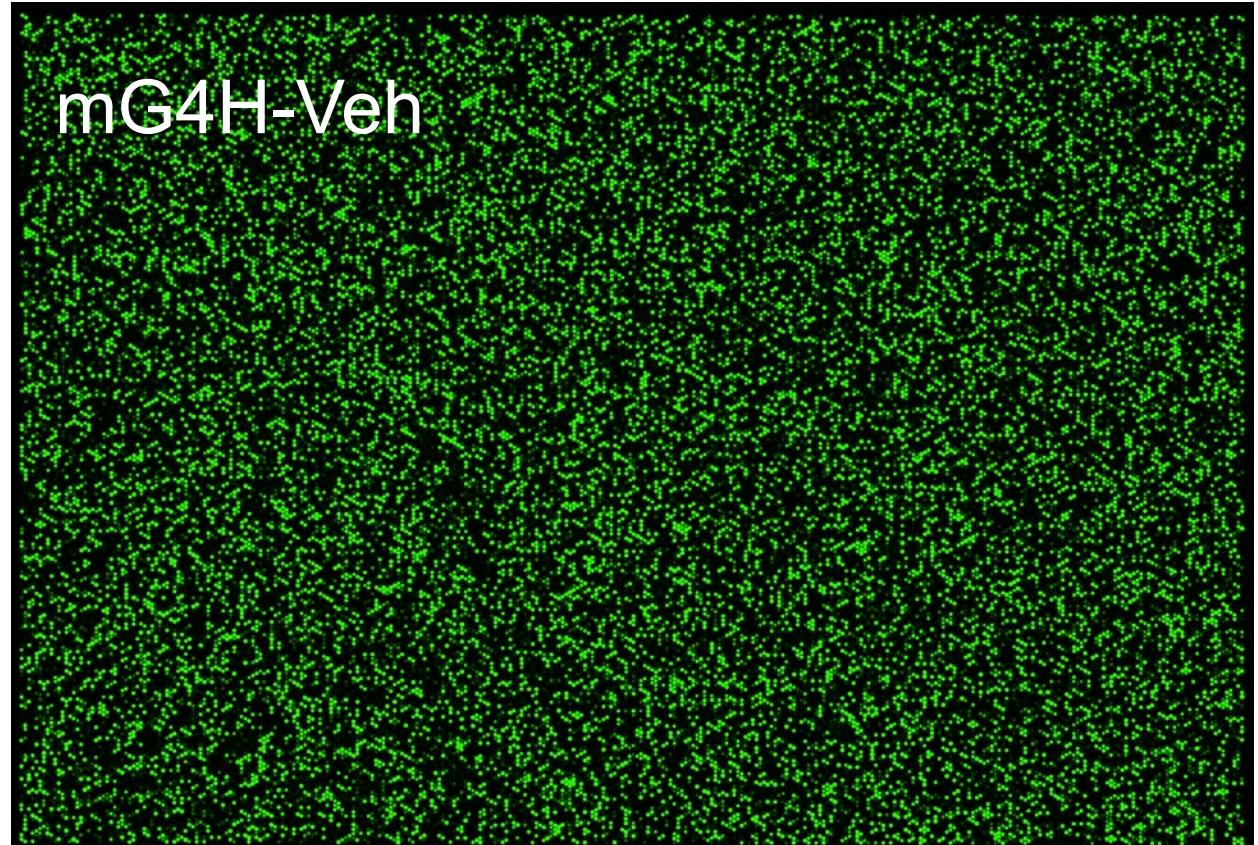
Isolated  
Mitochondria  
2D-PAGE  
Pro-Q  
Emerald



# Metabolite Modification of the Transcriptome



181.9 MB





# pathway analysis of Microarray

## Microarray 2-way ANOVA

Pathway	List	Genes	z-score
<b>Metabolic pathways</b>	194	1085	5.42
Focal adhesion	35	190	2.30
<b>Insulin signaling pathway</b>	27	135	2.49
Axon guidance	25	127	2.30
<b>PPAR signaling pathway</b>	22	76	4.20
Chagas disease	21	102	2.33
Hematopoietic cell lineage	21	83	3.39
Peroxisome	21	78	3.72
<b>Dilated cardiomyopathy</b>	20	86	2.88
<b>Hypertrophic cardiomyopathy</b>	20	83	3.06
Arrhythmogenic right ventricular cardiomyopathy	19	71	3.50
ECM-receptor interaction	19	79	2.97
<b>Glycolysis / Gluconeogenesis</b>	19	68	3.72
Pyrimidine metabolism	19	95	2.08
Valine, leucine and isoleucine degradation	18	47	5.21
Metabolism of xenobiotics by cytochrome P450	16	64	2.90
Leishmaniasis	15	64	2.53
Glutathione metabolism	14	53	2.95
p53 signaling pathway	14	66	2.02
Arginine and proline metabolism	13	52	2.61
Graft-versus-host disease	13	49	2.85
Lysine degradation	13	41	3.60
<b>Type II diabetes mellitus</b>	13	47	3.03
Fatty acid metabolism	12	46	2.68
<b>Type I diabetes mellitus</b>	12	53	2.12

