



GBSC 724: Advanced Topics in Metabolomics

Metabolomics in Models of Cardiovascular Disease

Wednesday, March 15, 2017

Adam R. Wende, Ph.D.

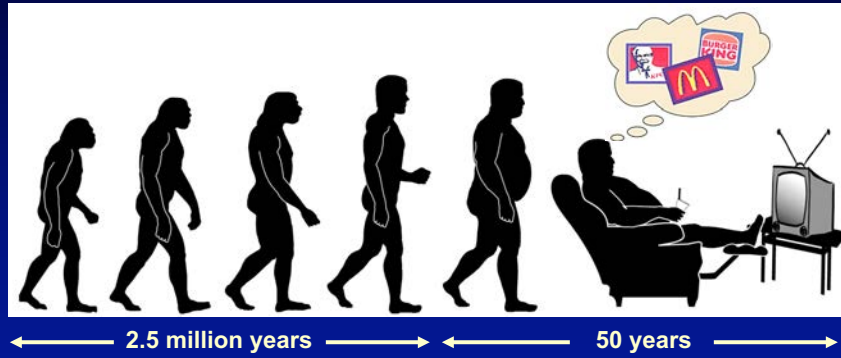
Assistant Professor
Division of Molecular and Cellular Pathology



Outline

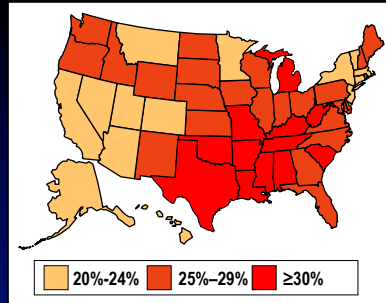
- Define the question and model to determine the connection between metabolism and diabetic heart disease.
- Identify the molecular mechanisms by which glucose directly alters molecular function using systems biology.
 - Transcriptomics
 - Proteomics
 - Metabolomics
 - Epigenetics (e.g. methylomics)

Obesity, Metabolic Syndrome, Diabetes, and Heart Failure

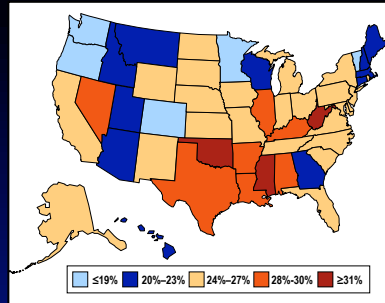


From: Roger Unger - UTSW

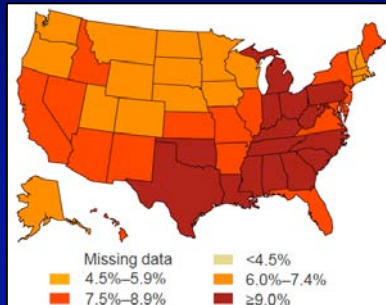
2010 – Obesity



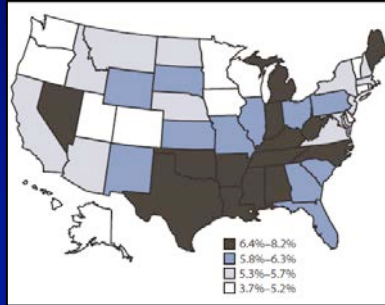
2010 – Physical Inactivity



2010 – Diabetes

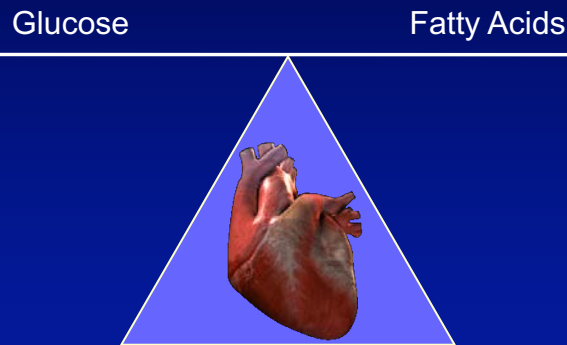


2010 – Heart Disease



www.cdc.gov/diabetes/statistics and www.cdc.gov/mmwr

Maintaining Cardiac Function Through Metabolic Substrate Balance



giphy.com

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.

lactate usage and a slight decline in that of pyruvate. There is no change in utilization of 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 of diabetes is not clear. Anesthesia, which was used in the dogs, may have played some part.

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

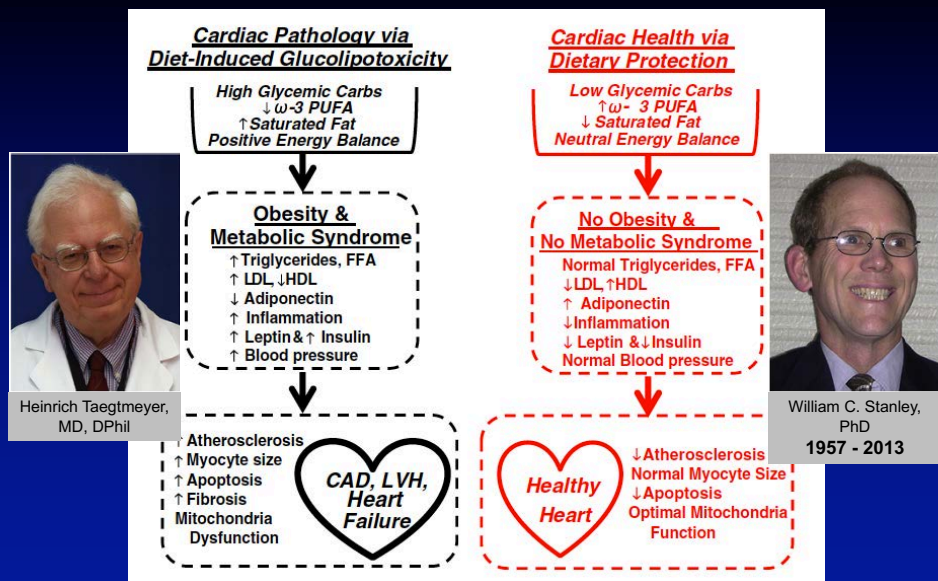
Metabolic Substrate Utilization in the Heart

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

	MV ₀ ₂	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

Point/Counterpoint - The Right Balance?



Taegtmeier and Stanley 2011 *J Mol Cell Cardiol* 50(1):2

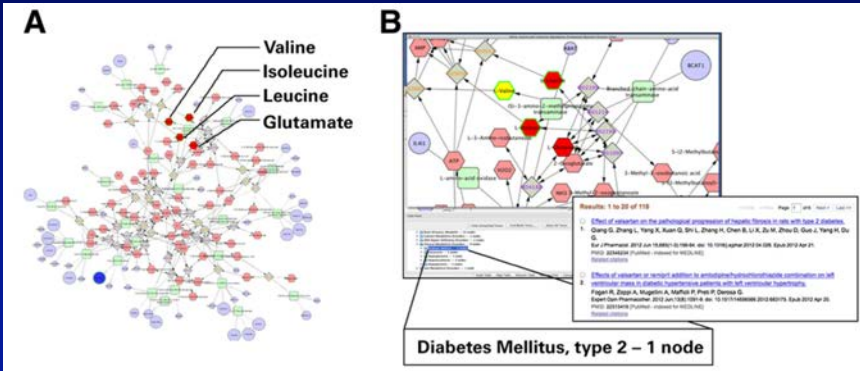
Diabetes and Metabolomics

Diabetes. 2015 Mar;64(3):718-732.

Metabolomics and Diabetes: Analytical and Computational Approaches.

Sas KM¹, Karnovsky A², Michailidis G³, Pennathur S⁴.

Metabolomics is an integral part for understanding disease processes ... information garnered in the biomarker investigations, future research should shed more light on disease pathogenesis and explore new treatment options.



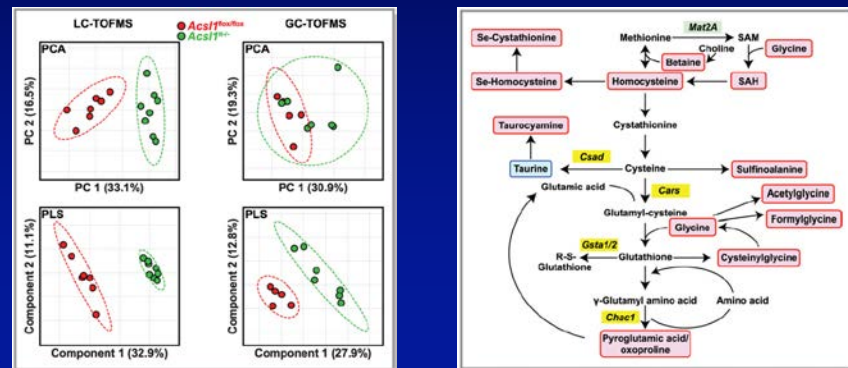
Heart failure and substrate switching

J Am Heart Assoc. 2015 Feb 24;4(2). pii: e001136. doi: 10.1161/JAHA.114.001136.

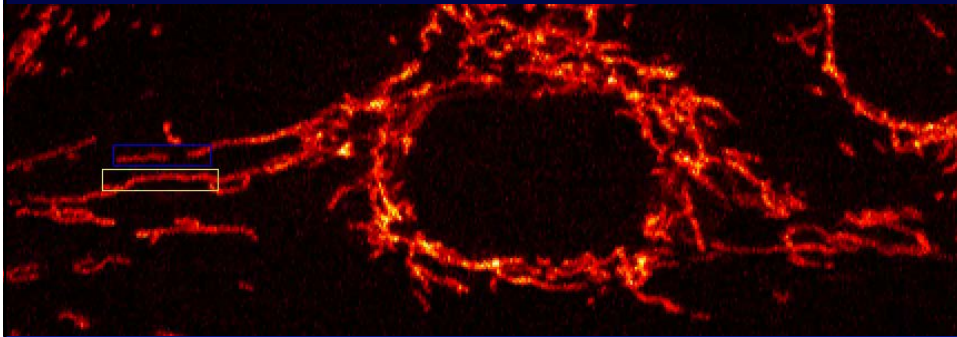
Cardiac energy dependence on glucose increases metabolites related to glutathione and activates metabolic genes controlled by mechanistic target of rapamycin.

Schisler JC¹, Grevengoed TJ², Pascual F², Cooper DE², Ellis JM², Paul DS², Willis MS³, Patterson C¹, Jia W⁴, Coleman RA².

The hypertrophy, oxidative stress, and metabolic changes that occur within the heart when glucose supplants FA as a major energy source suggest that substrate switching to glucose is not entirely benign.



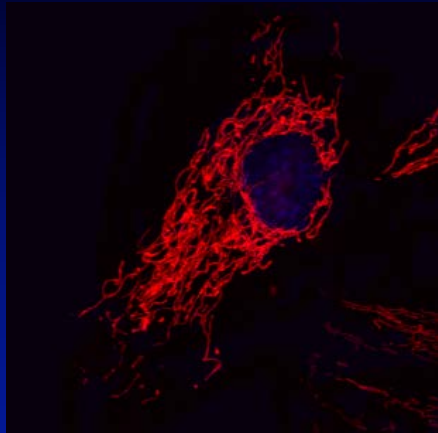
Mitochondria – a Dynamic Network



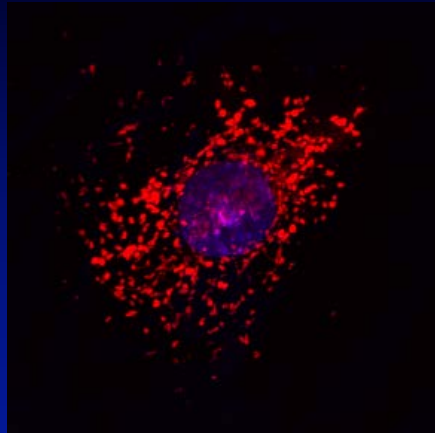
Fan ... Brooks 2010 *Free Radic Biol Med* 49(11):1646

