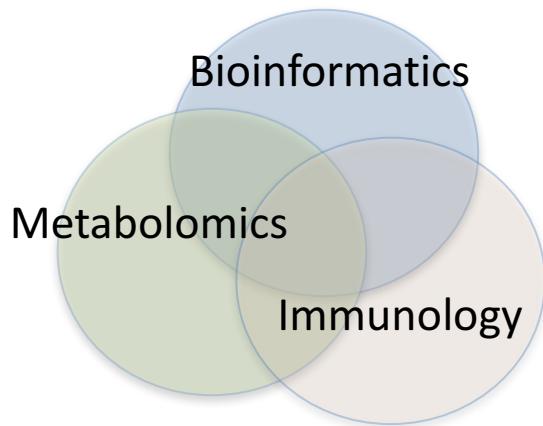


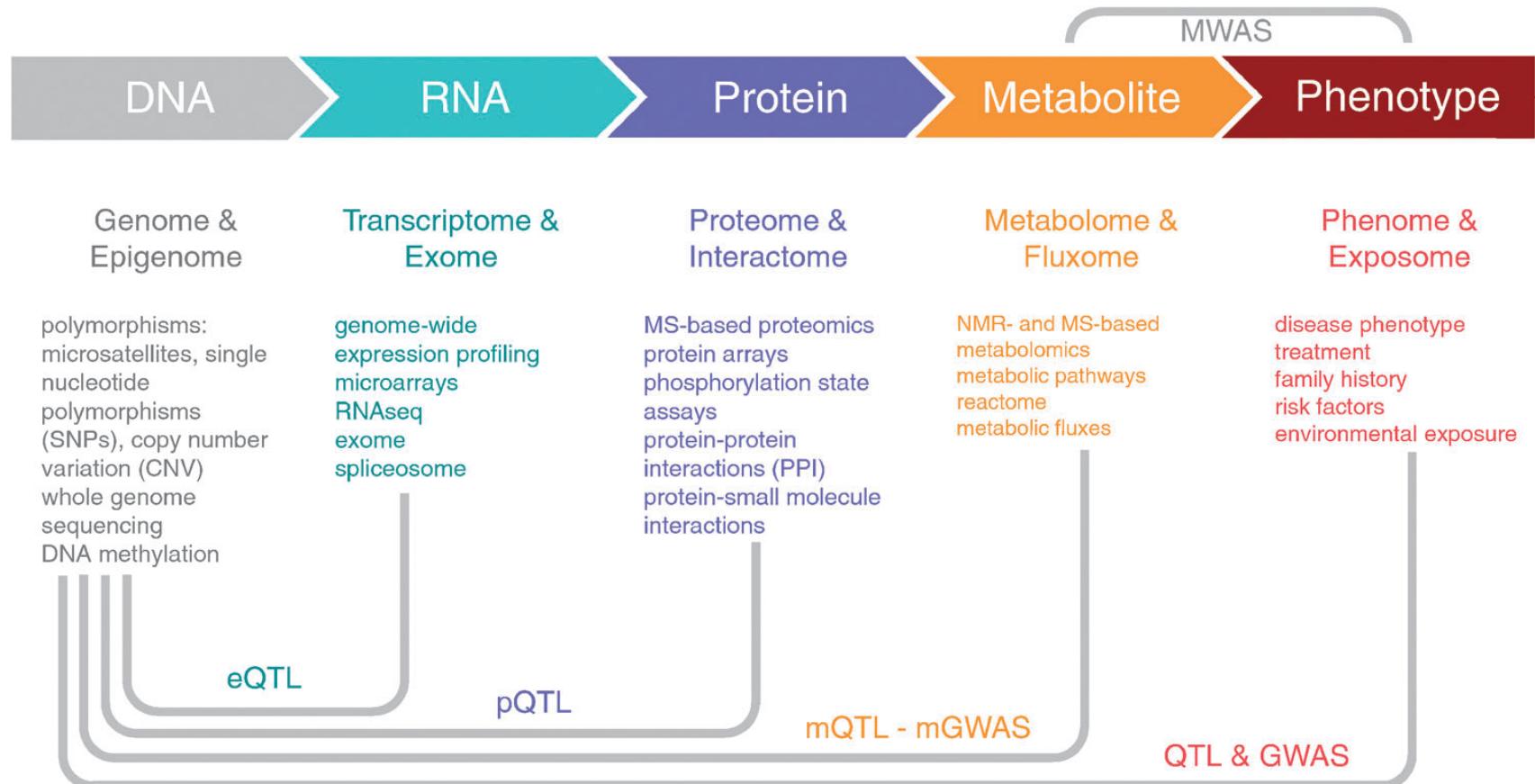


# Overview of Metabolomics Workflow



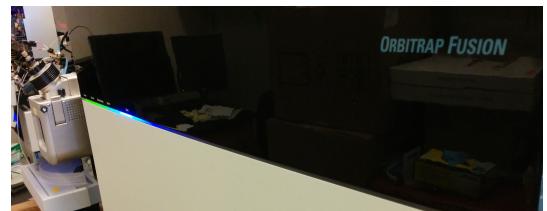
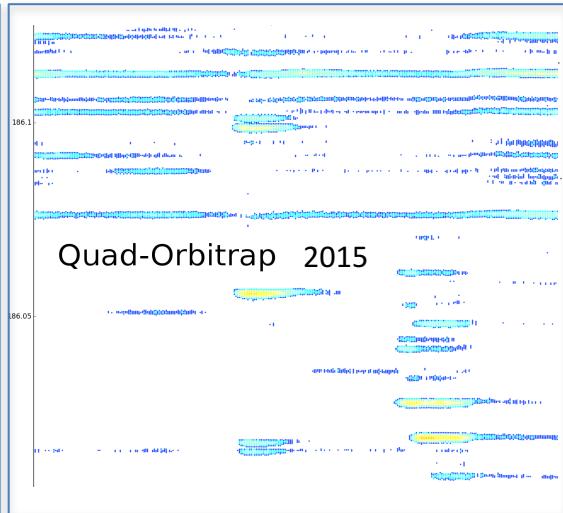
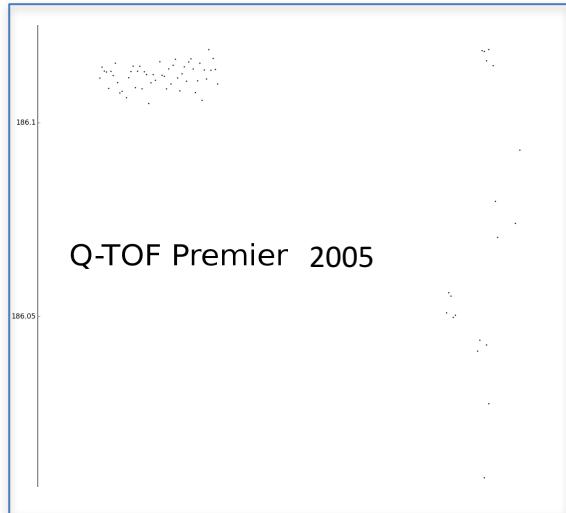
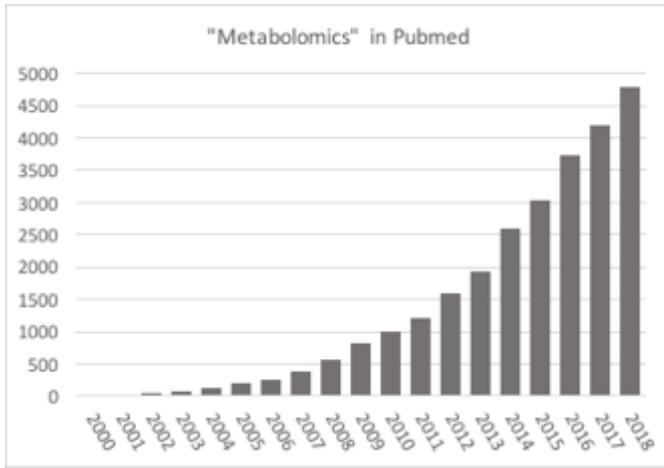
Shuzhao Li, Ph.D  
Assistant Professor  
Department of Medicine  
Emory University  
August 26, 2019

# Biomedical -omics data

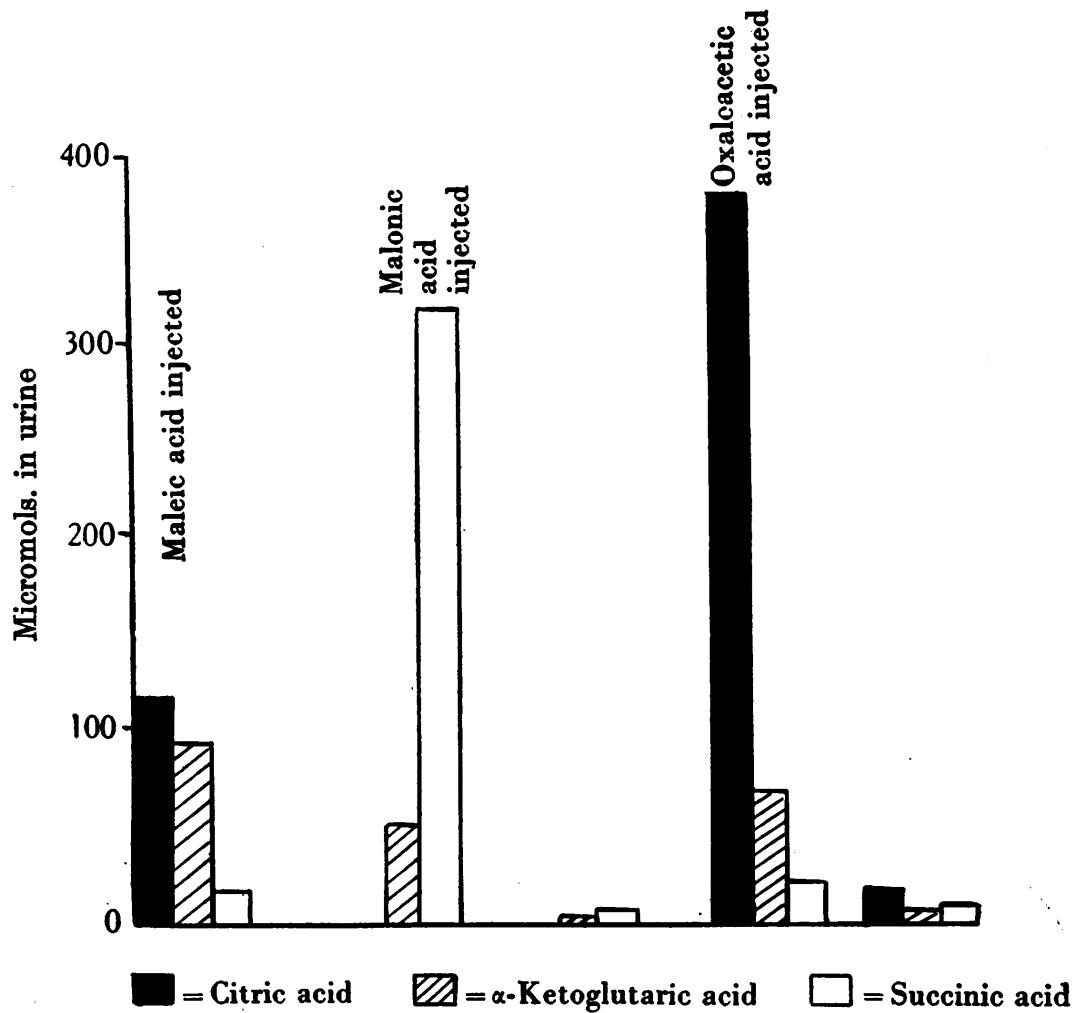
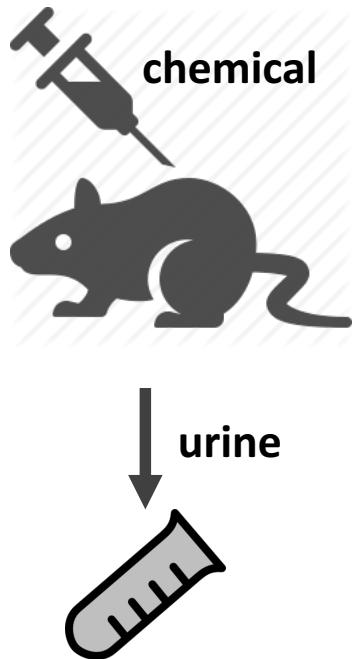