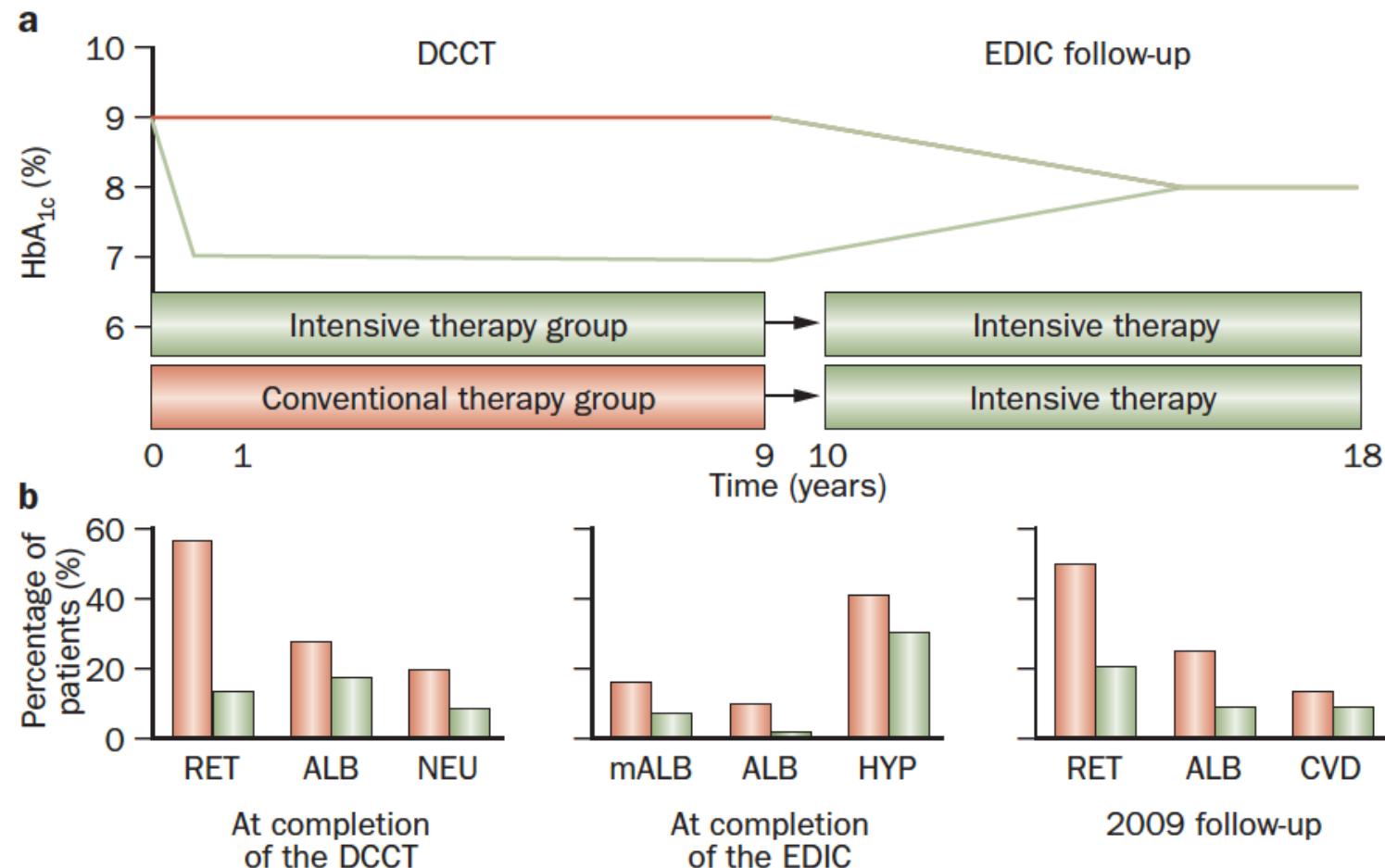
# From Human to Mouse and Back Again...