Mitochondria – too much fat

Serum

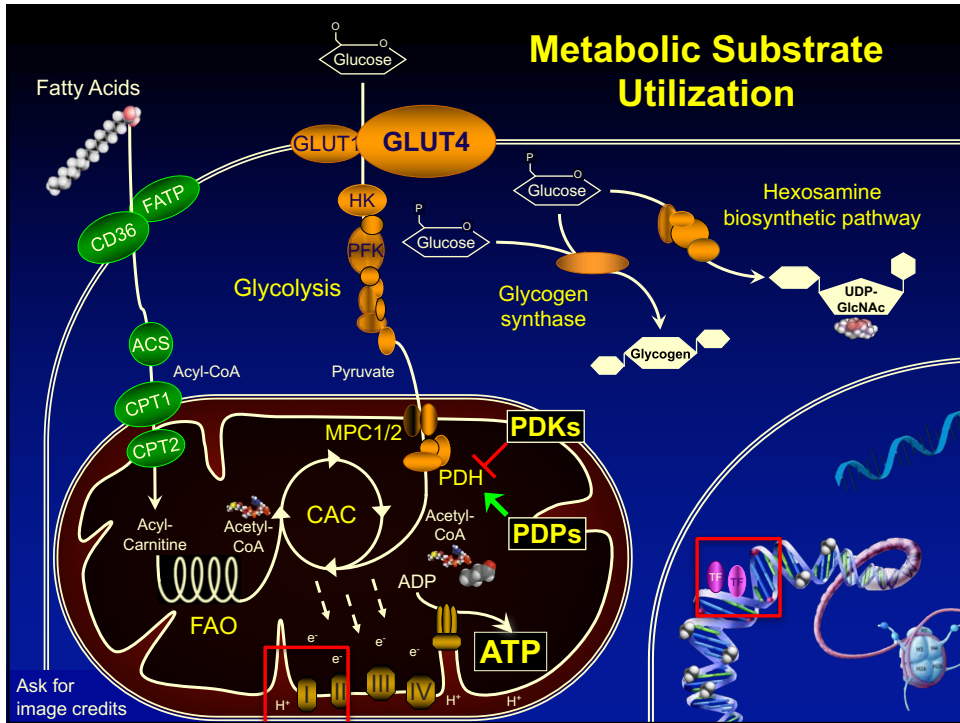


Serum + 500 μ M Palmitate

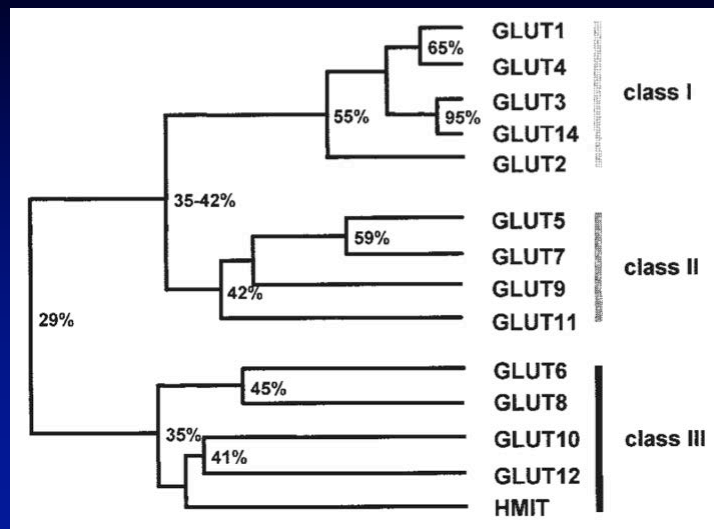


Adapted from Heiko Bugger





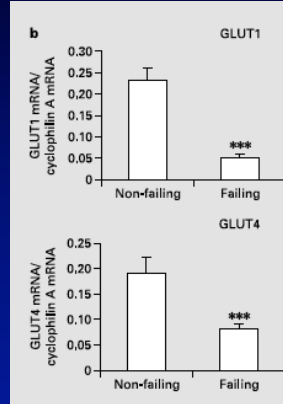
Facilitative Glucose Transporters: GLUTs "Solute Carrier Family, SLC2A"



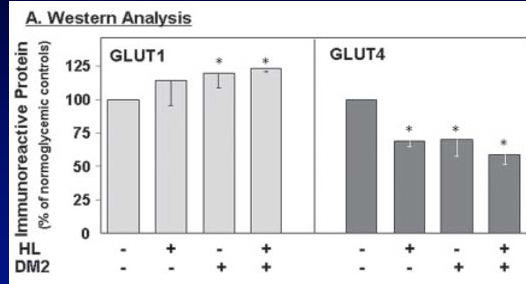
Scheepers ... Schurmann 2004 *J Parenter Enteral Nutr* 28:364

Changes in Human Heart GLUT Levels

RNA
Human heart failure



Protein
Human heart diabetes



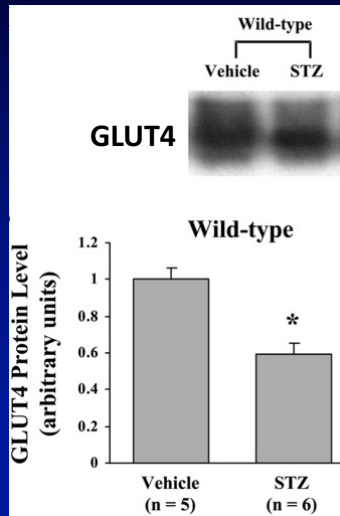
Biopsies obtained during coronary bypass surgery
HL = hyperlipidemia
DM2 = diabetes mellitus type 2

Razeghi ... Taegtmeier 2002 *Cardiology* 280(41):34786

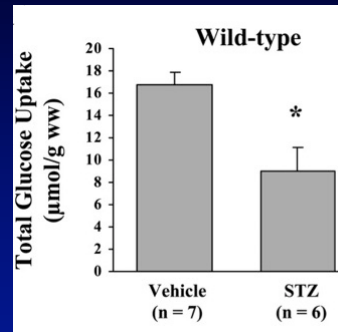
Armoni ... Karnieli 2005 *J Biol Chem* 280(41):34786

Glucose Utilization and Rodent Models of Type 1 Diabetes

Protein
Diabetic Mouse Heart

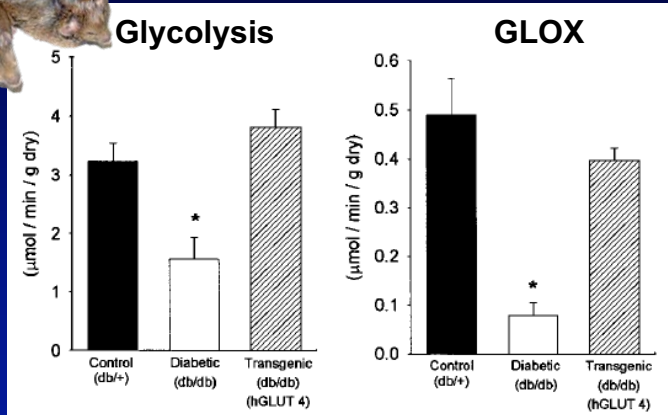
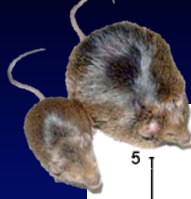


Glucose Uptake
Diabetic Mouse Heart



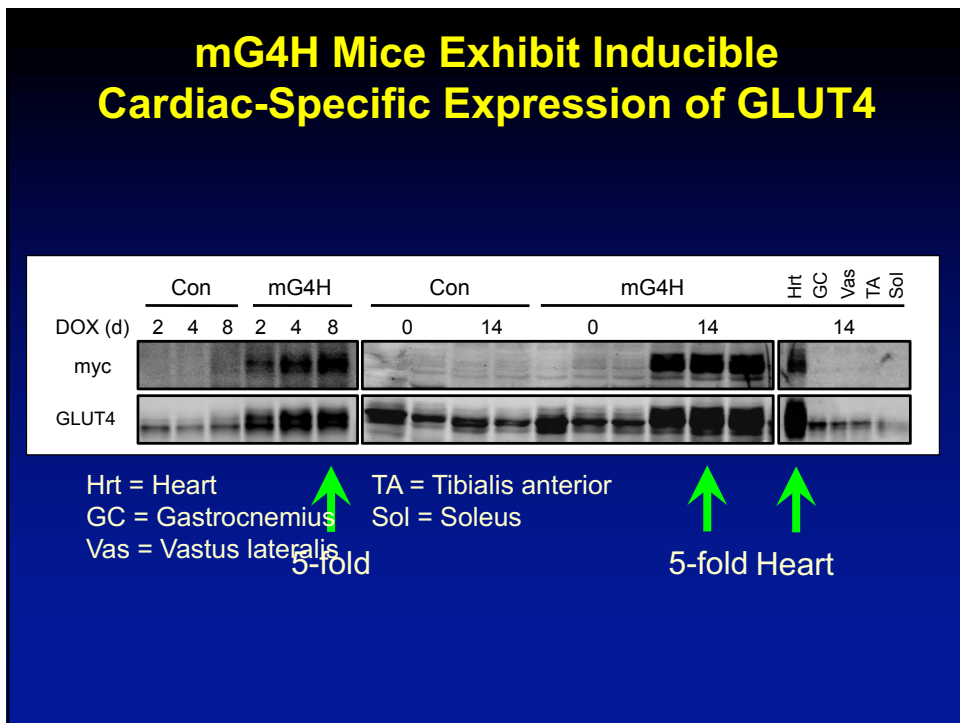
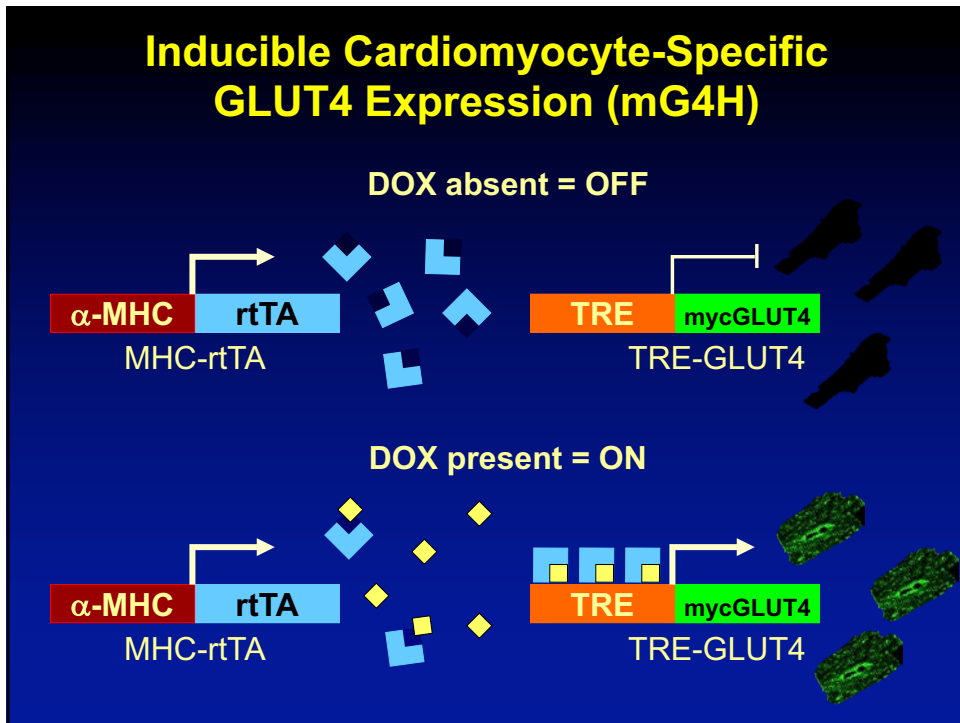
Panagia ... Clarke 2005 *Am J Physiol* 288:H2677

Constitutive GLUT4 Expression Prevents Development of Glucose Utilization Defects

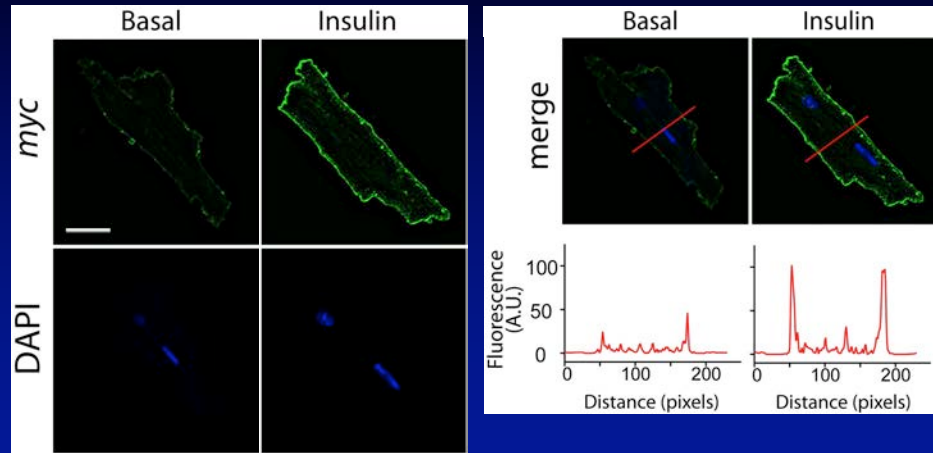


Belke ... Severson 2000 *Am J Physiol* 279:E1104

Question: Is the change in cardiac metabolic substrate flexibility adaptive or maladaptive?