# Advance of Metabolomics

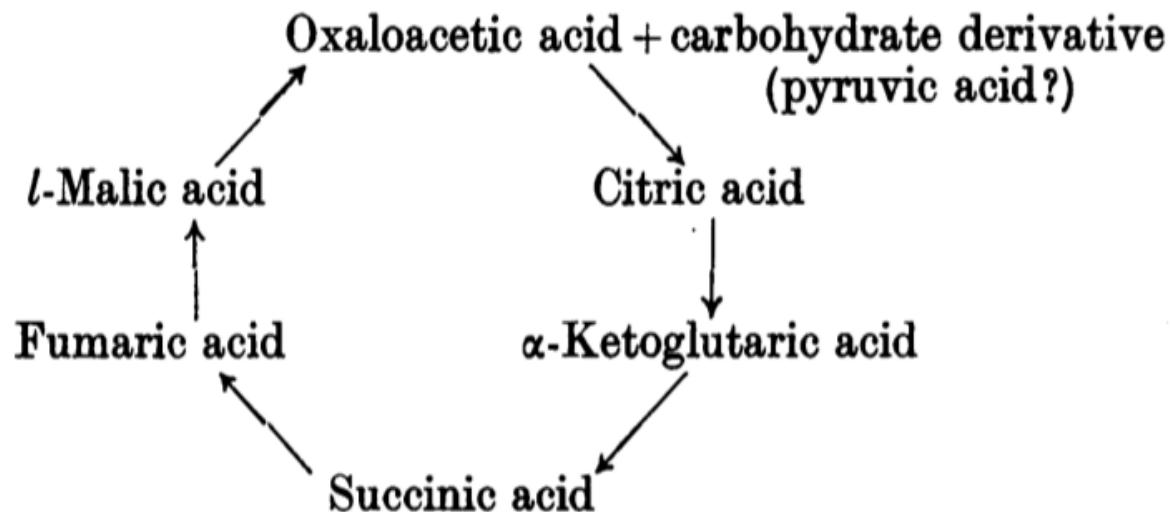
Number of papers



# Krebs et al, 1937



# Krebs Cycle, 1937



Krebs et al. 1938. Biochem Journal. 32:113

# Beadle 1941

## Genetic Control of Biochemical Reactions in *Neurospora*

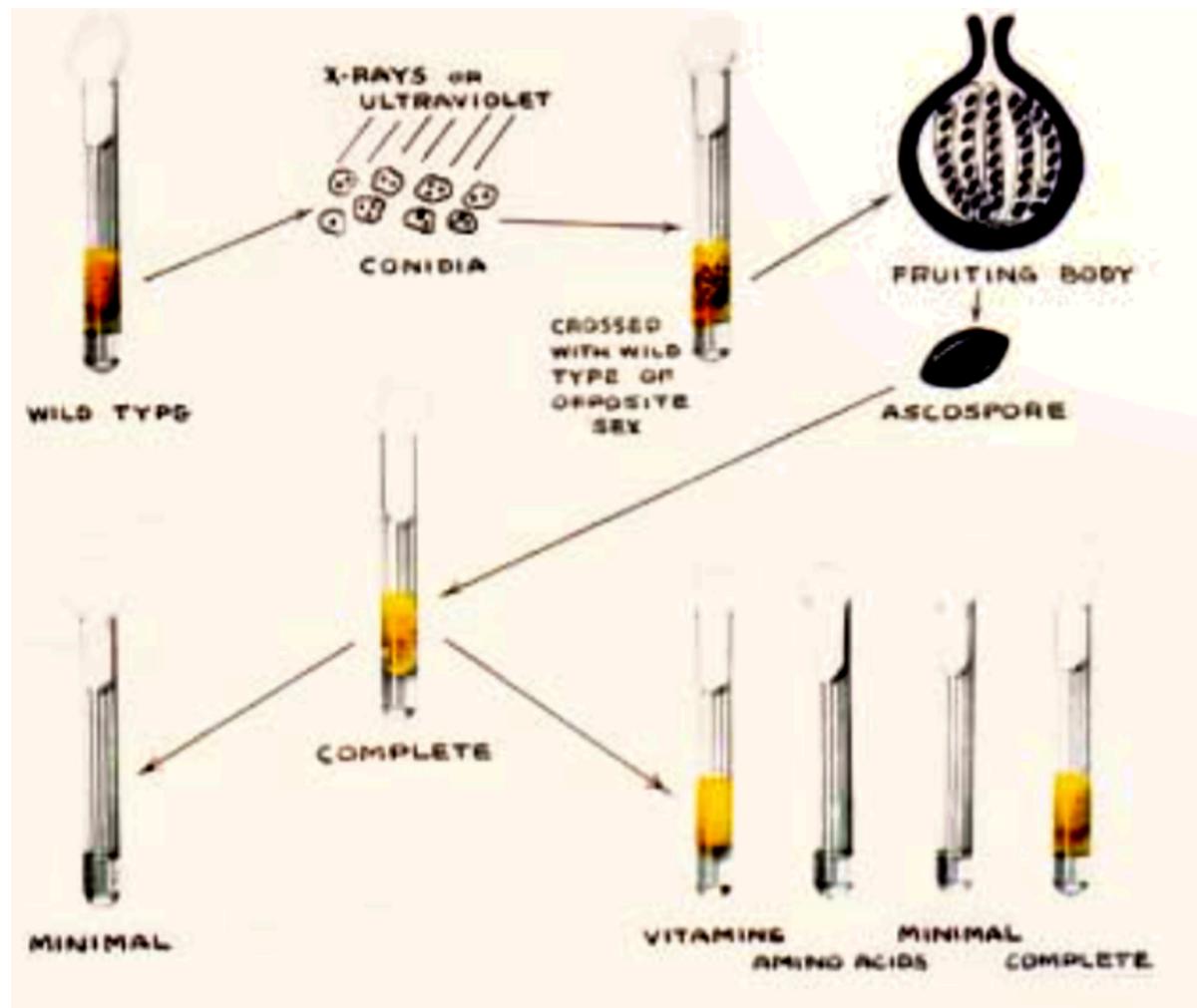


Figure from *Nature Reviews Genetics* 5, 949-954, 2004

# Related Nobel Prizes

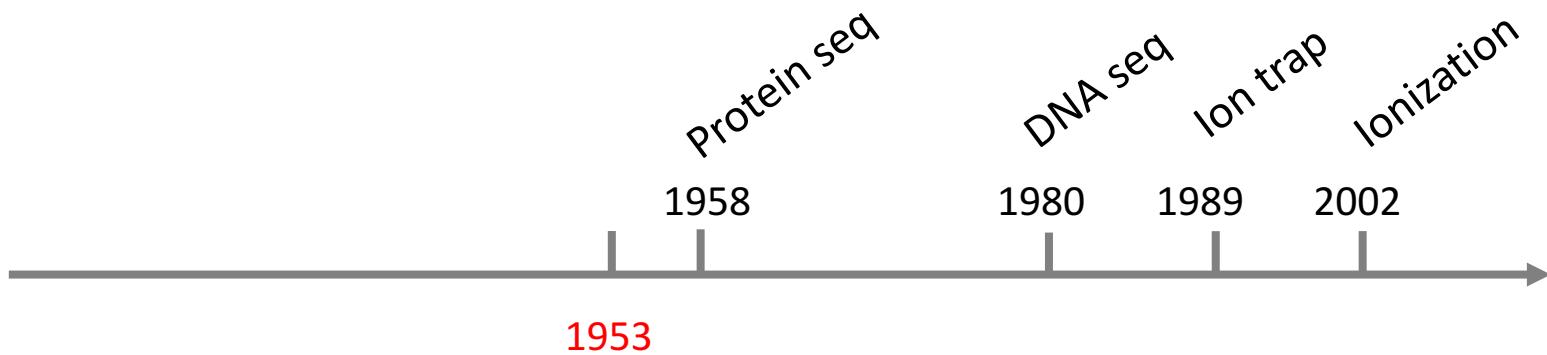
Genetic Control of Biochemical Reactions