Broad Institute Communications

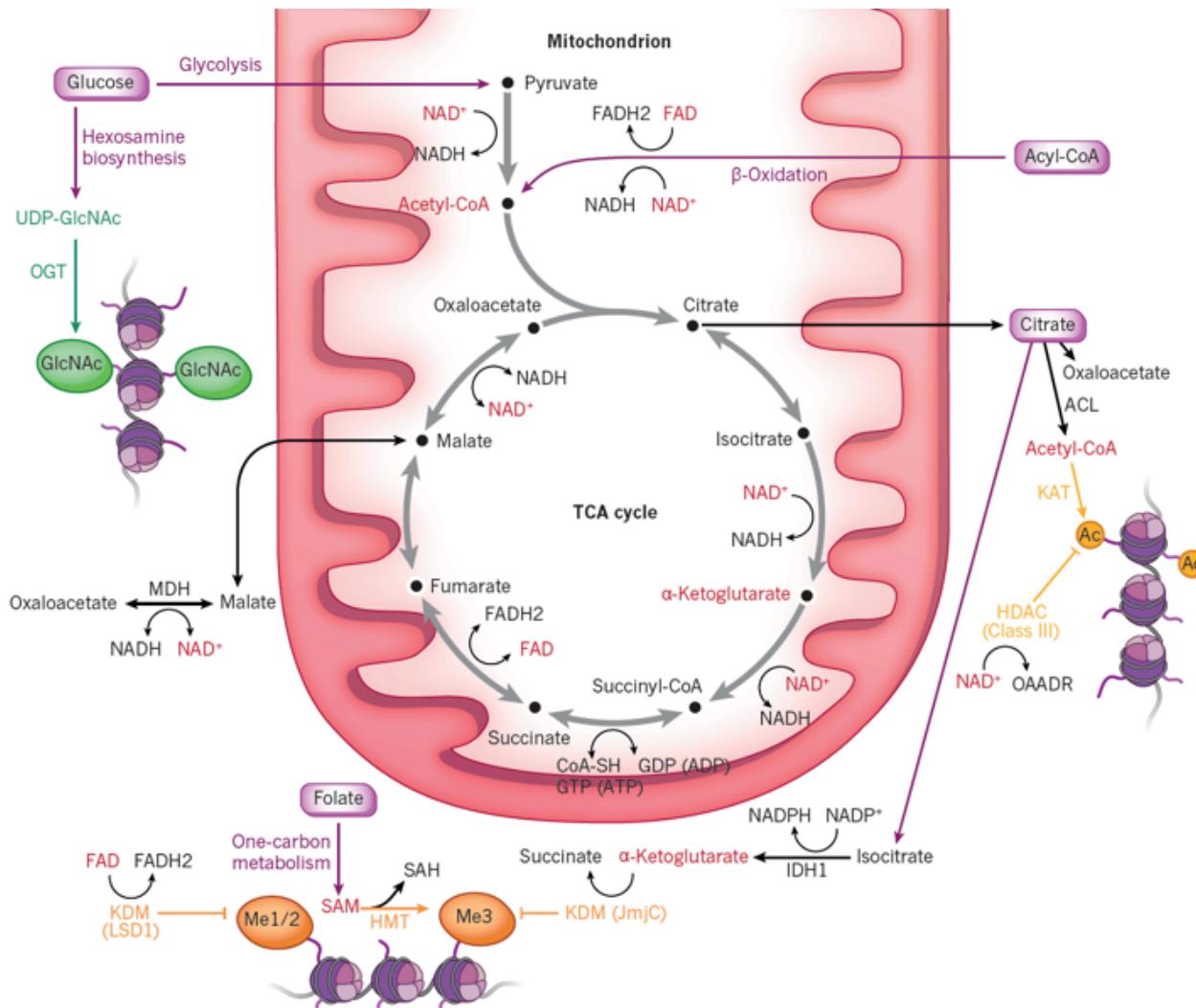
# Epigenetics - Memory

## EDIC: Epidemiology of