Insulin-induced GLUT4 Vesicle Fusion and Exofacial Myc-Epitope Exposure

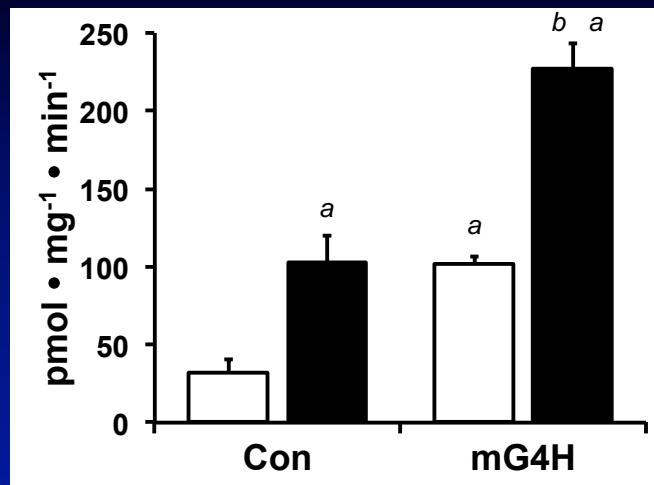


Ariel Contreras-Ferrat
Wende ... Abel *in prep*

GLUT4 Induction Increases Basal and Insulin-Stimulated Glucose Uptake

Cardiac
Myocytes
2-DG
Uptake

□ Basal
■ 0.1 nM Ins



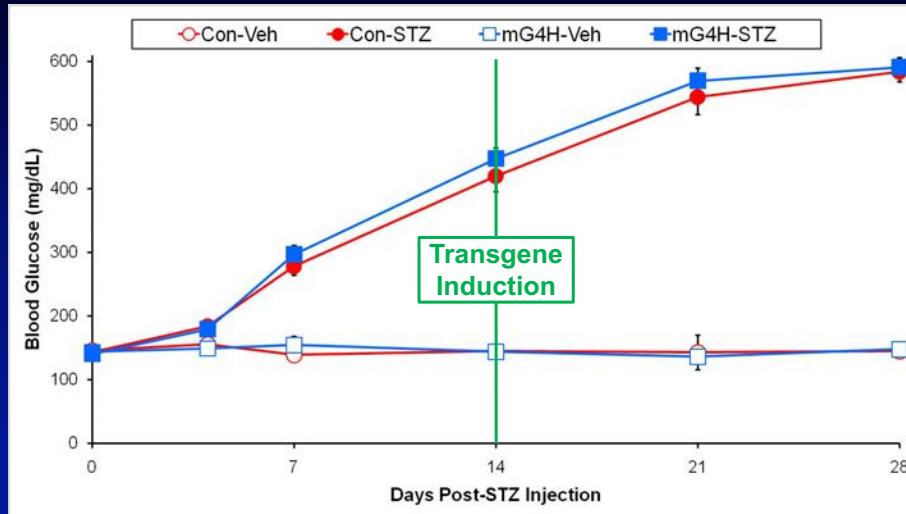
$n = 3 - 4$

^a $P < 0.01$ vs. Con-Basal

^b $P < 0.001$ vs. All

Renata O. Pereira
Wende ... Abel *in prep*

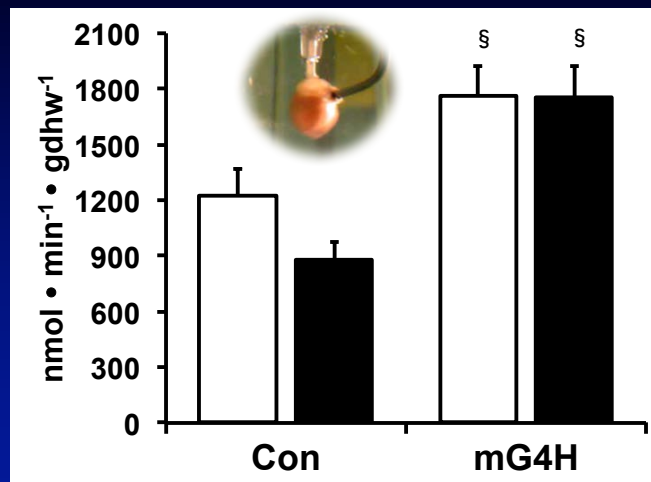
Streptozotocin (STZ)-Induced Hyperglycemia is Not Altered by Transgene Induction



GLUT4 Induction Increases Glycolysis and Rescues Diabetic Cardiac Glycolytic Defects

Isolated Working Hearts Glycolysis

□ Vehicle
■ STZ



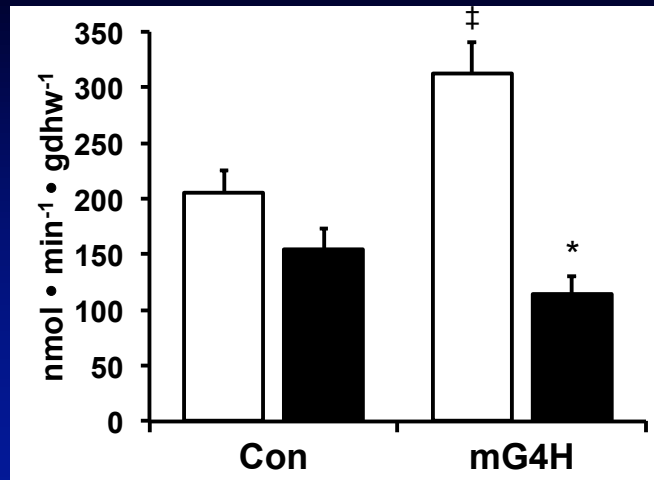
$n = 6 - 10$
§ $P < 0.01$ vs. Con

Joseph Tuinei
Wende ... Abel *in prep*

GLUT4 Induction Increases GLOX but Accelerates Diabetic Cardiac GLOX Defects

Isolated Working Hearts
Glucose Oxidation (GLOX)

□ Vehicle
■ STZ



$n = 6 - 10$

[‡] $P < 0.001$ vs. All

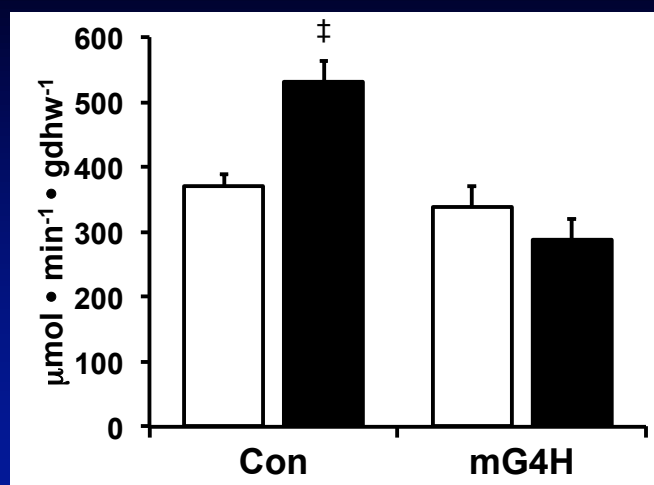
^{*} $P < 0.01$ vs. Veh

Joseph Tuinei
Wende ... Abel *in prep*

GLUT4 Induction Prevents Increased Cardiac POX in Diabetes

Isolated Working Hearts
Palmitate Oxidation (POX)

□ Vehicle
■ STZ

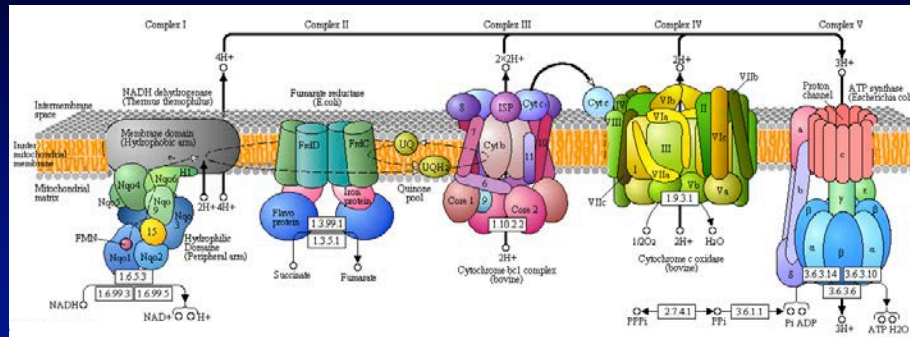


$n = 5 - 13$

[‡] $P < 0.001$ vs. All

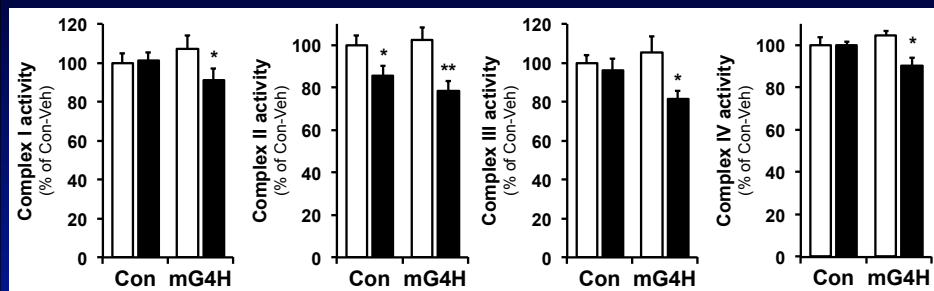
Joseph Tuinei
Wende ... Abel *in prep*

Oxidative Phosphorylation



www.genome.jp/kegg/pathway.html

GLUT4 Induction Accelerates Development of Mitochondrial Dysfunction



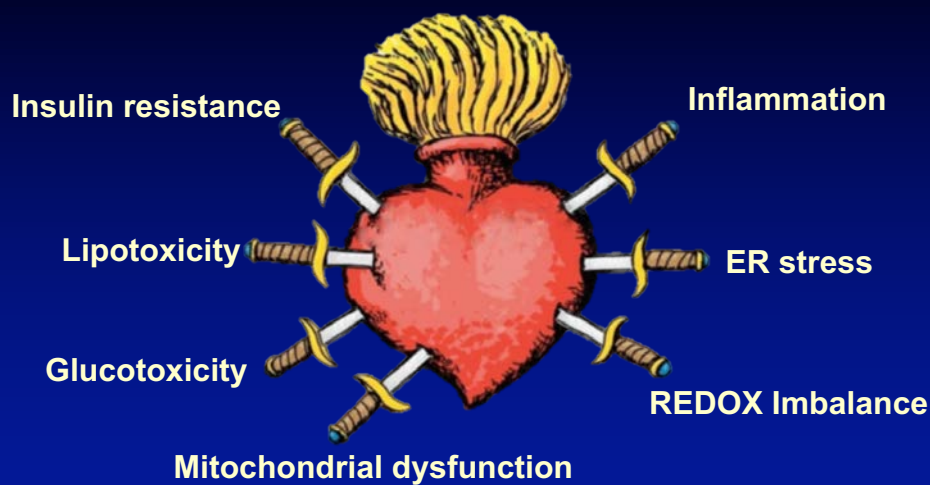
$n = 3 - 4$
* $P < 0.05$

Oleh Khalimonchuk
Wende ... Abel *in prep*

Conclusion – Part 1

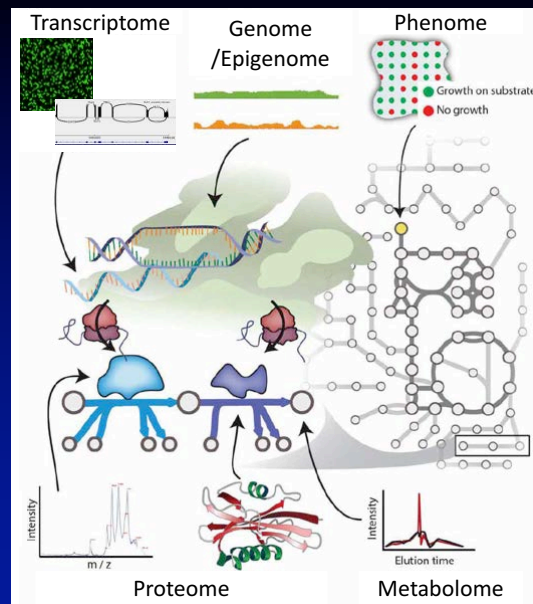
In the context of diabetes, enhancing glucose delivery by expression of GLUT4 accelerates the progression of mitochondrial dysfunction.