1958

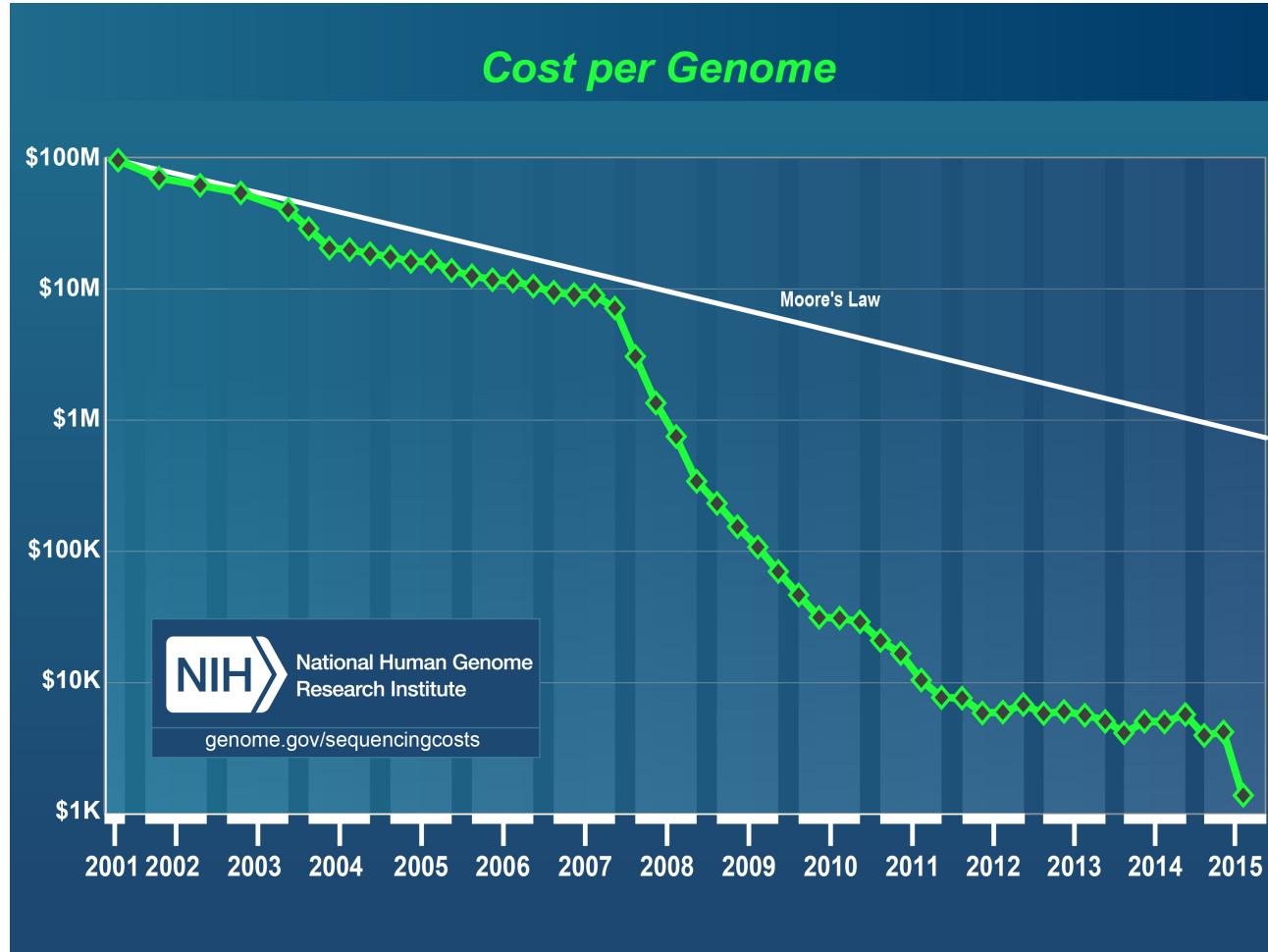
1953

Krebs Cycle

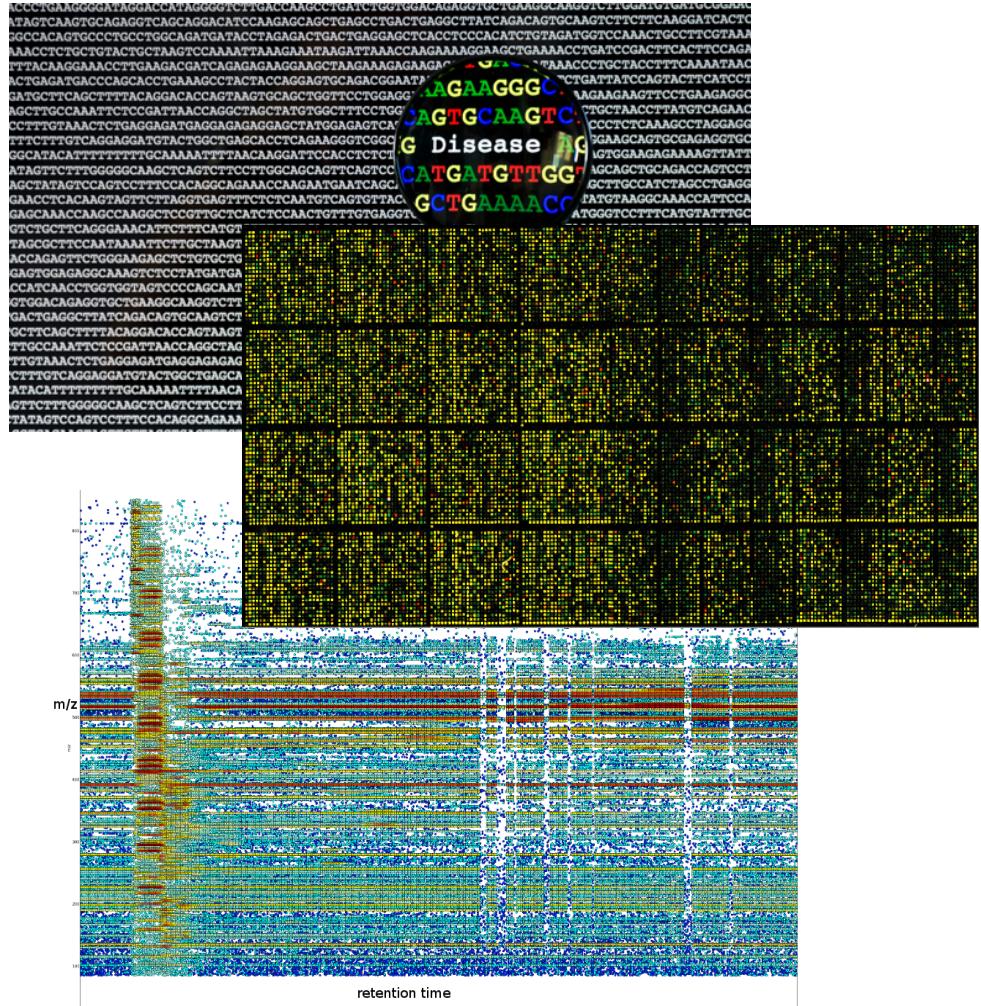
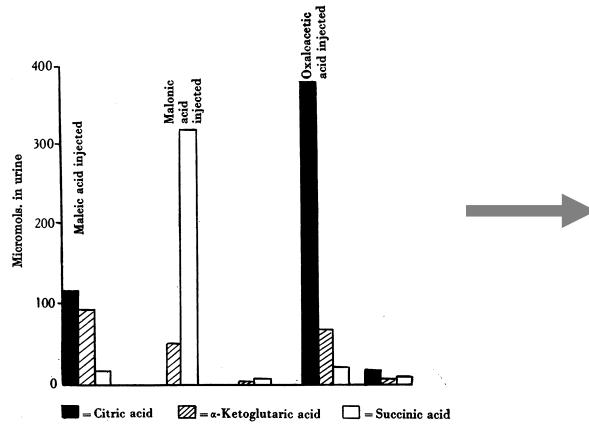
# Related Nobel Prizes



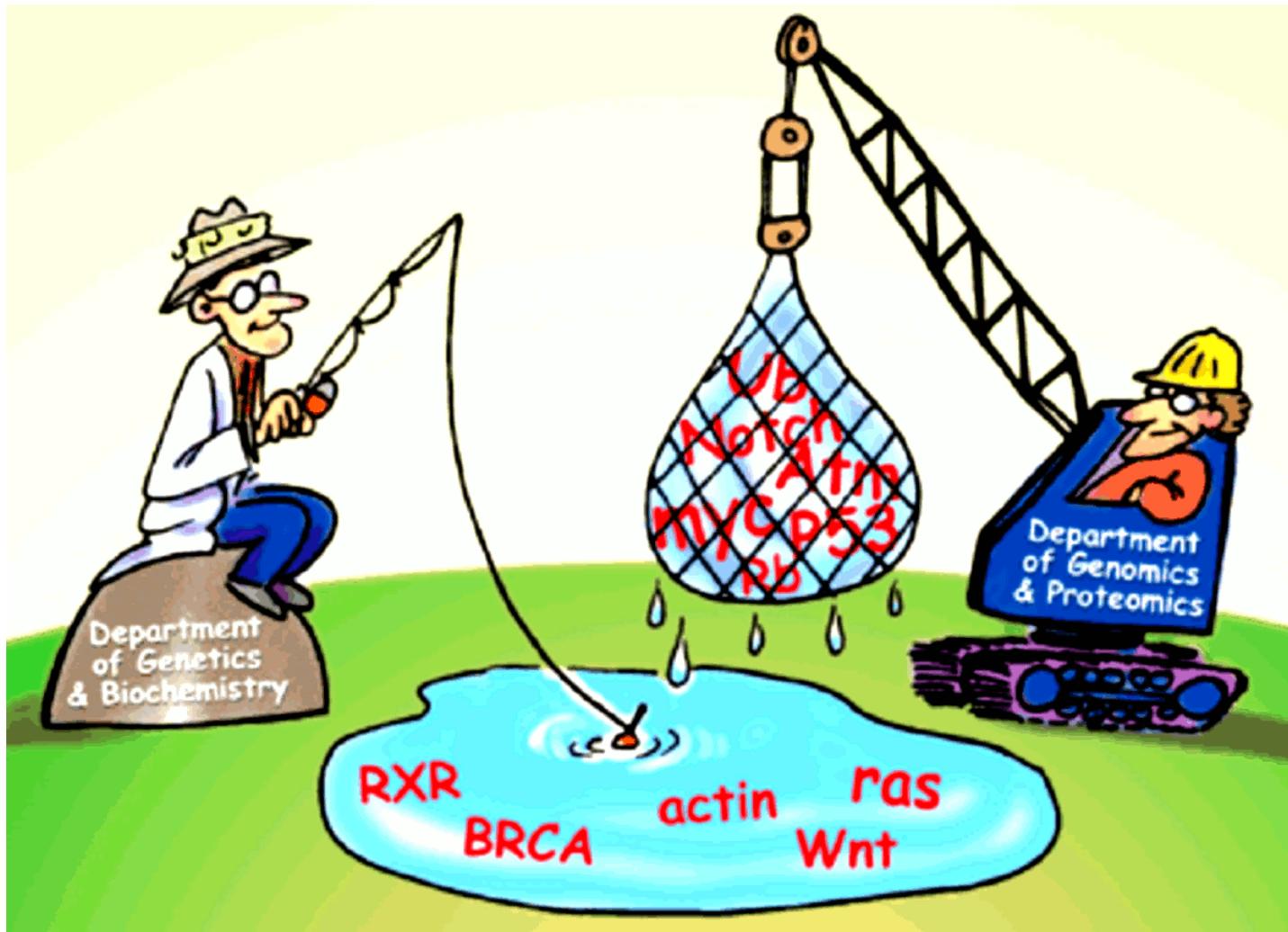
# Economy of data generation (genomics)