Diabetes Interventions Trial



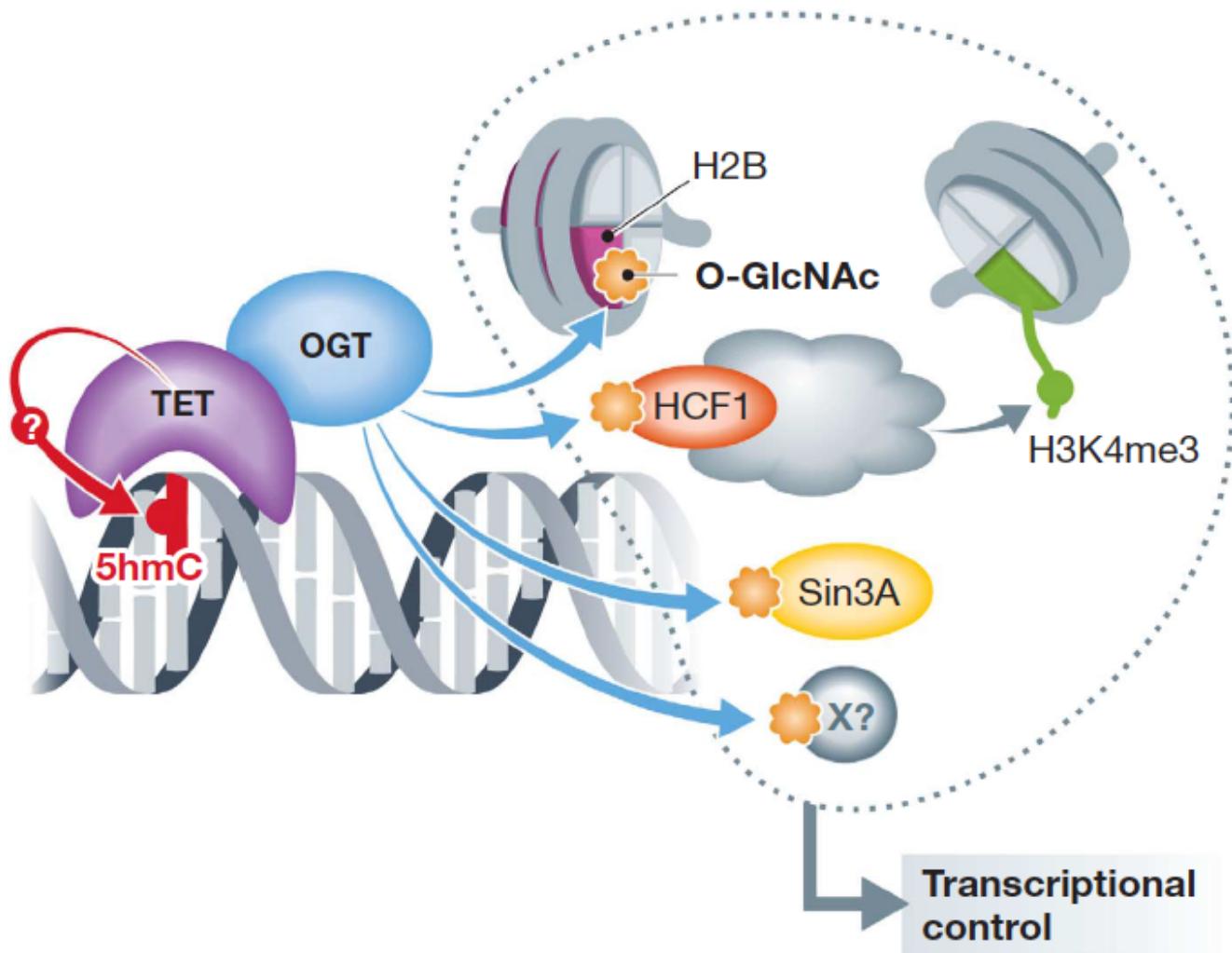
Pirola ... El-Osta 2010 Nat Rev Endocrinol 6(12):665