Diabetic Cardiomyopathy “Death by a Thousand Cuts...”



Adapted from Wende, Symons, and Abel 2012 *Curr Hypertens Rep* 14(6):517

Systems Biology



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

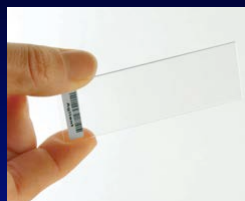
Glucometer, ELISA, GC-MS,
HPLC, NMR, fluxomics, etc.

Genome / Epigenome

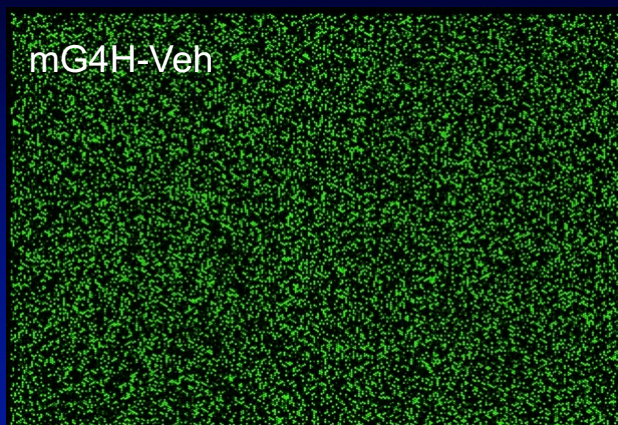
Southerns, sequencing,
GenBank, ENCODE,
ChIP-seq, bsDNA-seq, etc.

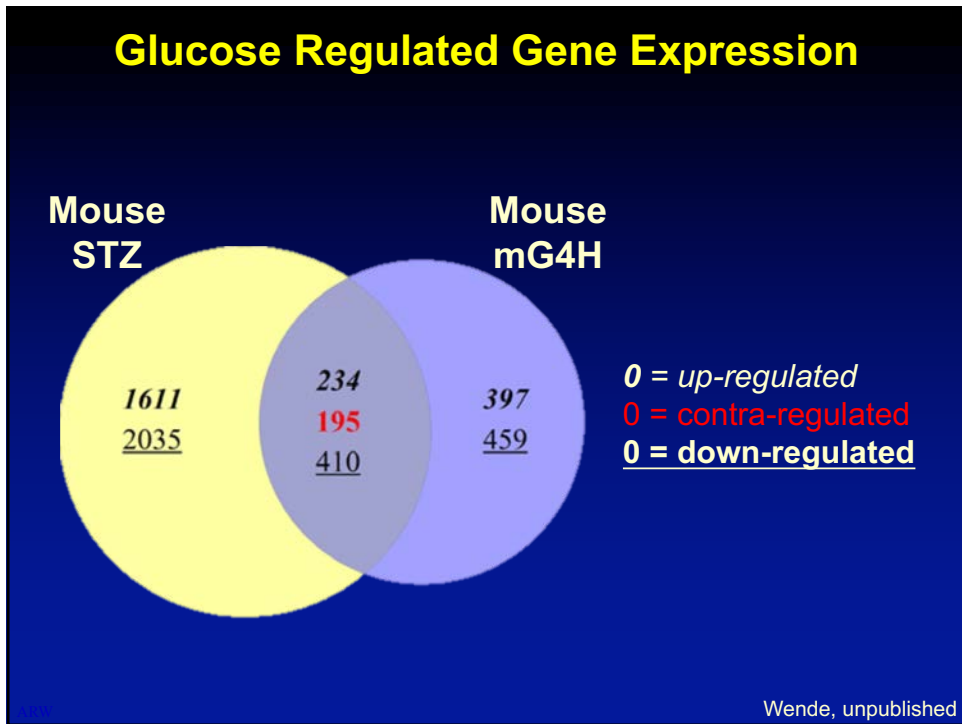
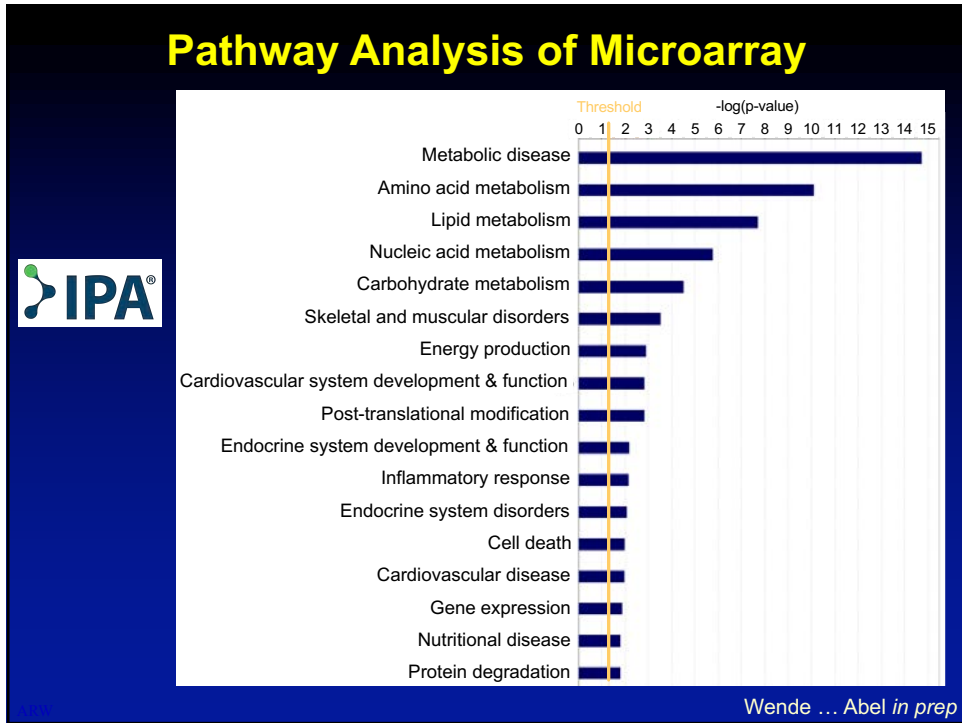
Adapted from Lewis and Abdel-Haleem 2013 *Front Physiol* 4:237

Transcriptomic Analysis Using the Agilent SurePrint G3 60K Microarray

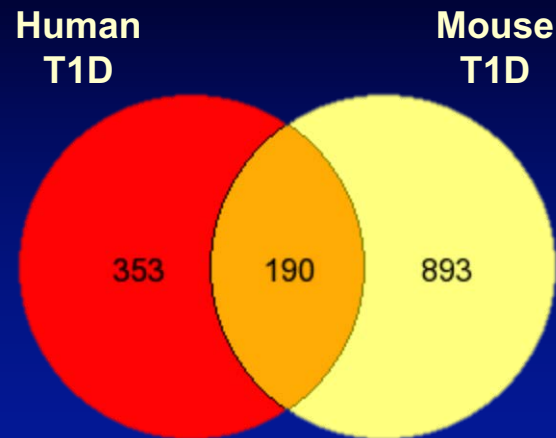


181.9 MB



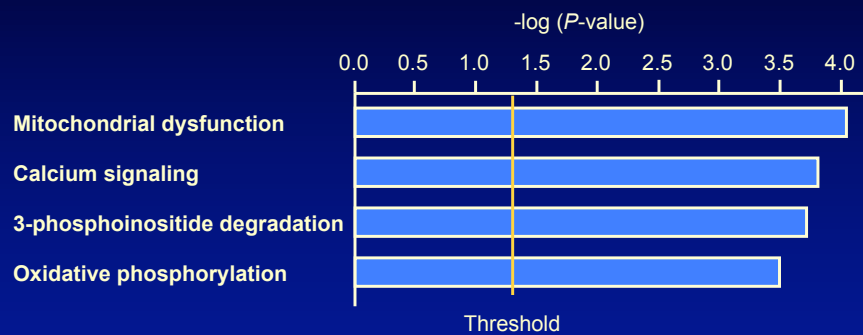


Species Conservation of Gene Expression Changes in Diabetes



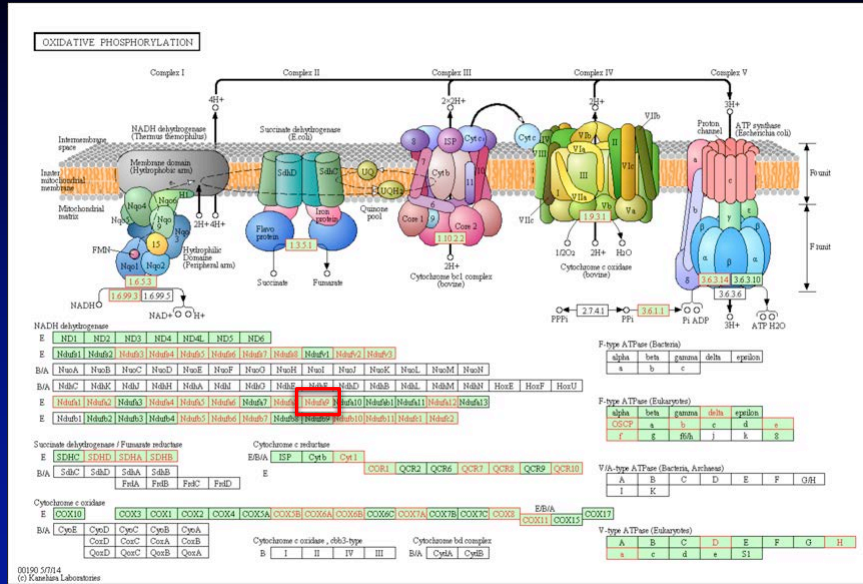
Drakos ... Wende, unpublished

Species Conservation of Gene Expression Changes in Diabetes



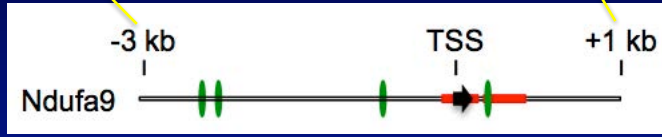
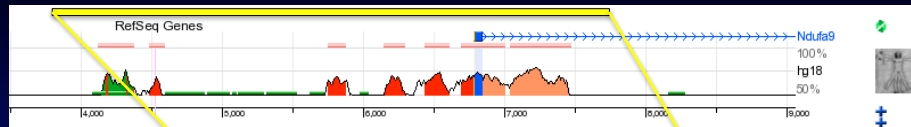
Drakos ... Wende, unpublished