# The data deluge



# When science goes -omics

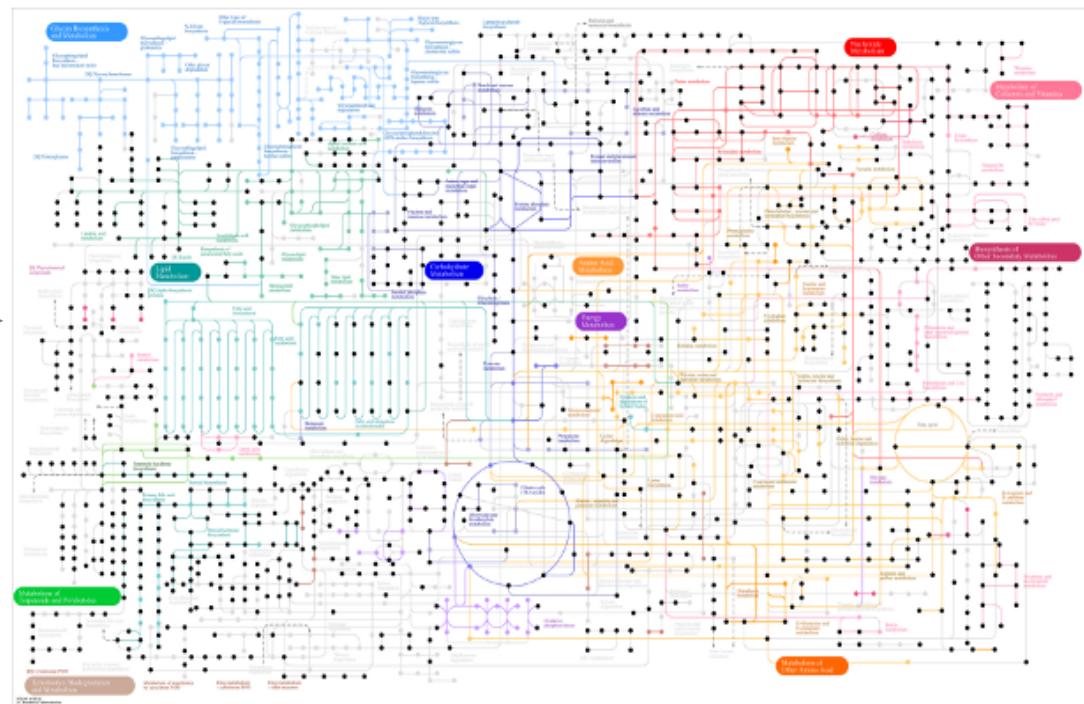
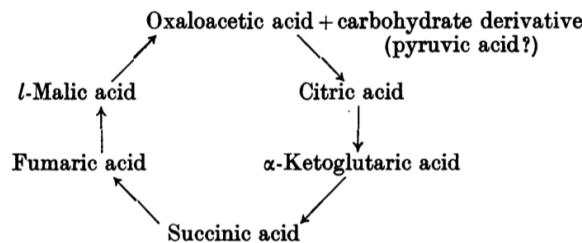


Science (2001) 291:1221

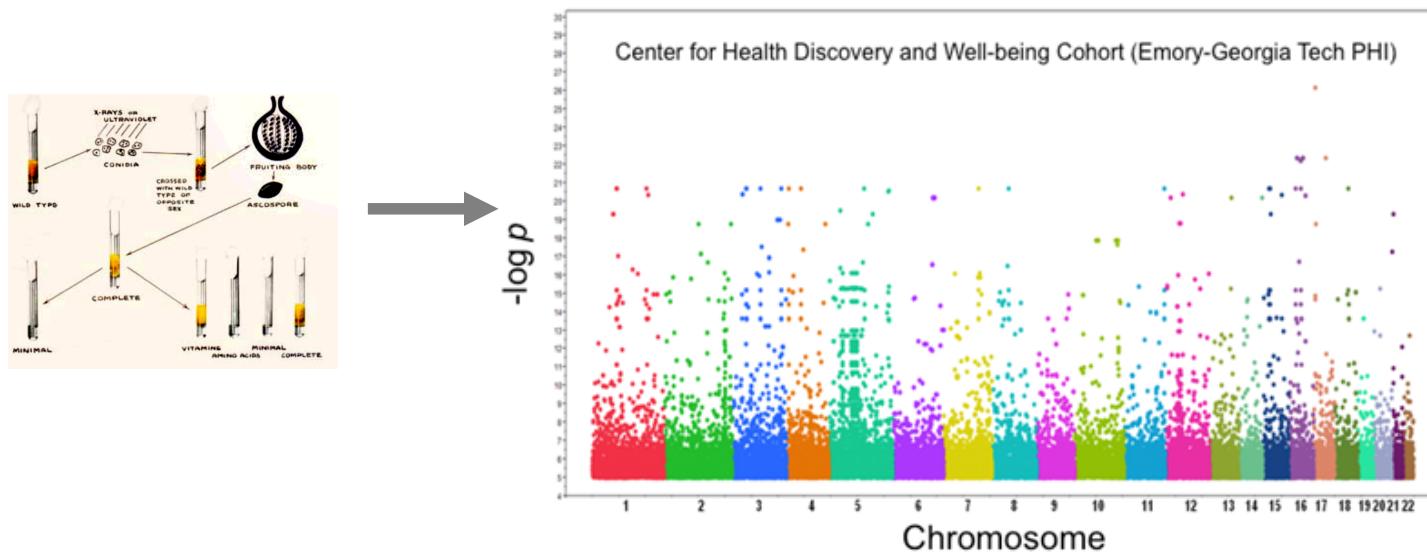
# Working with big data

- Global profiling may support many competing hypotheses
- False discovery rate
- Latent structures in data
- Statistical analysis and inference, machine learning
- Combining prior knowledge and domain knowledge
- Integration of different types of data and evidence
- Simulation