# Metabolite Signaling to Epigenetics



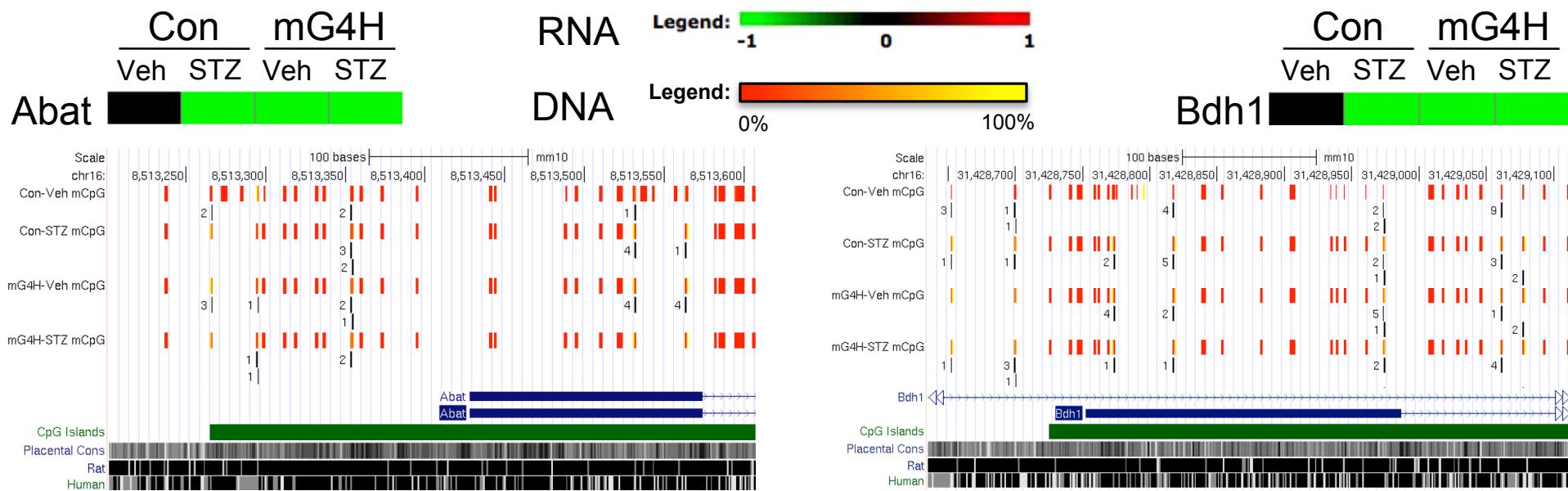
Gut and Verdin 2013 Nature 502:489

# How does GlcNAc fit in?



Mariappa ... Aalten 2013 *EMBO J* 32:612

# Metabolite Modification of the Methylome



# Human/Mouse Comparisons

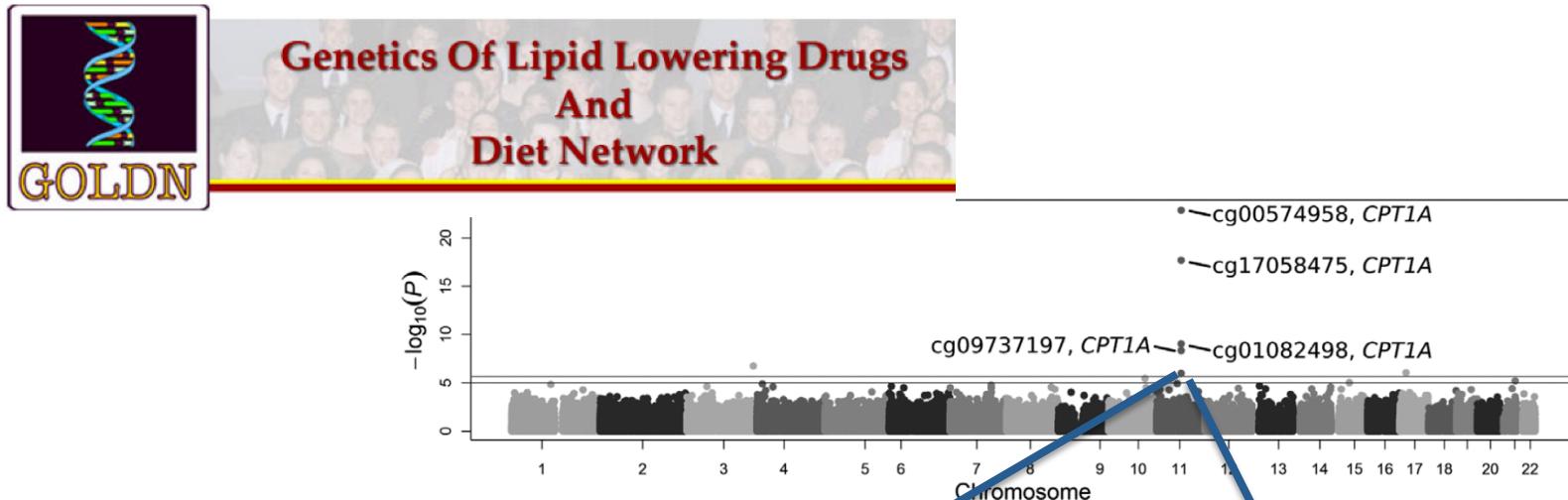
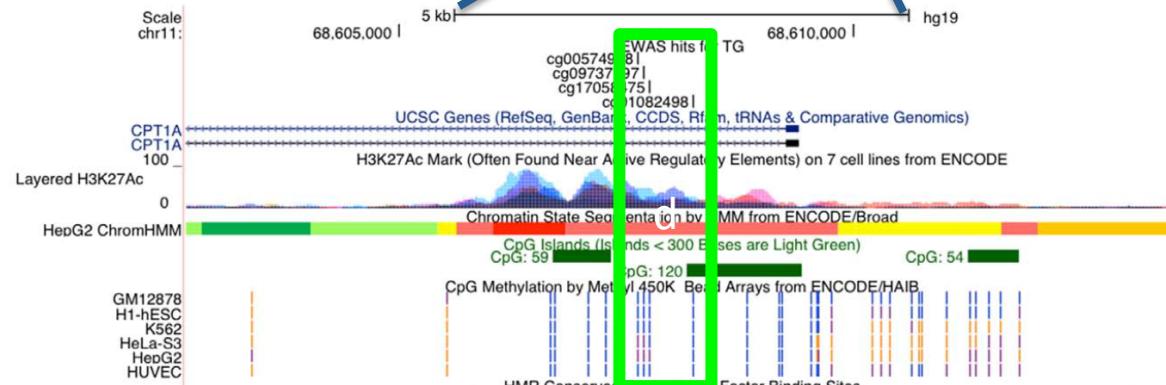