Oxidative Phosphorylation



GeneSifter using KEGG

Ndufa9 Gene Promoter Structure



KEY

TSS = Transcription start site

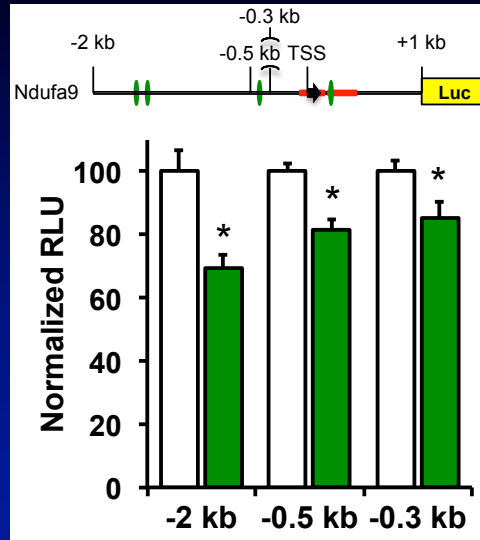
█ = CpG island

█ = Sp1 RE

<http://ecrbrowser.dcode.org>

Ndufa9 Gene Promoter Mapping

Transient Transfection Promoter Activity



Glucose

□ 5.5 mM

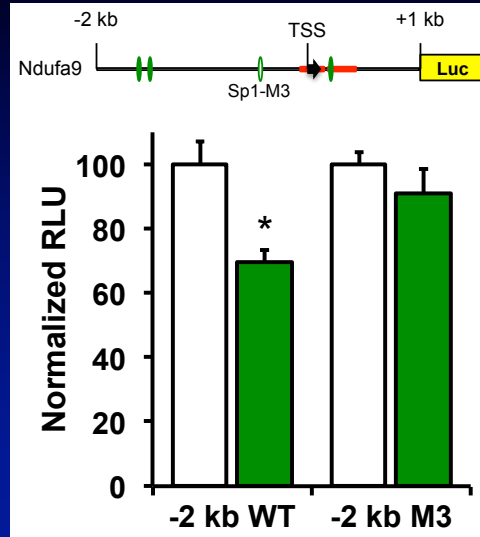
■ 25 mM

C₂C₁₂ Myotubes
n = 9
* P < 0.05

Wende ... Abel *in prep*

Ndufa9 Gene Promoter Mapping

Transient Transfection Promoter Activity



Glucose

□ 5.5 mM

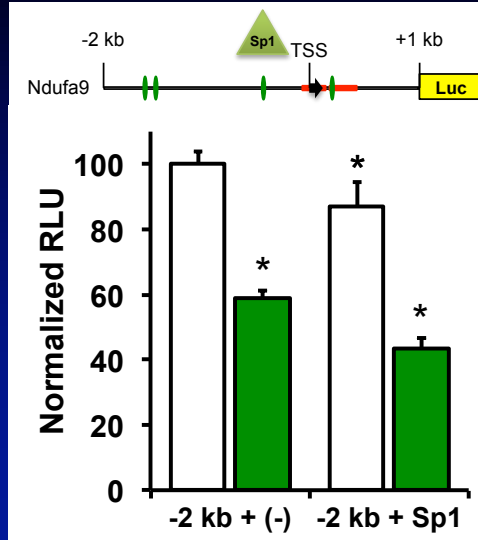
■ 25 mM

C₂C₁₂ Myotubes
n = 9
* P < 0.05

Wende ... Abel *in prep*

Ndufa9 Gene Promoter Mapping

Transient Transfection Promoter Activity



Glucose

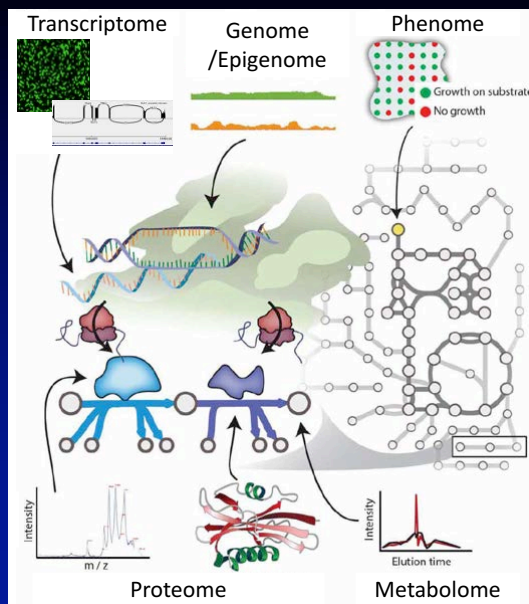
□ 5.5 mM

■ 25 mM

C₂C₁₂ Myotubes
n = 9
* P < 0.05

Wende ... Abel *in prep*

Systems Biology



Phenome

Obesity, diabetes, heart failure, BHI, etc.

Transcriptome

Northern, 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

the journal of biological chemistry
jbc
2014 THEMATIC MINIREVIEW SERIES
Nutrient Regulation of Cellular Metabolism & Physiology by O-GlcNAcylation
ASBMB AMERICAN SOCIETY FOR BIOCHEMISTRY AND MOLECULAR BIOLOGY

O-GlcNAcylation

Research Topic

30 years old: O-GlcNAc reaches age of reason - Regulation of cell signaling and metabolism by O-GlcNAcylation.

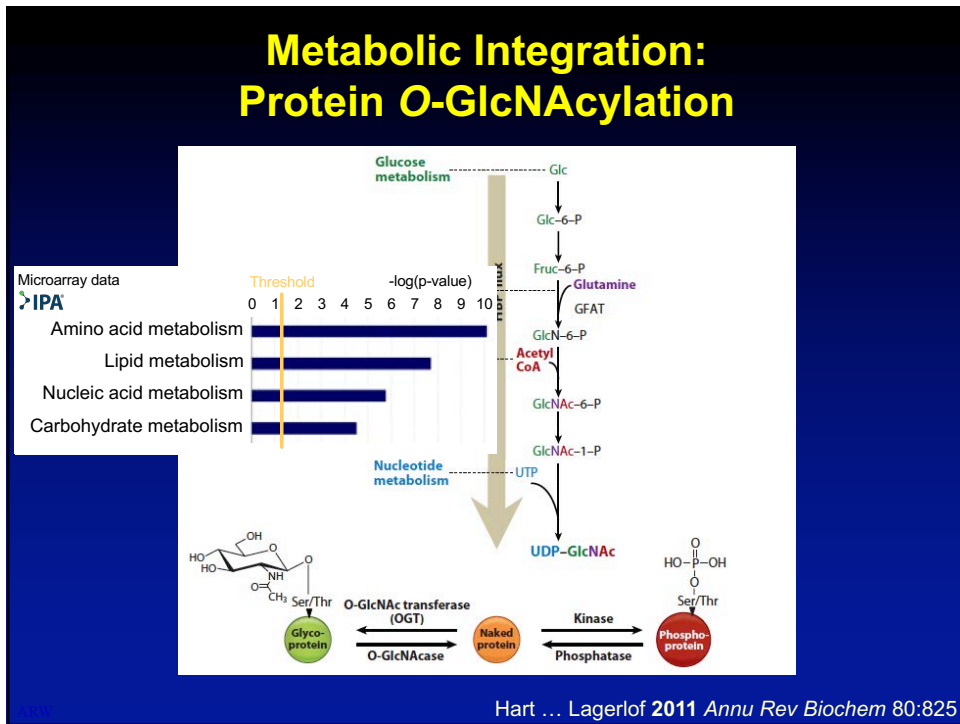
Download Ebook PDF Download Ebook EPUB

Like: 1 Comment: 0 Share: 0

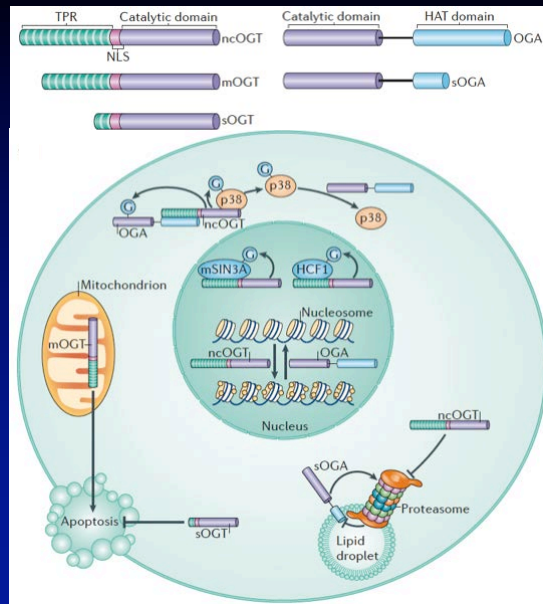
f 23 t 4 g+ 2 in 3 < 65

Overview **13** Articles **63** Authors Impact Comments

35,029 VIEWS

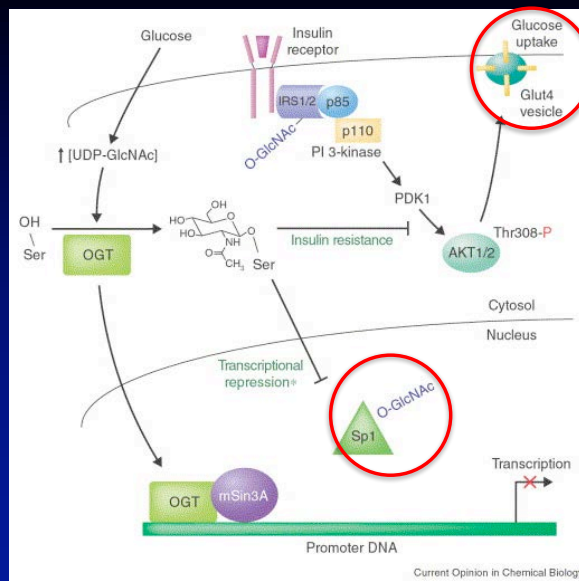


O-GlcNAc Cycling



Hanover ... Love 2012 *Nat Rev Mol Cell Biol* 13(5):312

GlcNAc Regulation of Sp1

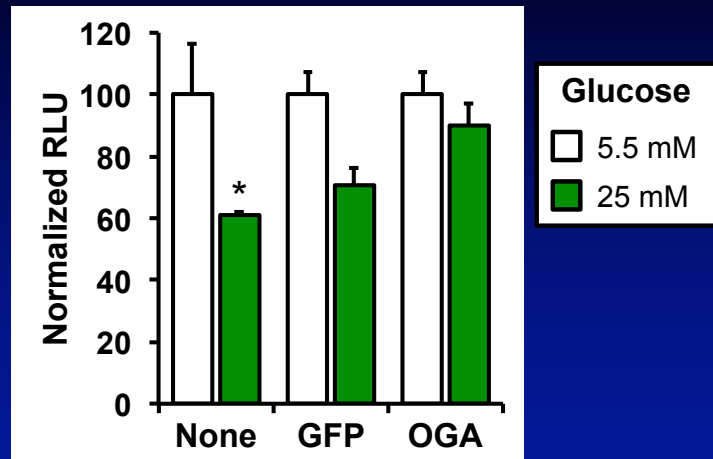


Current Opinion in Chemical Biology

Vosseller ... Hart 2002 *Curr Opin Chem Biol* 6(6):851

GlcNAcylation Regulates *Ndufa9* Gene Expression

Transient
Transfection
Promoter
Activity



C₂C₁₂ Myotubes
n = 3
* *P* < 0.05

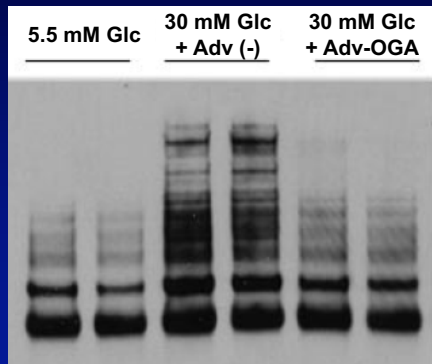
Li Wang
Wende ... Abel *in prep*

Conclusion – Part 2

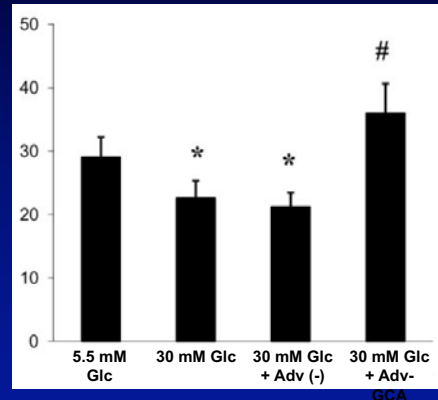
Enhanced glucose delivery regulates oxidative capacity via transcriptional mechanisms including GlcNAcylation of transcription factors.