# Newer metabolomics cover majority of metabolic pathways

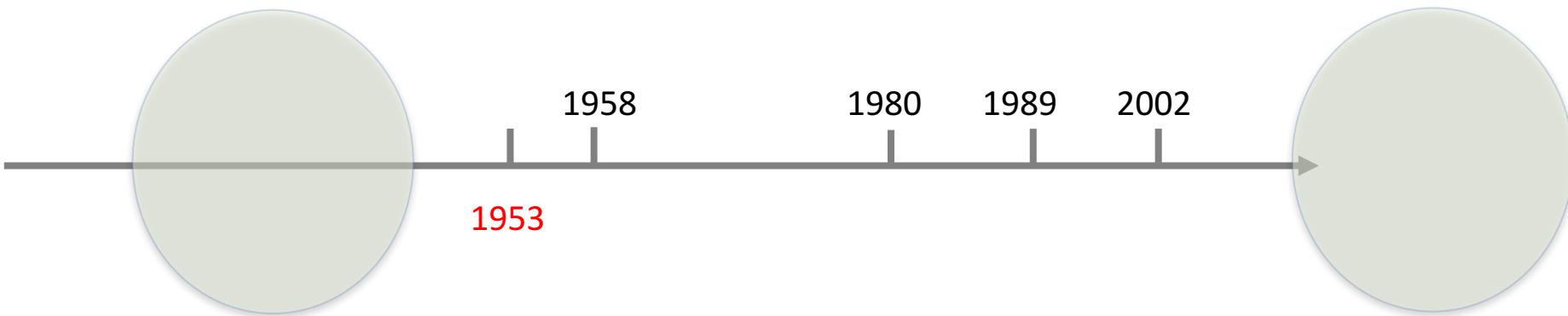


# $G \times M$ , Metabolome wide association of genetic variations



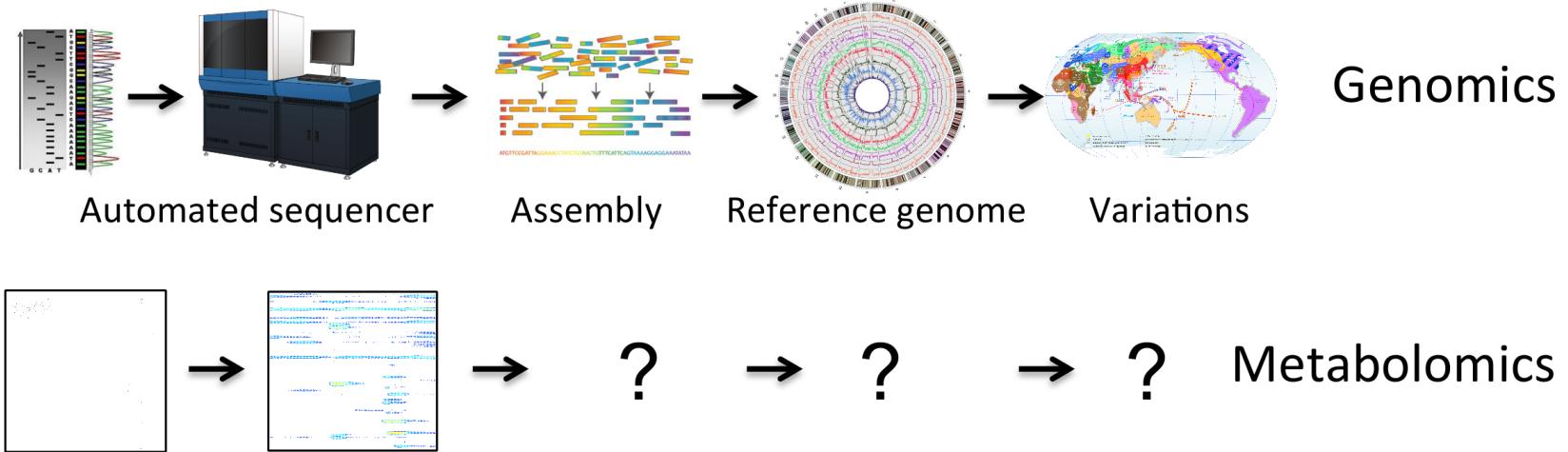
Courtesy: Dean P. Jones and Greg Gibson

# Biochemistry 1.0

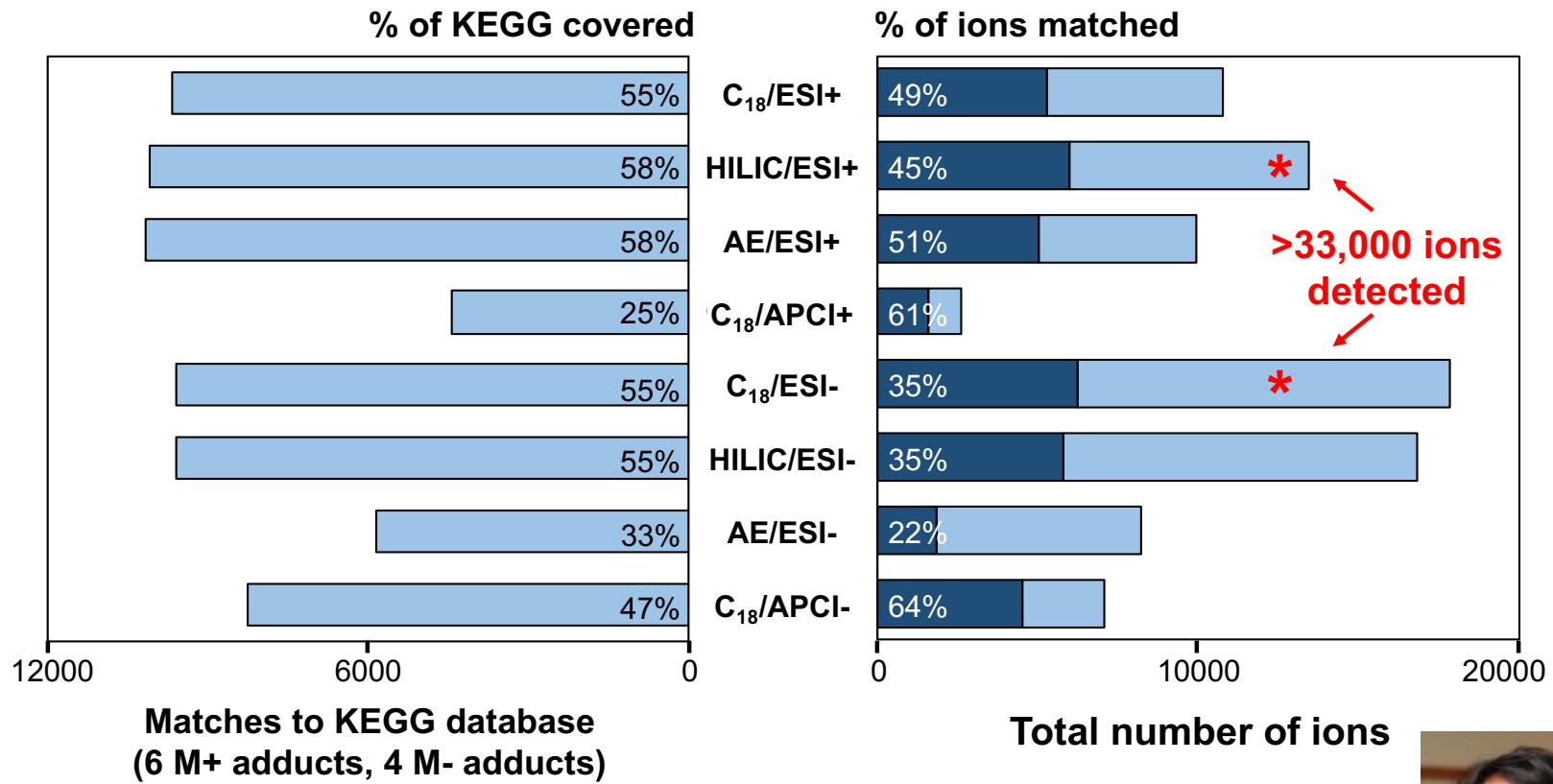


# Biochemistry 2.0

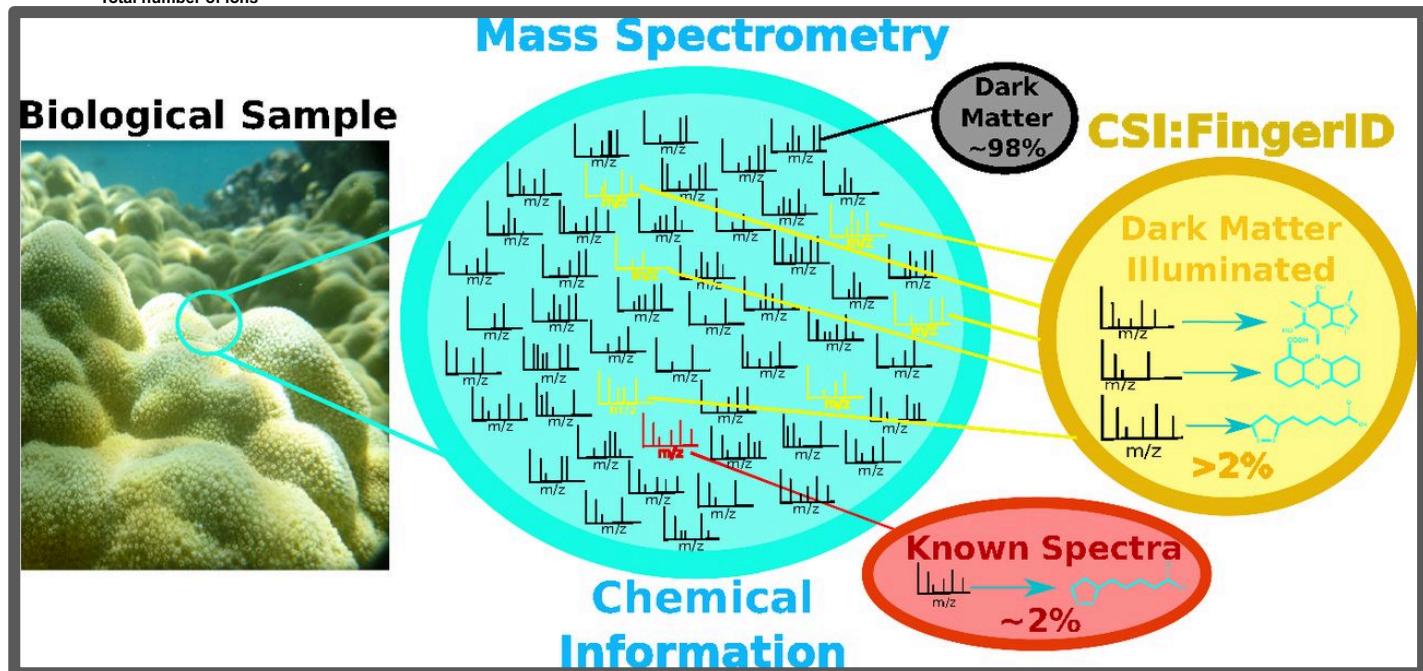
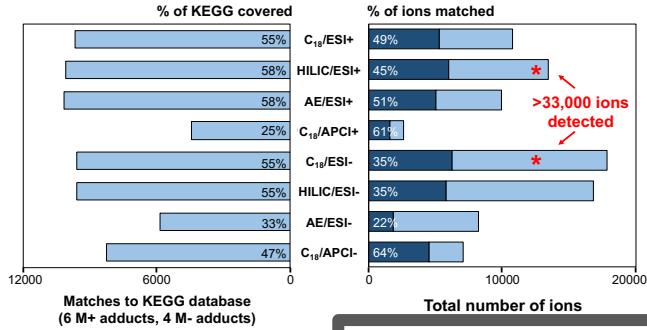
# Challenges and Opportunities of Metabolomics



# Most metabolomic peaks are unidentified



# Dark matters of Metabolomics



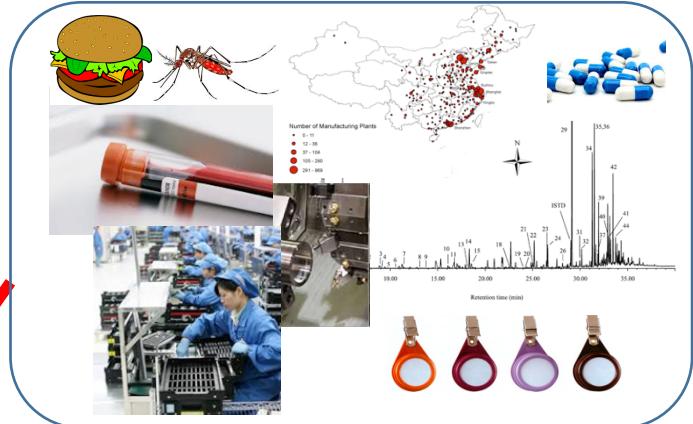
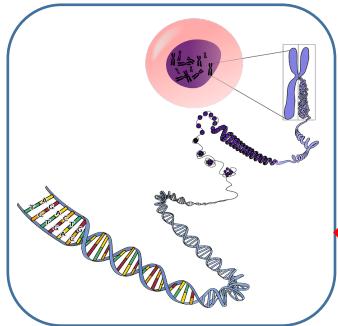
# Metabolomics is Biochemistry 2.0

- Catalog unidentified chemicals in humans
- Update of metabolic models, reactions and pathways (new maps)
- Signaling and regulatory roles of metabolites and chemicals
- Linking genomes and environmental exposures (exposome), including microbial products, diet and drugs
- Enabling precision medicine with digital health
- Data and computation intensive