Figure 2. Epigenome-wide association Manhattan plot for VLDL-C in the discovery dataset ( $n=991$ ). VLDL-C indicates very-low-density lipoprotein cholesterol.



Irvin ... Arnett 2014 Circulation 130:565

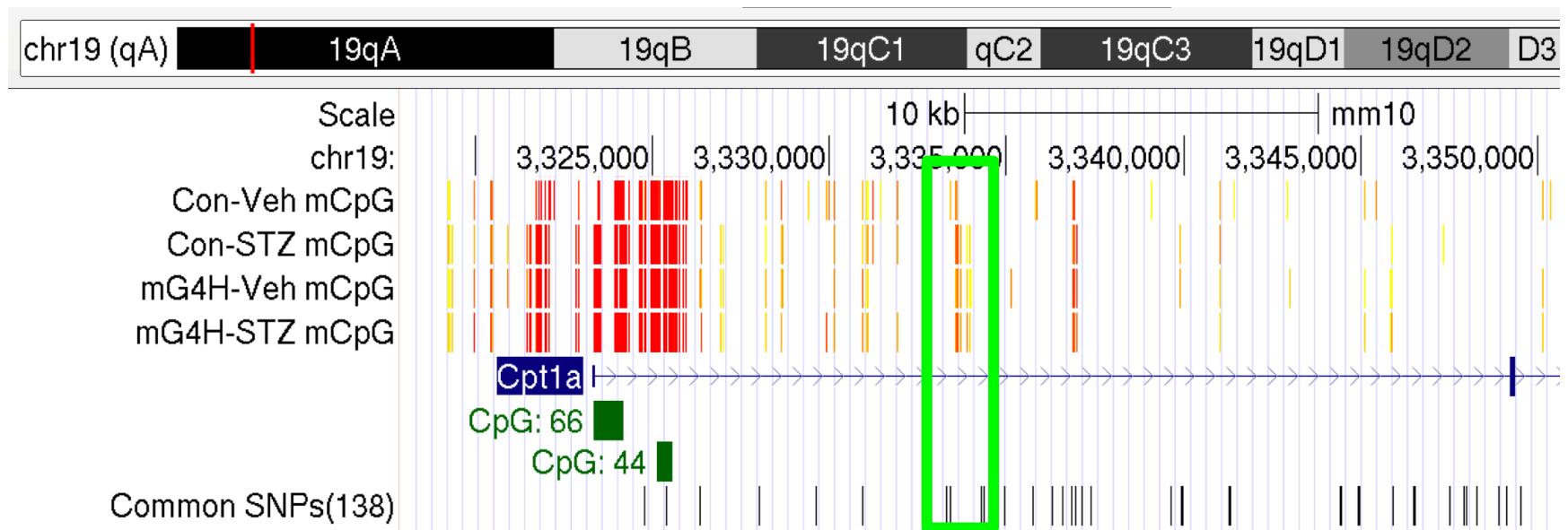
# Human/Mouse Comparisons

## Mouse Gene Expression

Con Con mG4H mG4H    GENE  
Veh STZ Veh STZ

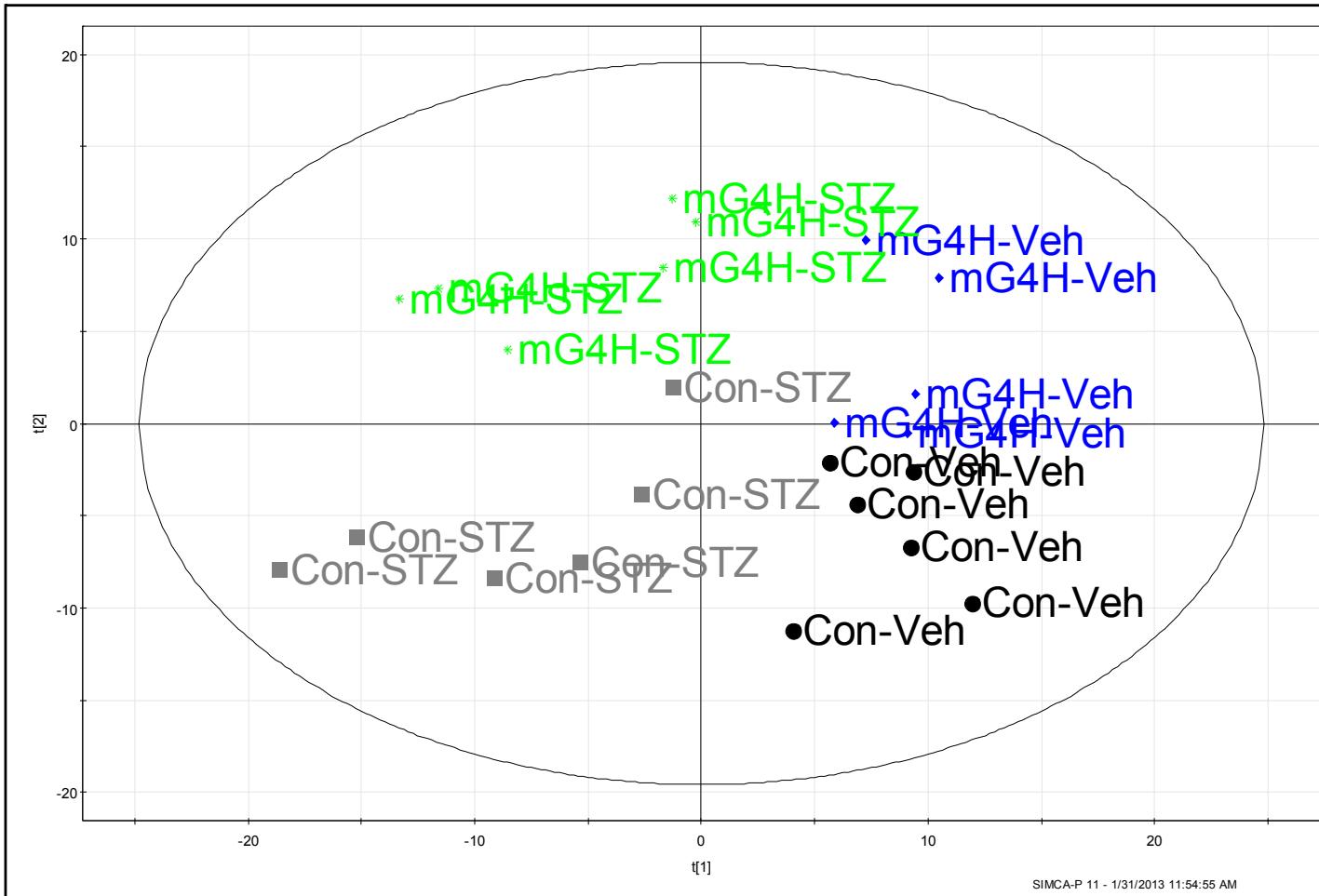


## Mouse DNA Methylation

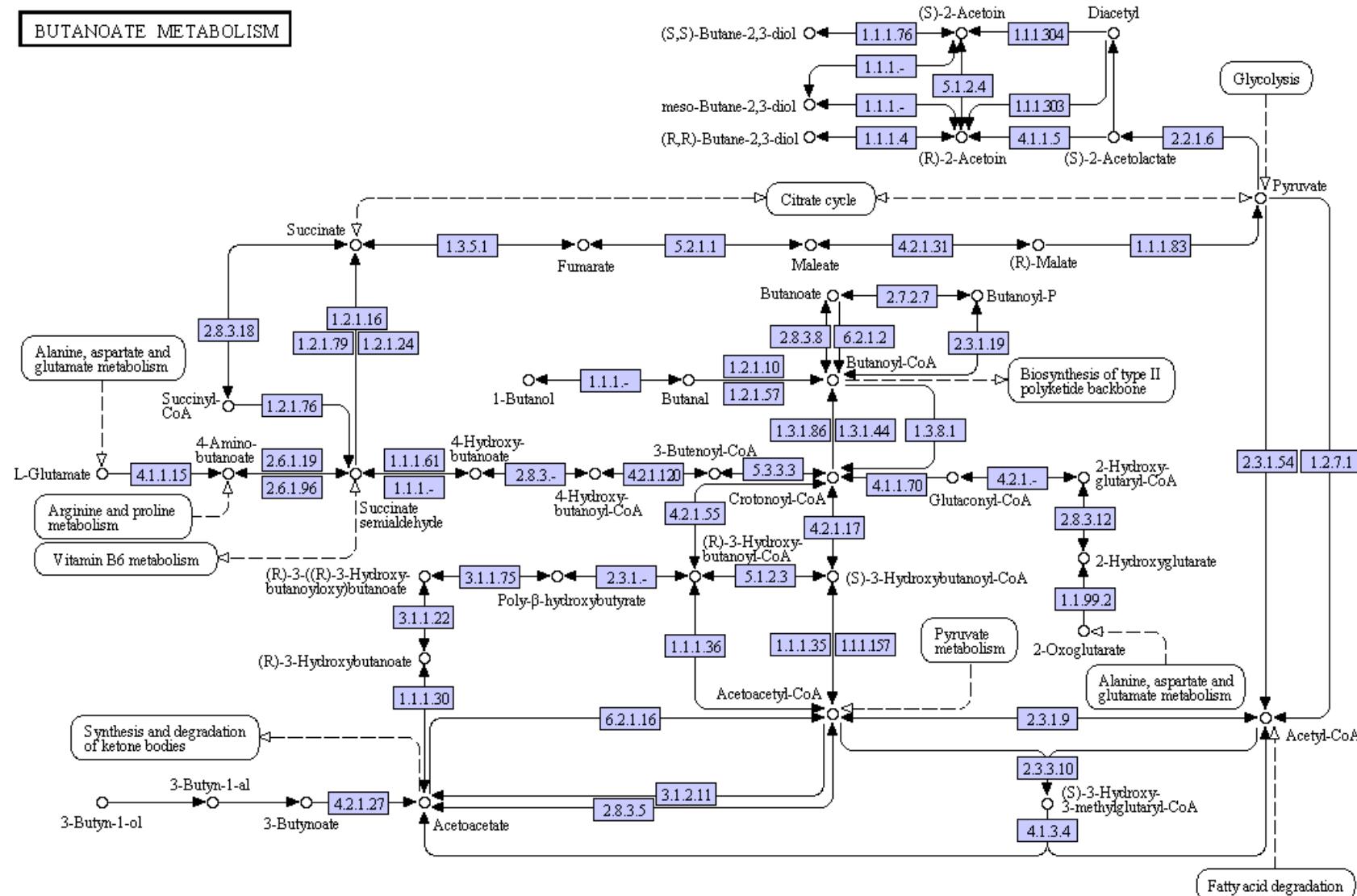


Wende, unpublished

# Metabolite Modification of the Metabolome



# Metabolite Modification of the Metabolome



Metabolomics

# Acknowledgements

## Wende Lab

Manoja Brahma – Postdoc  
Mark C. McCrory – Manager  
Brenna G. Nye – Undergrad  
Mark Pepin – MSTP  
Lamario J Williams – Undergrad

## E. Dale Abel – Utah to Iowa

John C. Schell – MD/PhD student  
Joseph Tuinei – Industry

## Hansjörg Schwertz – GU, Germany



JDRF 51002608



Molecular & Cellular Pathology  
Redox Biology  
Diabetes Center  
Cardiovascular Center



## UAB Collaborators

Steve Barnes – Metabolomics  
John C. Chatham – GlcNAc  
David Crossman – Genomics/Informatics  
Farah D. Lubin – Epigenetics

## Zymo Research

Keith Booher – DNA methylation  
Hunter Chung – Informatics

## U of U Cores

James Cox – Metabolomics  
Brett Milash – Genomics/Informatics  
Krishna Parsawar – Proteomics



R00 HL111322