Mitochondrial Protein O-GlcNAcylation and Neonatal Cardiomyocyte Metabolic Function

Mitochondrial Protein
O-GlcNAcylation



Complex I Activity

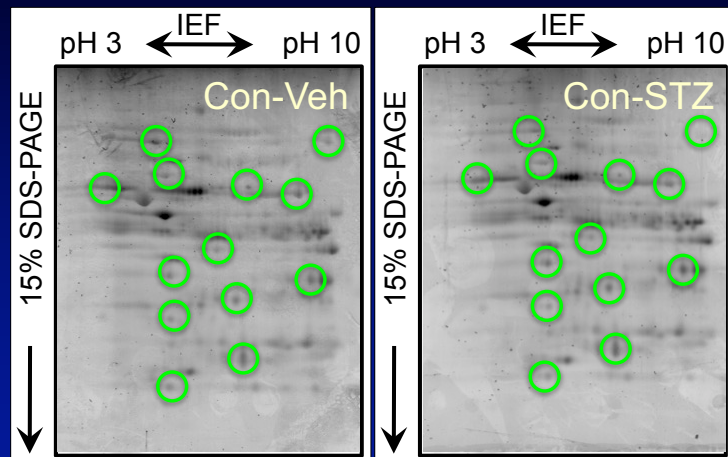


O-GlcNAcylation of NDUFA9

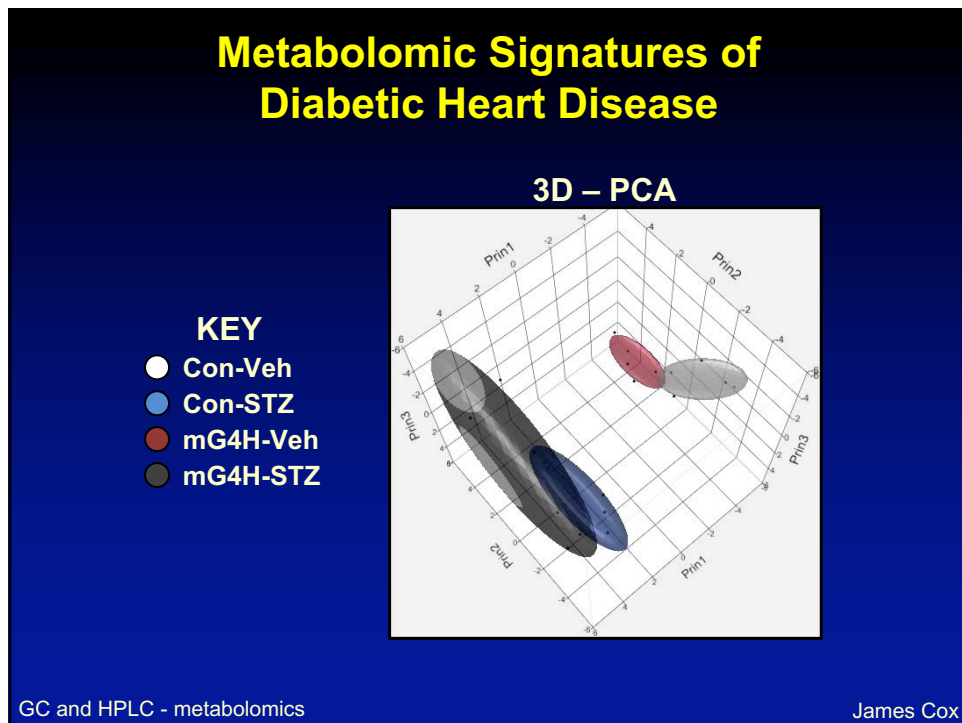
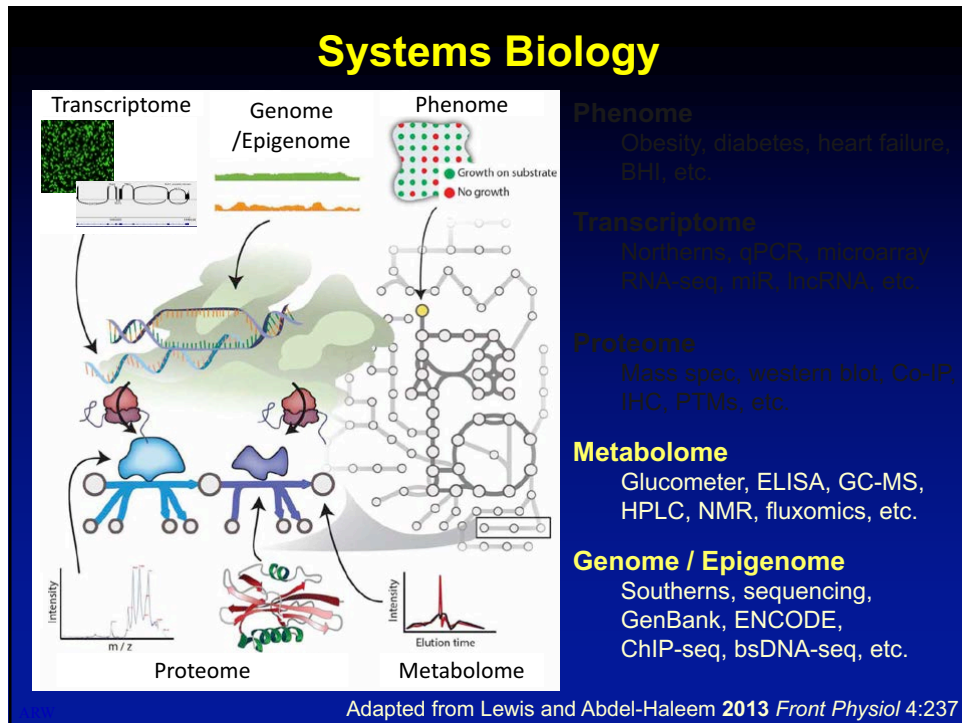
Hu ... Dillmann 2009 *J Biol Chem* 284(1):547

GLUT4 Induction Alters the Cardiac Mitochondrial Glycoproteome

Isolated
Mitochondria
2D-PAGE
Pro-Q
Emerald



Hansjörg Schwertz
Wende, unpublished



Studies on Myocardial Metabolism*

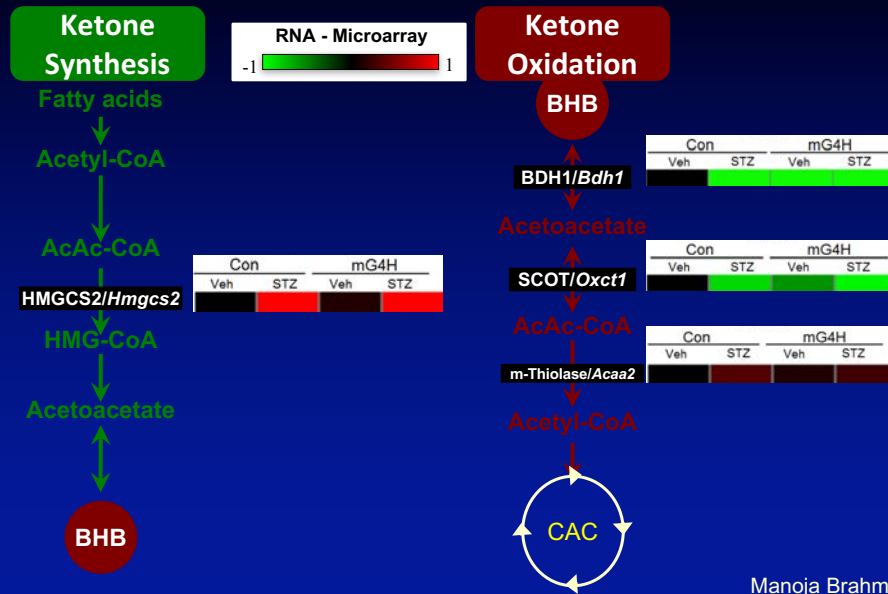
IV. Myocardial Metabolism in Diabetes

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

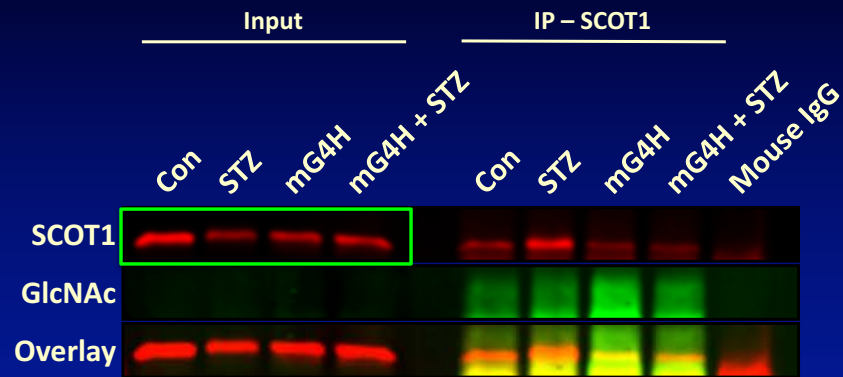
lactate usage and a slight decline in that of pyruvate. There is no change in utilization of amino acids by the heart in both species. Myocardial glucose consumption is reduced in dog and man relative to the elevation in blood glucose concentration. The myocardial usage of ketones is slightly increased 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 of diabetes is not clear. Anesthesia, which was used in the dogs, may have played some part.

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

GLUT4 Induction Alters Cardiac Ketone Utilization Genes



GLUT4 Induction Alters Cardiac Ketone Protein GlcNAcylation

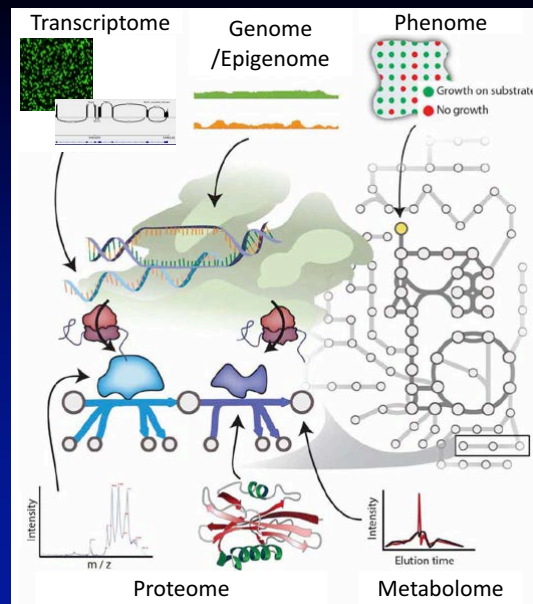


Manoja Brahma

Conclusion – Part 3

Enhanced cardiac glucose delivery alters metabolic flux through other pathways and regulates the mitochondrial proteome via O-GlcNAcylation.

Systems Biology



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

From Human to Mouse and Back Again



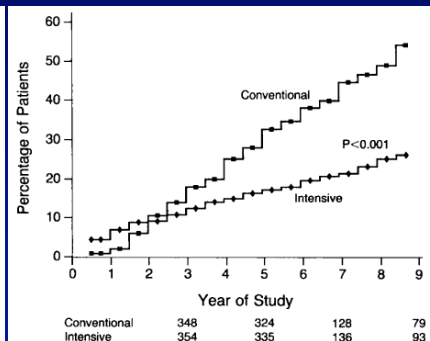
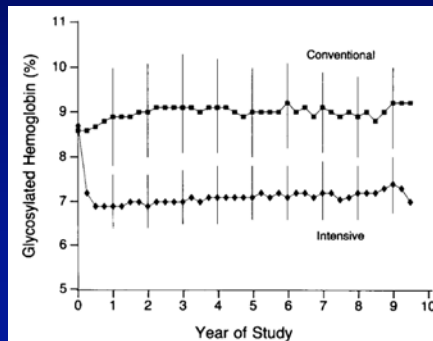
Broad Institute Communications

Role of Epigenetics in Gene Expression



Epigenetics - Programming DCCT: Diabetes Control and Complications Trial

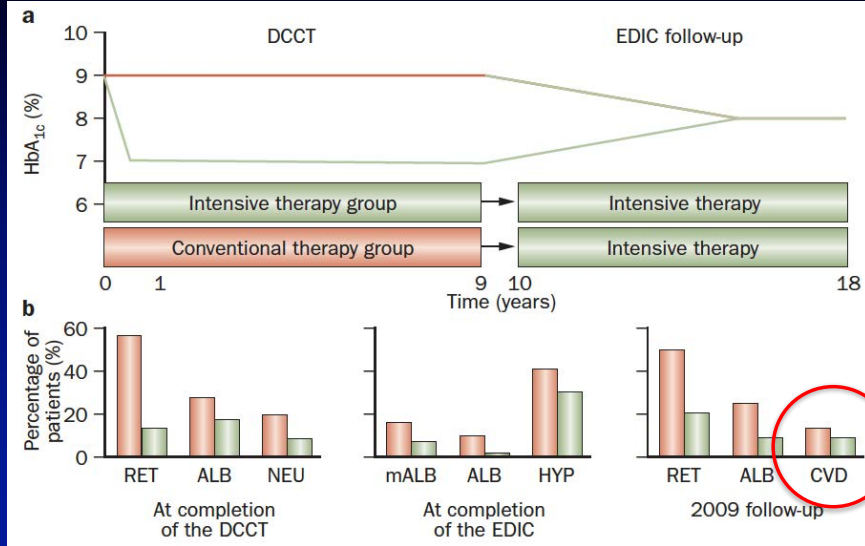
The New England
Journal of Medicine
©Copyright, 1993, by the Massachusetts Medical Society
Volume 329 SEPTEMBER 30, 1993 Number 14
THE EFFECT OF INTENSIVE TREATMENT OF DIABETES ON THE DEVELOPMENT AND PROGRESSION OF LONG-TERM COMPLICATIONS IN INSULIN-DEPENDENT DIABETES MELLITUS
THE DIABETES CONTROL AND COMPLICATIONS TRIAL RESEARCH GROUP*