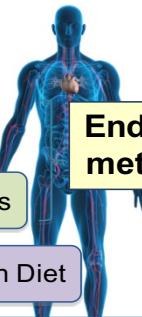
# G × M × E

# Environment

## Genome



## Gene function



Core Biological Metabolome

Microbiome-related Chemicals

Non-nutritive Chemicals in Diet

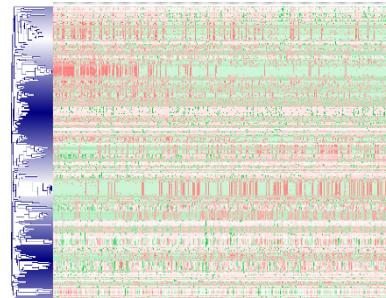
Supplements and Pharmaceuticals

**Environmental metabolome**

Commercial Products

Environmental Chemicals

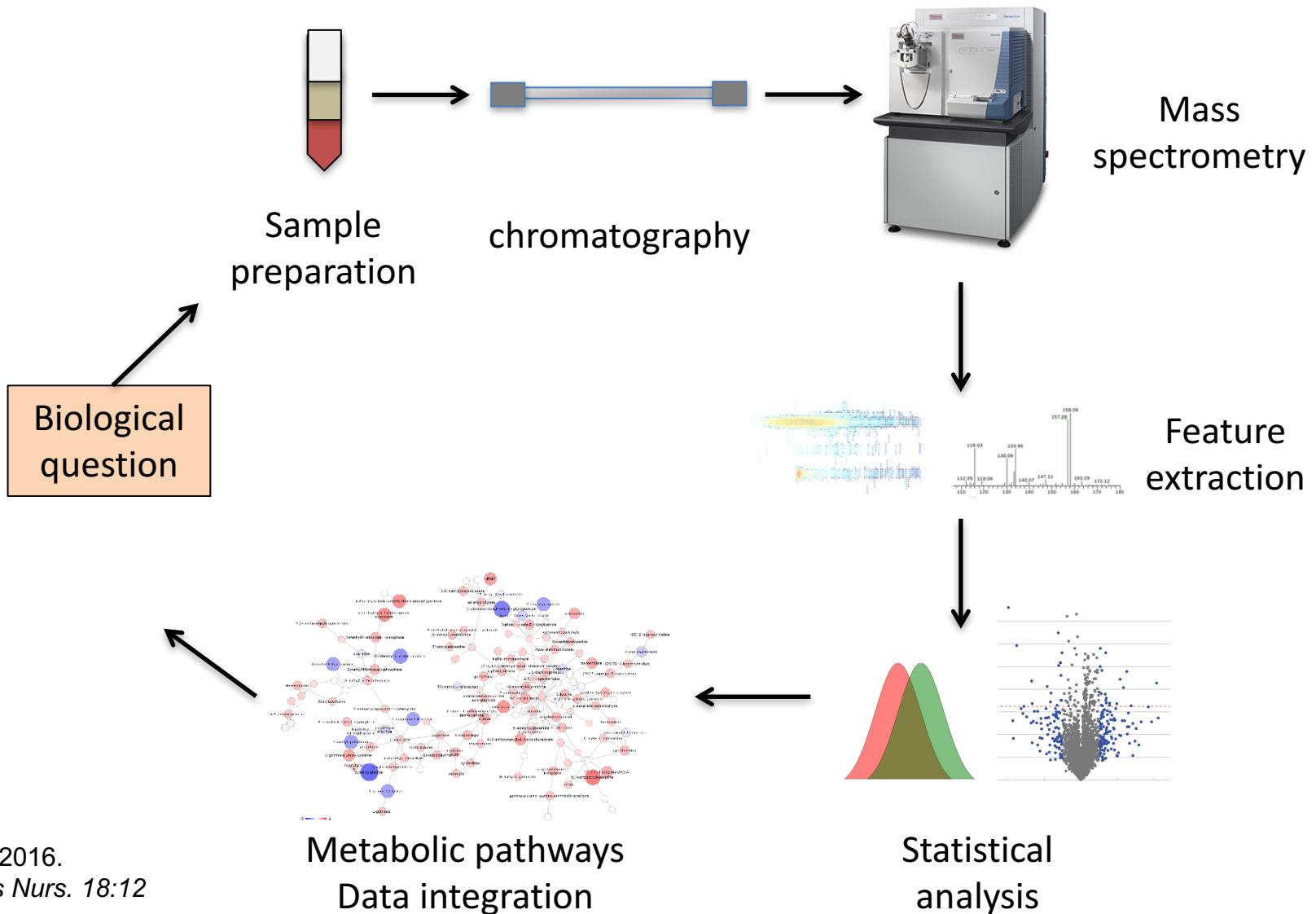
## Molecular response



## Metabolome

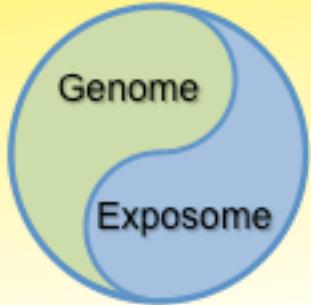
Body burden

# Metabolomics workflow (simplified)

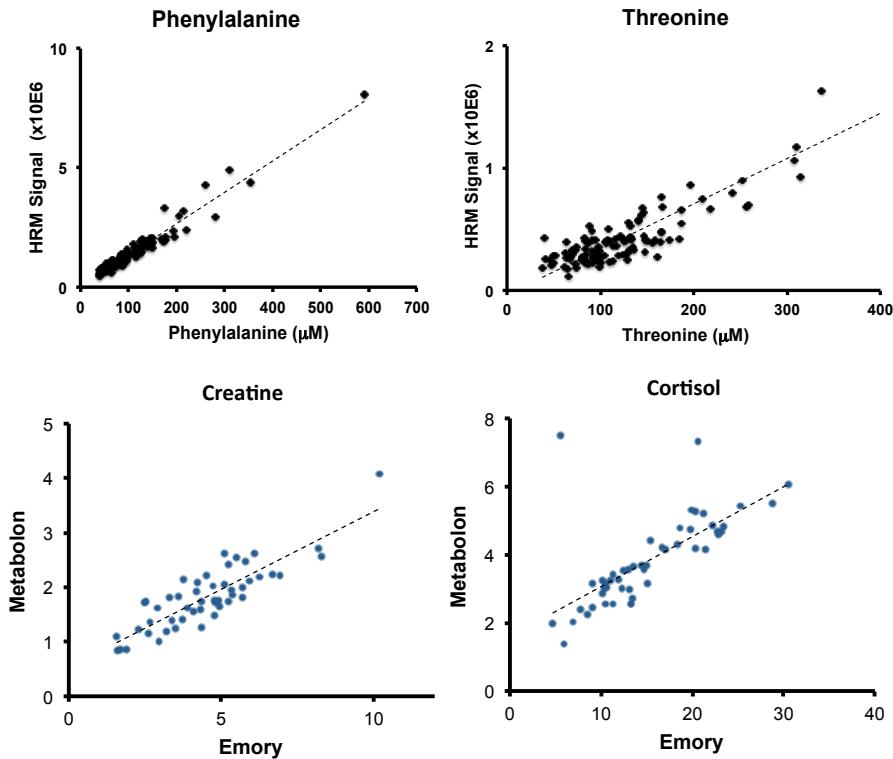


# Metabolomics technologies / platforms

- NMR vs Mass spectrometry
- Targeted vs untargeted Metabolomics
- LC vs GC
- MS vs MS<sup>n</sup>
- IM, EC, ...
- Isotope tracing



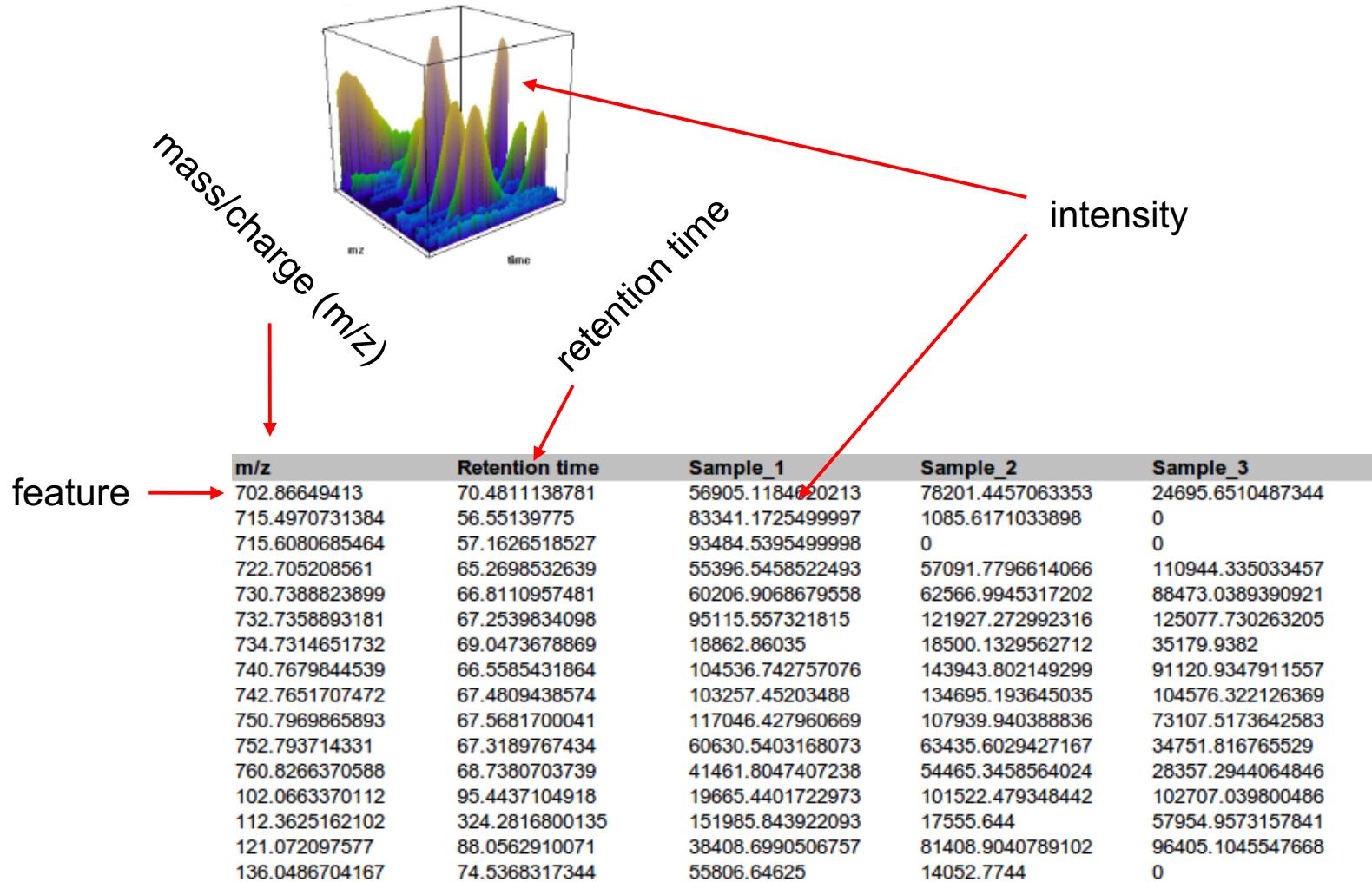
# Accurate quantification in high-throughput mode shown by cross-laboratory comparisons



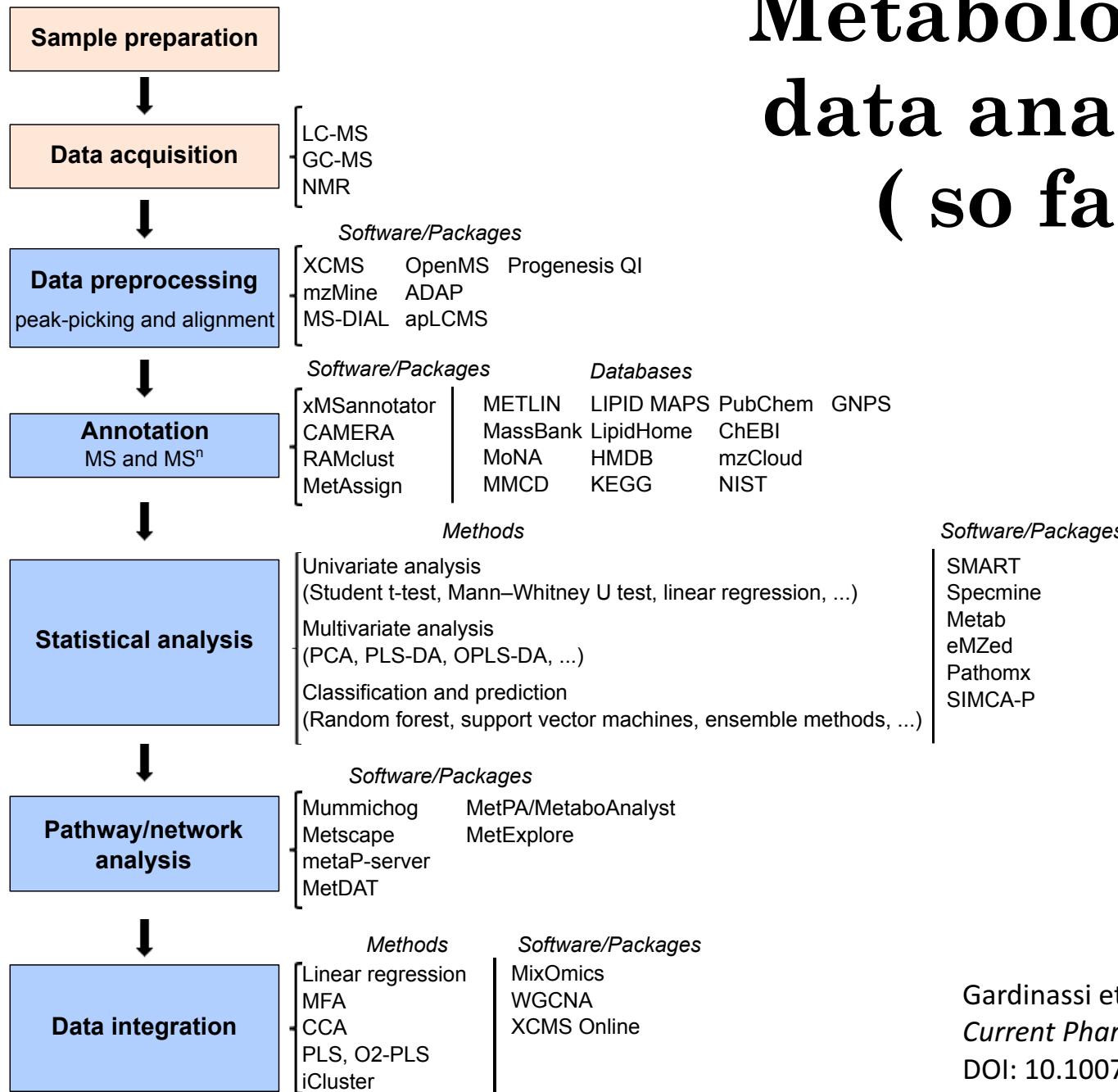
## HRM metabolite quantification in 30 orphan samples

Identity	Mean $\pm$ SD ( $\mu\text{M}$ )	HMDB ( $\mu\text{M}$ )
Arginine	148 $\pm$ 39	60 to 140
Glycine	280 $\pm$ 62	212-329
Histidine	100 $\pm$ 12	75 to 143
Ornithine	83 $\pm$ 28	54 to 94
Phenylalanine	131 $\pm$ 18	48 to 88
Threonine	136 $\pm$ 22	102 to 260
Tryptophan	56 $\pm$ 7	44 to 78
Tyrosine	84 $\pm$ 23	54 to 143
Glucose	4310 $\pm$ 1153	3900 to 6100
Kynurenine	2.0 $\pm$ 0.4	1.4 to 2.4
Carnitine	52 $\pm$ 9	30 to 57
Creatinine	93 $\pm$ 13	59 to 109
Creatine	16 $\pm$ 8	8.4 to 65

# Metabolomics data (LC-MS)



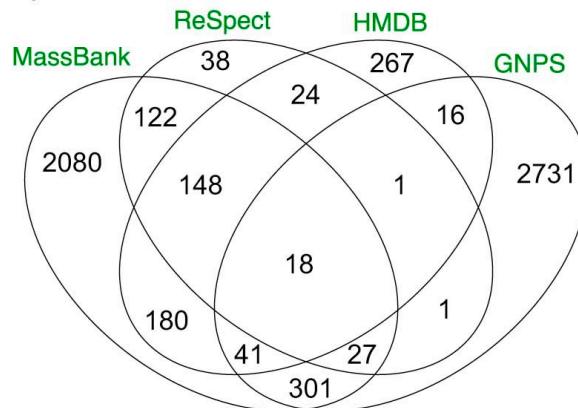
# Metabolomics data analysis ( so far)



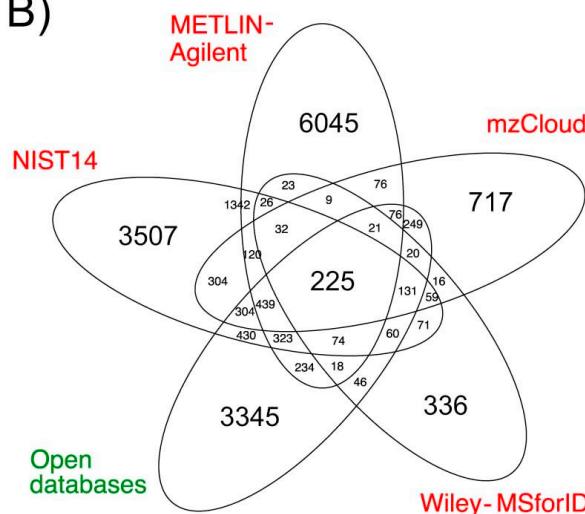
Gardinassi et al (2017)  
*Current Pharmacology Reports*,  
DOI: 10.1007/s40495-017-0107-0.

# Mass Databases (2016)

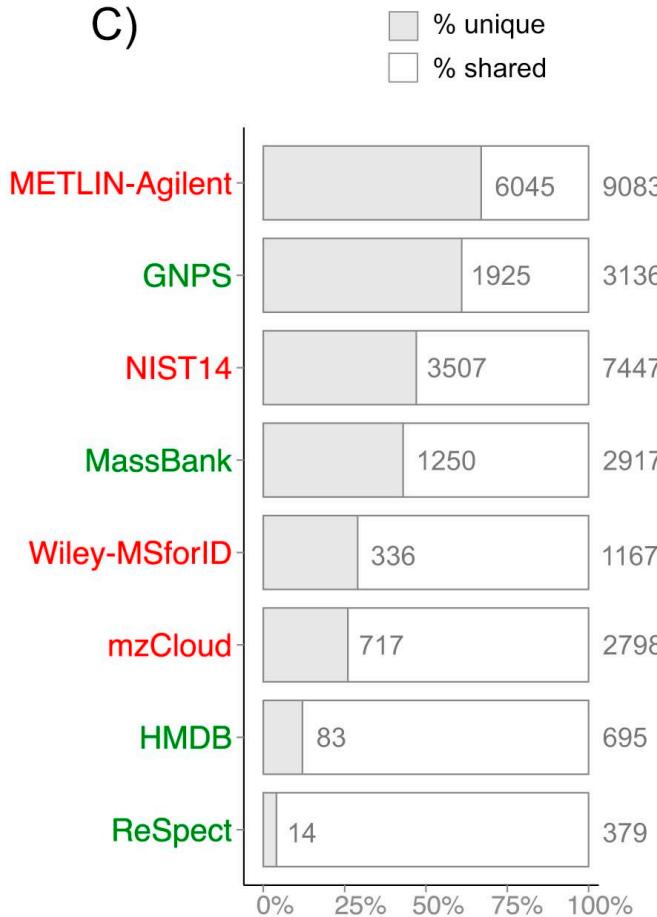
A)



B)



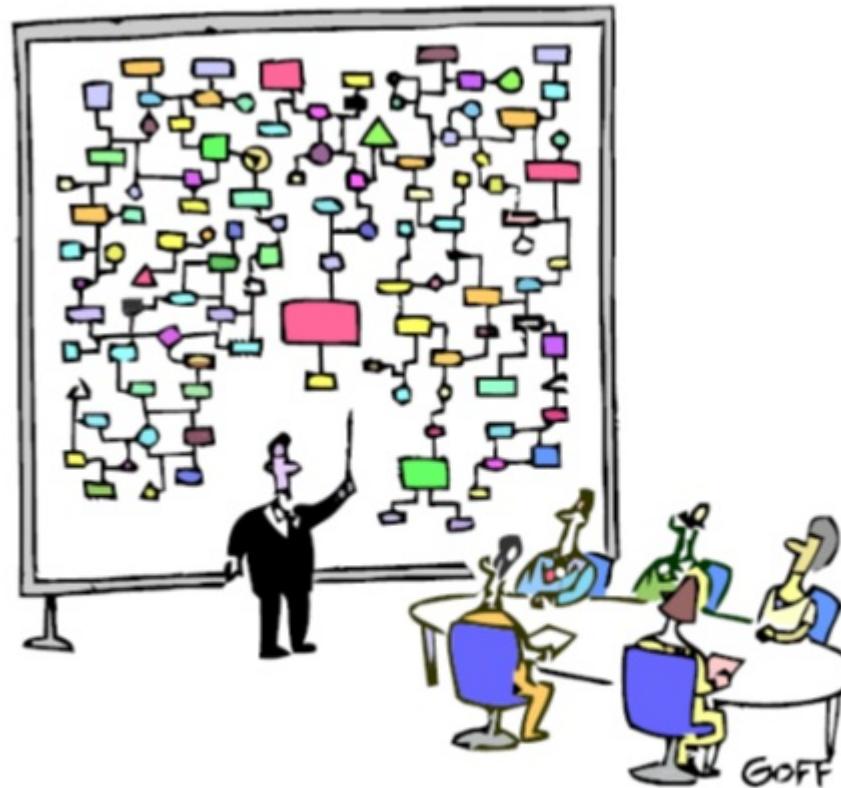
C)



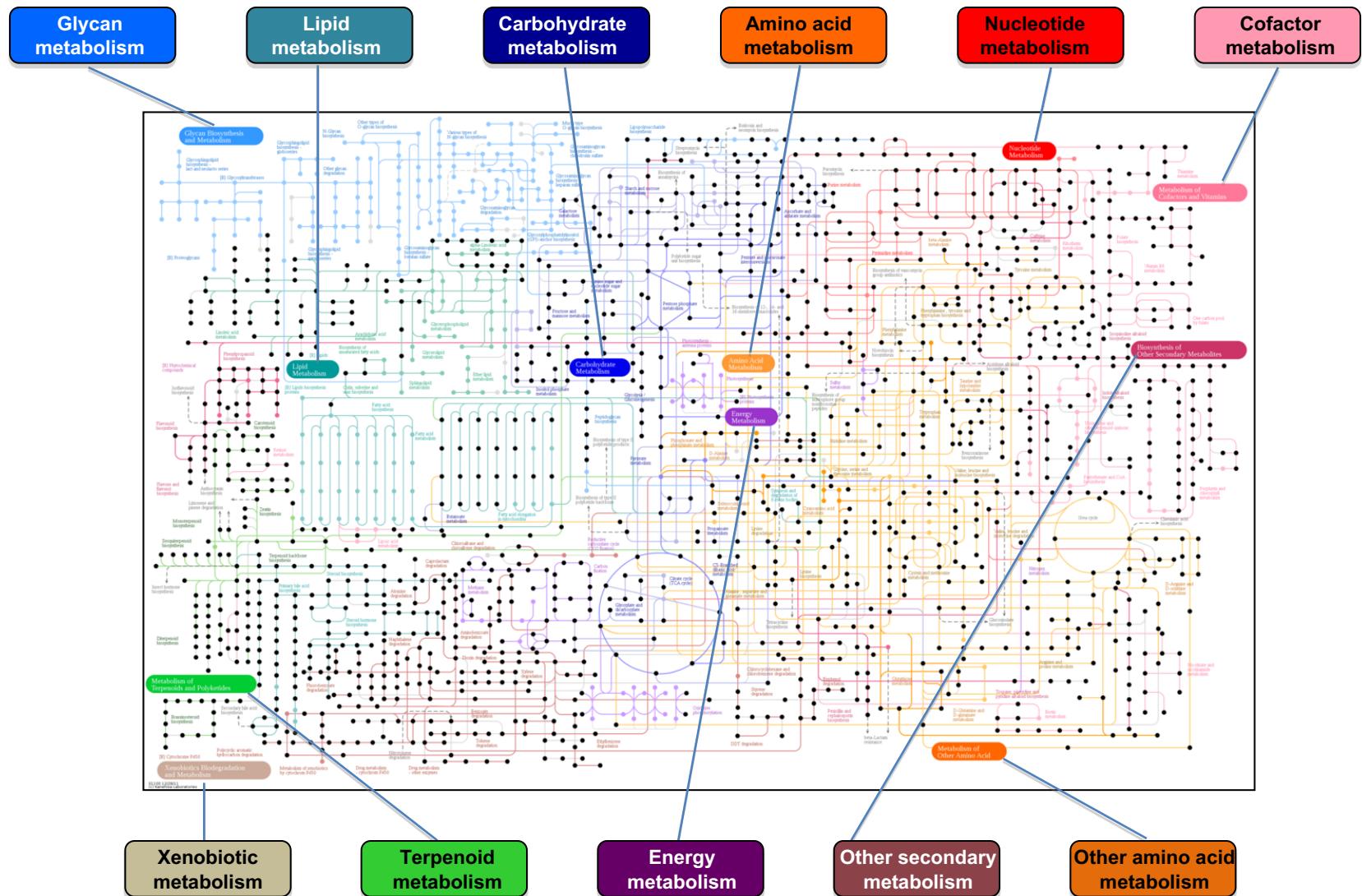
(A) Venn diagram showing the overlap between open mass spectral databases (HMDB, MassBank, GNPS, and ReSpect). (B) Venn diagram showing the overlap between five commercial databases (Agilent METLIN PMD, mzCloud, NIST 14, and Wiley MS) and open databases described in A. (C) Number and percentage of unique and shared compounds (i.e., InChIKey) with  $\text{MS}^n$  ( $n \geq 2$ ) data in each database in relation to all eight resources.