Conventional	348	324	128	79
Intensive	354	335	136	93

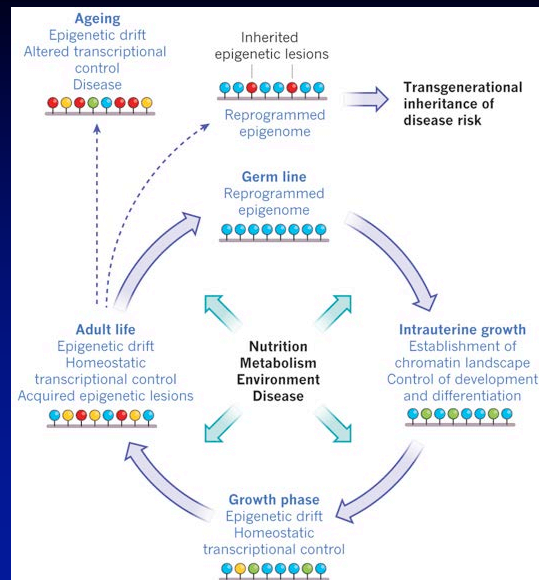
Epigenetics - Memory

EDIC: Epidemiology of Diabetes Interventions Trial



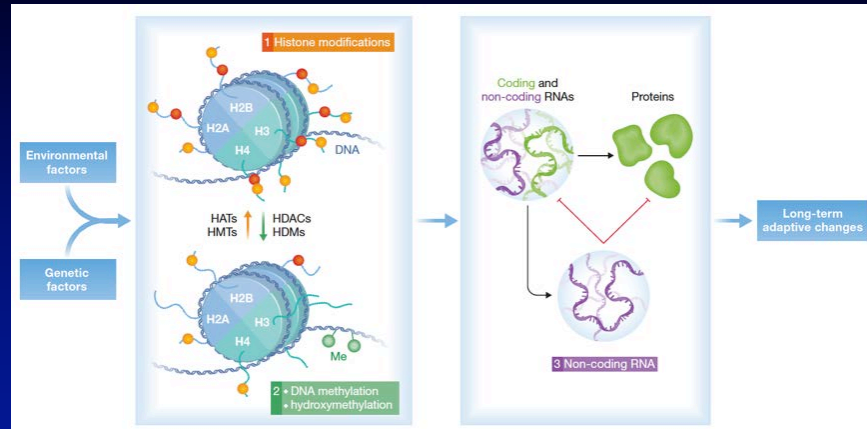
Pirola ... El-Osta 2010 *Nat Rev Endocrinol* 6(12):665

Epigenetics: Transgenerational and Drift



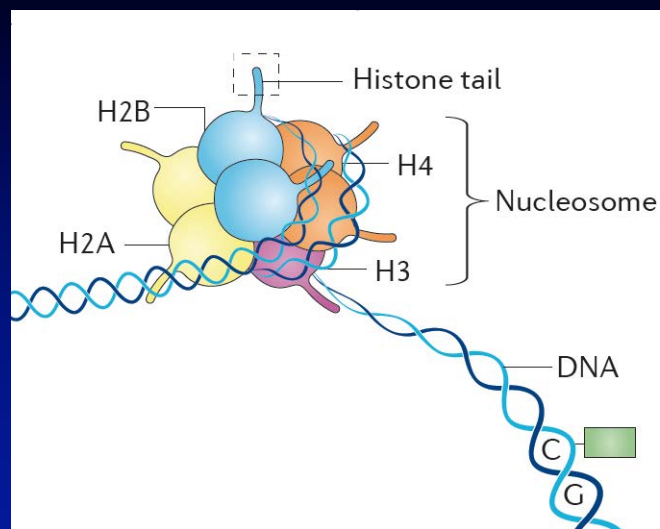
Gut and Verdin 2013 *Nature* 502:489

Epigenetic Code



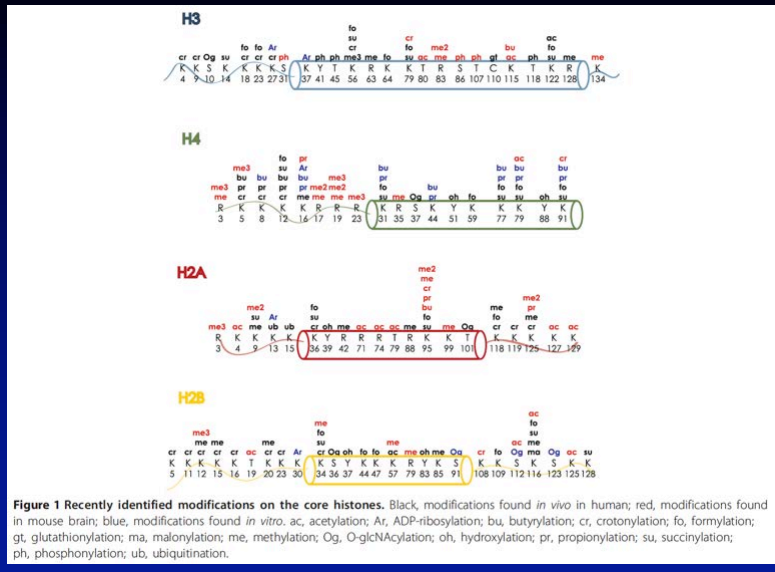
Fischer 2014 *EMBO J* 33(9):945-489

Chromatin Regulation



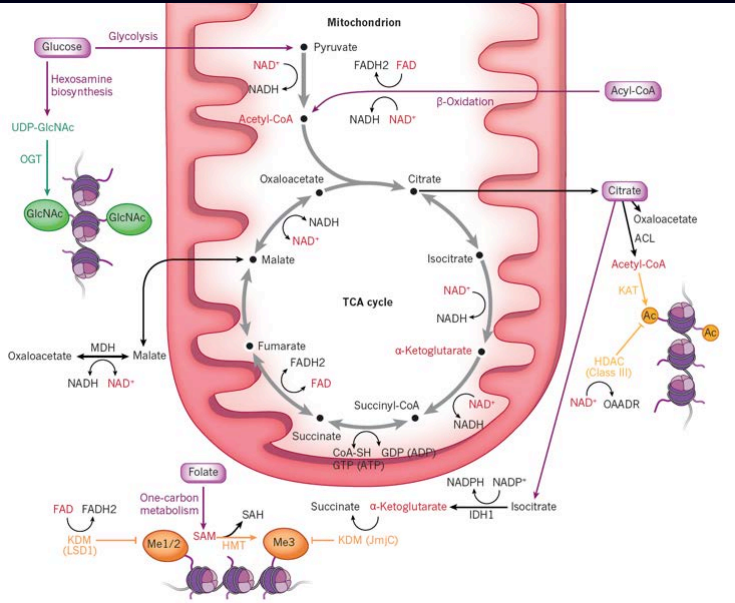
Gräff and Tsai 2013 *Nat Rev Neurosci* 14(2):97

How do metabolites fit in?



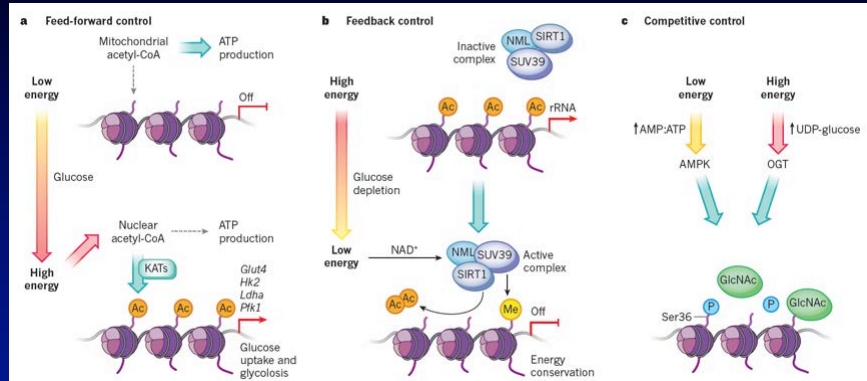
Arnaudo ... Garcia 2013 Epigenetics Chromatin 6(1):24

Metabolite Signaling to Chromatin



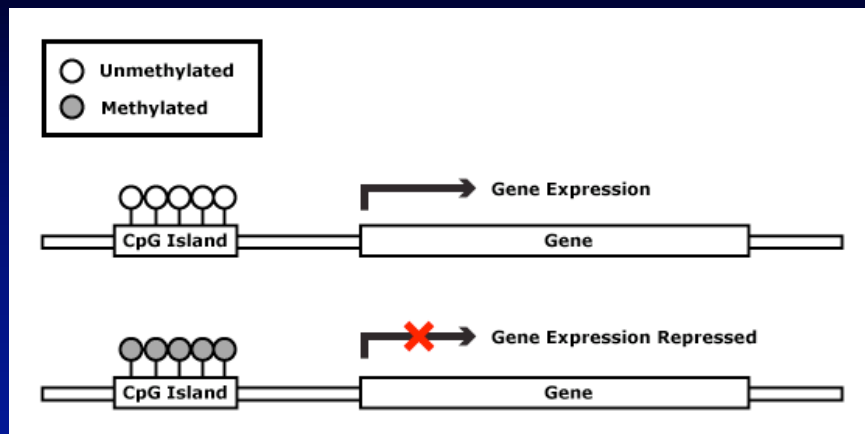
Gut and Verdin 2013 Nature 502:489

How does GlcNAc fit in?



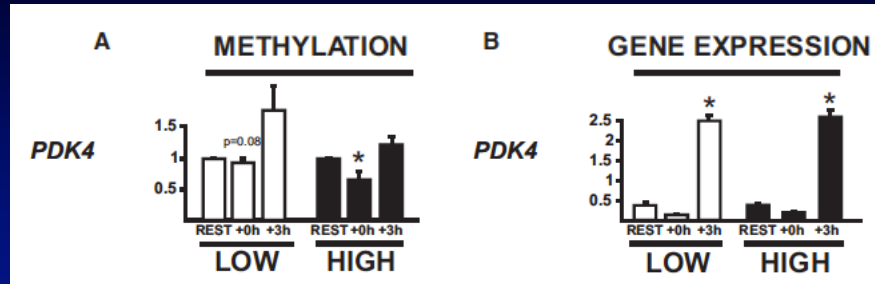
Gut and Verdin 2013 *Nature* 502(7472):489

DNA Methylation 101



ucsf.edu

Exercise Alters DNA Methylation of Key Metabolic Genes



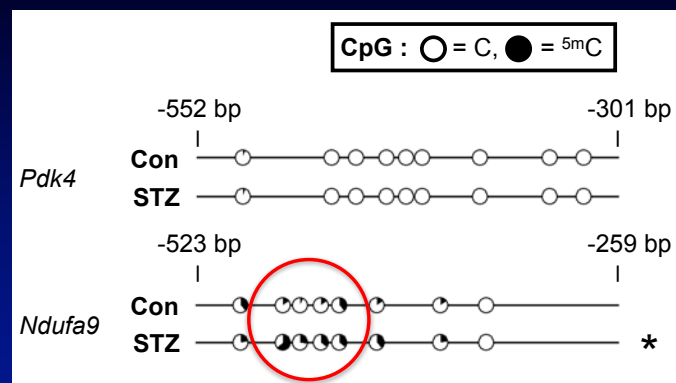
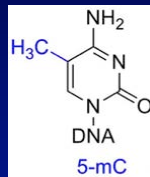
Low = 40% VO_{2peak} High = 80% VO_{2peak}

Subjects fasted overnight and then consumed a high carbohydrate diet 4 hr prior to exercise.

Barres and Zierath 2012 *Cell Metab* 15:405

Diabetes Regulated Cardiac DNA Methylation

Targeted
bsDNA-seq
5-mCpG

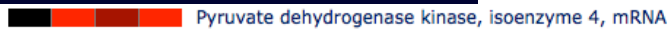


Heart, LV
 $n = 10$
* $P < 0.05$

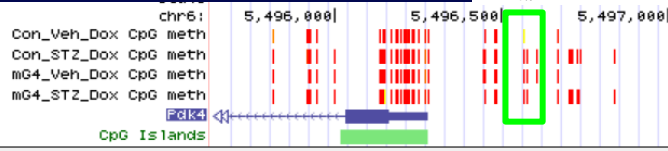
Wende, unpublished

Methylation and Expression

RNA – microarray



Methylation – genome sequencing



Protein – western blot



GeneSifter and Zymo/UCSC Genome Browser

Other Human/Mouse Comparisons



Genetics Of Lipid Lowering Drugs And Diet Network

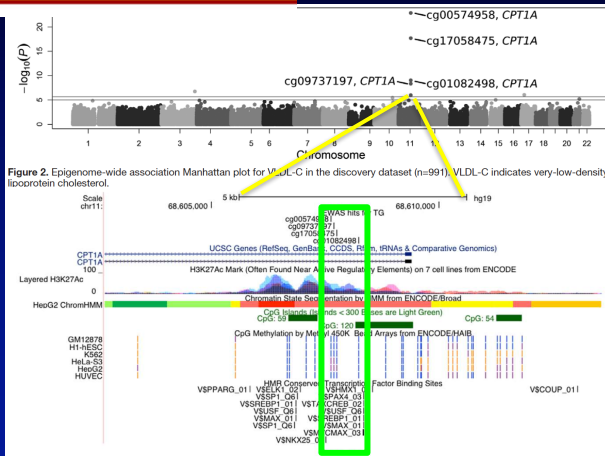
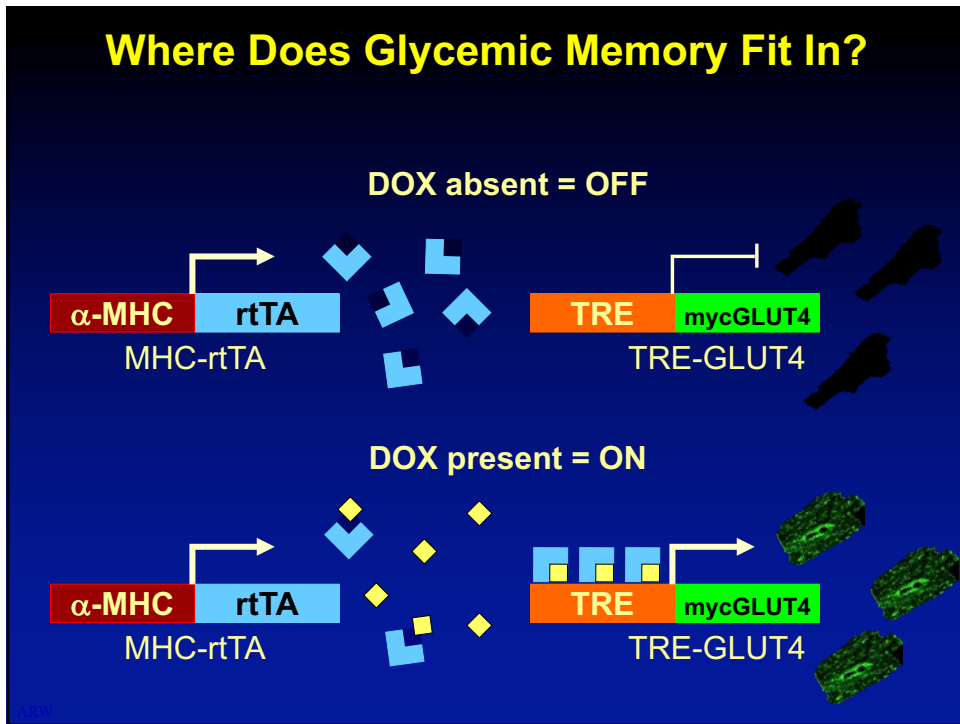
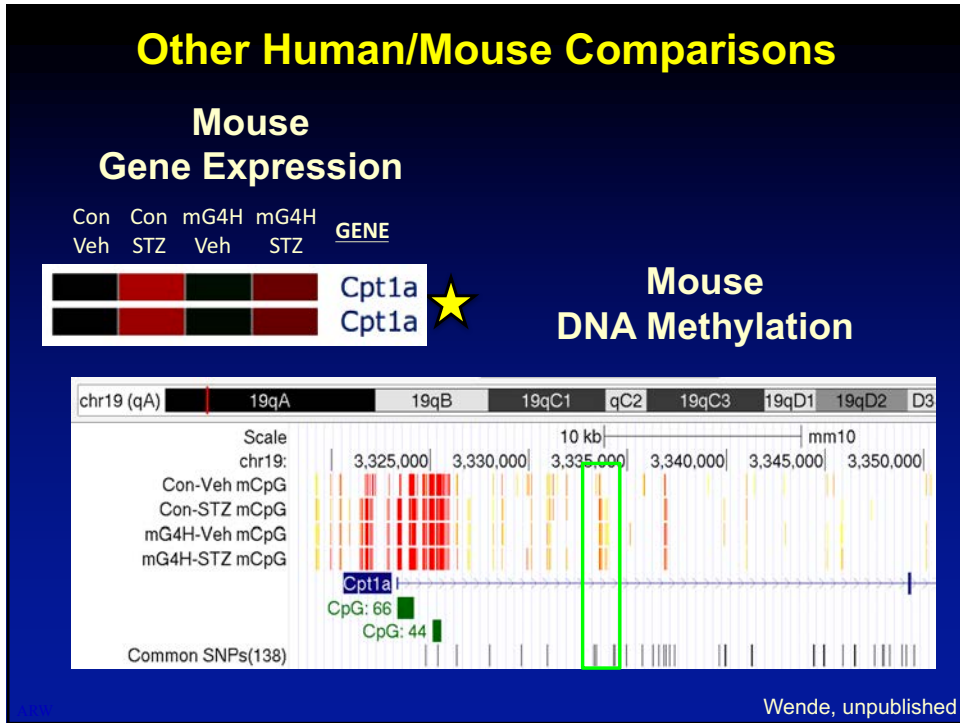


Figure 2. Epigenome-wide association Manhattan plot for VLDL-C in the discovery dataset (n=991). VLDL-C indicates very-low-density lipoprotein cholesterol.

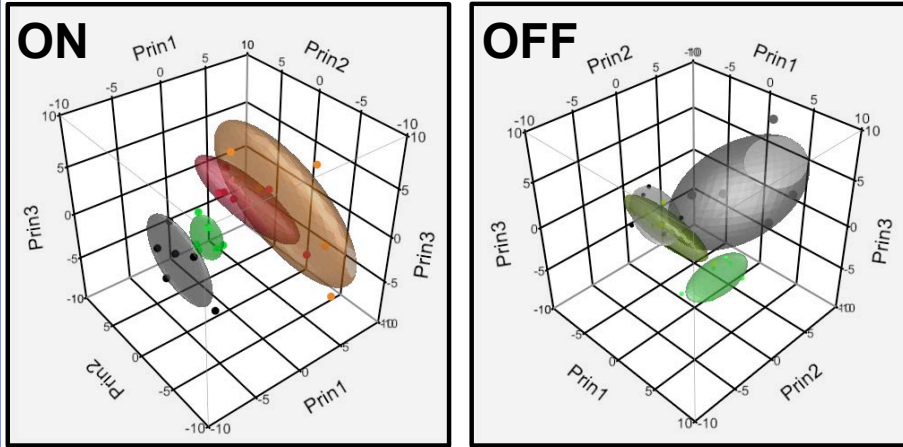
Figure 3. ENCODE annotation of the promoter region and intron 1 of *CPT1A*. Top CpGs for TG are positioned within the gene along with CpG islands, cell line chromatin state (ChromHMM), cell line methylation at CpG sites on the Methy450 Beadchip according to Hudson Alpha Institute for Biotechnology (HAIB; note blue, purple, and orange highlights correspond to low, medium and high methylation state, respectively), and HMR conserved transcription factor binding sites. CpG indicates cytosine-(phosphate)-guanine; and TG, triglyceride.

Irvin ... Arnett 2014 *Circulation* 130:565



Metabolomics

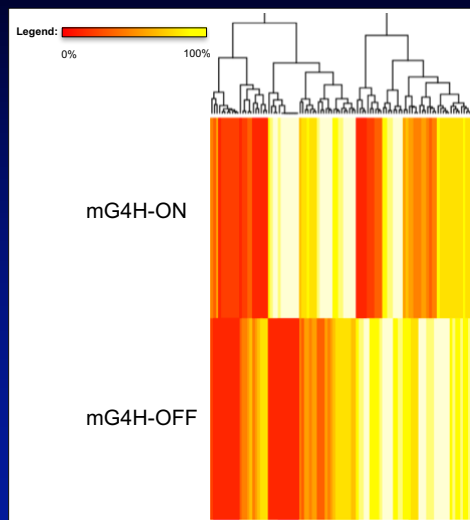
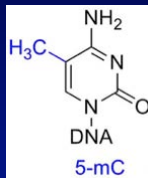
Con-Veh
 Con-STZ
 mG4H-STZ
 mG4H-Veh
 mG4H-1wk
 mG4H-2wk



Wende, unpublished

Glucose Cycling Alters Epigenetic Programming

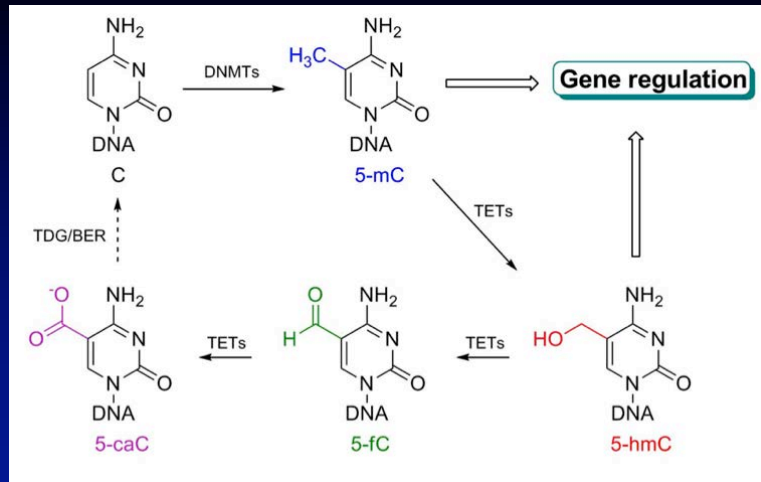
**Genomewide
 bsDNA-seq
 5-mCpG**



Heart, LV

Zymo Research
 Wende, unpublished

Background



5-hmC

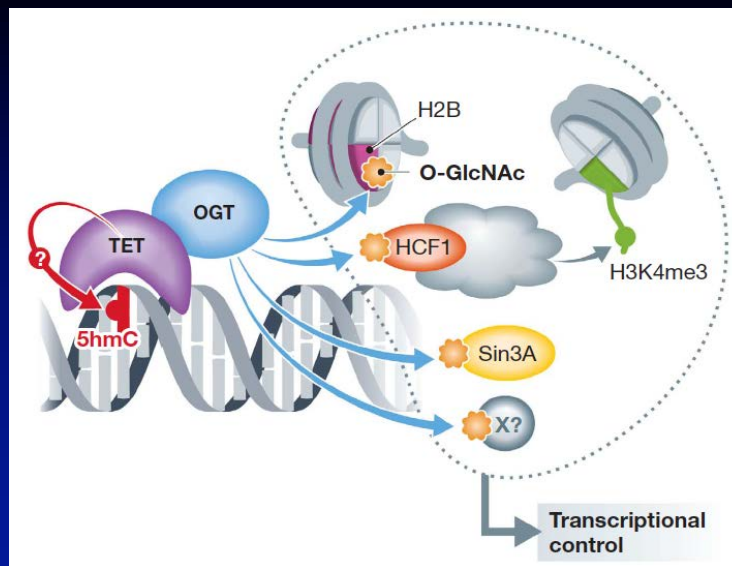
Wyatt and Cohen **1952** *Nature* 170(4338):1072

Kriaucioni and Heintz **2009** *Science* 324(5929):929

Tahiliani ... Rao **2009** *Science* 324(5929):930

<http://chemistry.uchicago.edu/faculty/faculty/person/member/chuan-he.html>

How does GlcNAc fit in?



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

Conclusion – Part 4

Cellular glucose fluctuations regulates the epigenome via histone modifications and controlling the machinery for DNA methylation.

Sugar Gumming Up the Works



giphy.com

Overall Summary

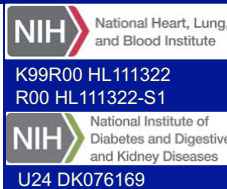
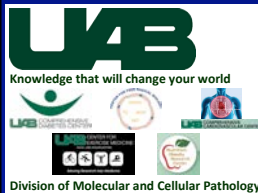
Using combined methylomics, transcriptomics, proteomics, and metabolomics we have begun to define the mechanism of glucotoxicity.

Acknowledgements

Wende Lab



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



UAB Collaborators

Steve Barnes
 John C. Chatham
 David K. Crossman
 Steve M. Pogwizd
 Martin E. Young

E. Dale Abel

John C. Schell
 Joseph Tuinei
 many others...

Stavros G. Drakos

Nikos A. Diakos

Hansjörg Schwertz

Oleh Khalimonchuk – UNL

Other Colleagues & Mentors past and present