Vinaixa, M., Schymanski, E.L., Neumann, S., Navarro, M., Salek, R.M. and Yanes, O., 2016. Mass spectral databases for LC/MS-and GC/MS-based metabolomics: state of the field and future prospects. *TrAC Trends in Analytical Chemistry*, 78, pp.23-35.

# Lot of pathways are known



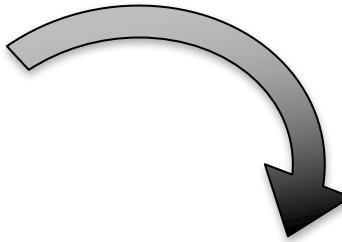
"And that's why we need a computer."



# Towards new metabolic maps



1482

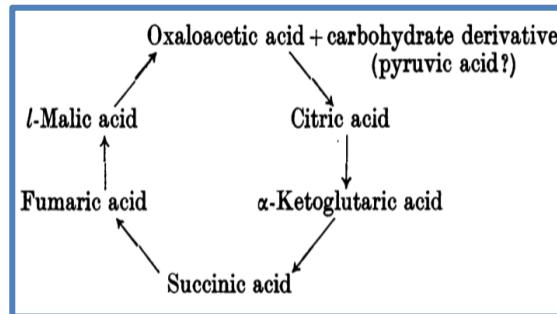
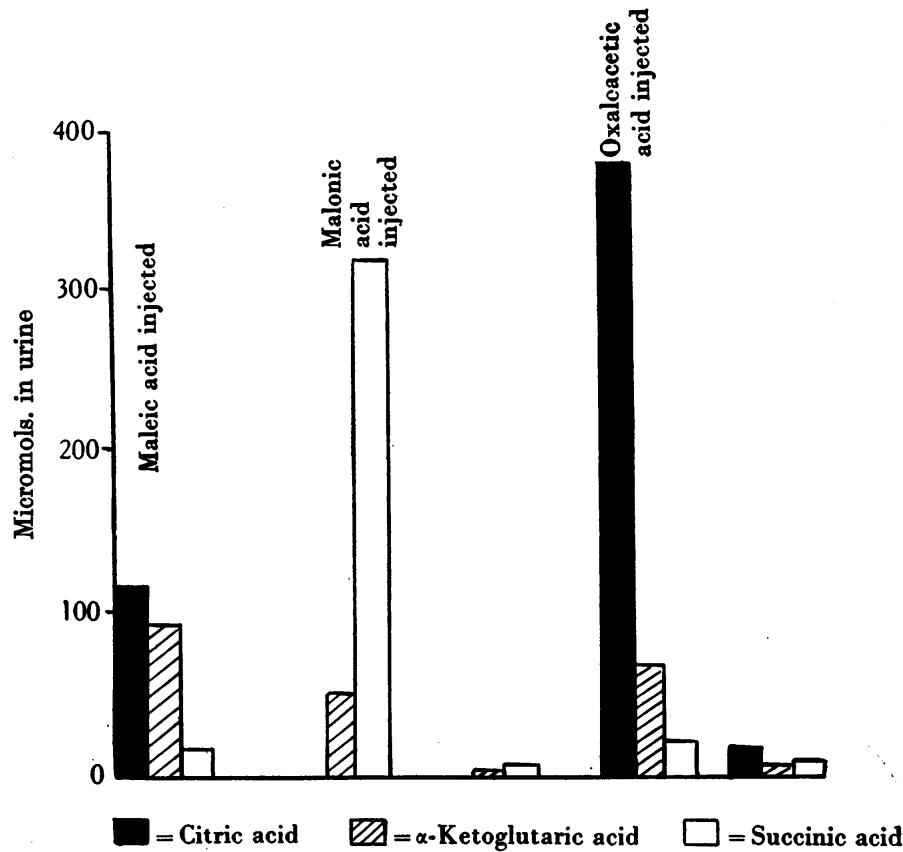
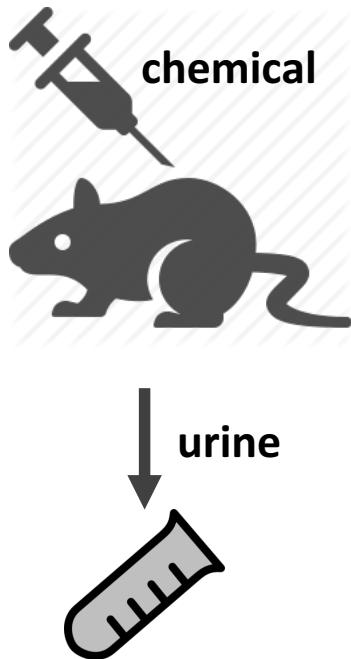


Political Map of the World, January 2015



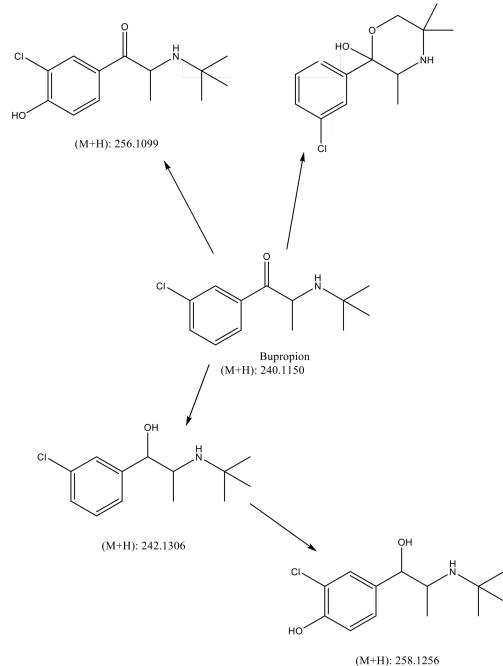
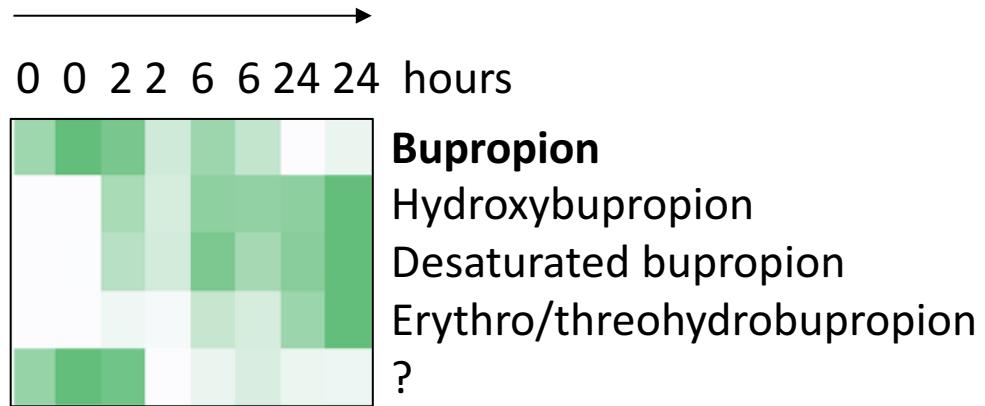
2015

# Charting metabolic pathways



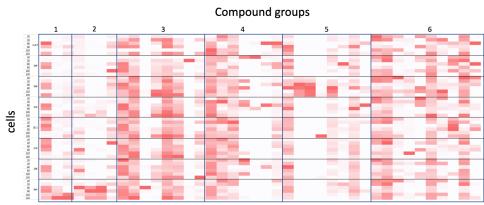
Krebs et al. 1938.  
Biochem Journal. 32:113

# Charting metabolic pathways (2019)



Jones/Li/Morgan/Miller  
NIEHS U2C ES030163

# Charting metabolic pathways



- Capturing unknown; tandem mass spec; isotope tracing in cells and whole organisms
- Combination with genetic tools
- Domain applications, tumors, microbiome, immune cells
- Azimuth database; Interfacing with computational reconstruction of biochemical networks

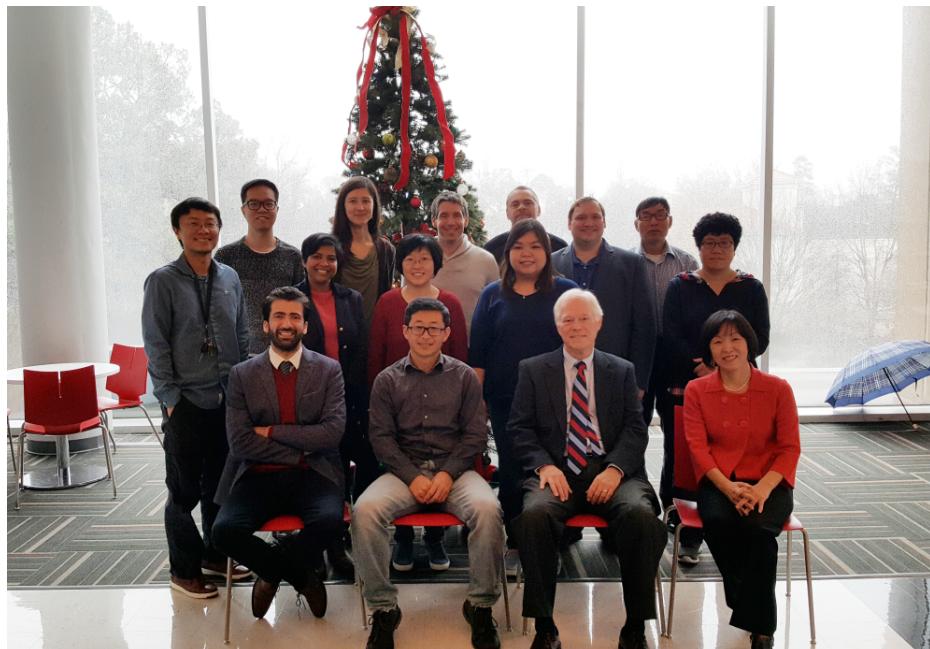
# Objectives

- Understanding of experimental design and statistics
- Using Jupyter notebooks for data analysis
- Web tools such as XCMS Online, MetaboAnalyst and Mummichog server
- Metabolite reporting standards
- Discussion of research projects

# Acknowledgement

## CHDS

Barbara Cohn  
Piera Cirrilo  
Nickilou Krigbaum  
**Stanford University**  
Bali Pulendran  
**University of Colorado**  
Myron Levin  
Adriana Weinberg



## Emory University

Rafi Ahmed  
Mark Mulligan (NYU)  
Nadine Roushafel  
Mary Galinski  
Judith Fridovich-Keil  
Taylor Fischer  
Miriam Vos

Dean P. Jones  
Young-Mi Go  
Douglas Walker (MSSM)  
ViLinh Tran  
Bill Liang  
Luiz Gardinassi  
Ken Liu  
M. Ryan Smith  
Tianwei Yu  
Edward Morgan  
Choon Myung Lee  
Grant Singer

**Funding support** from US National Institutes of Health U19 AI090023, HHSN272201300018I, UH2 AI132345, U01 CA235493, U2C ES030163, P30ES019776, U2C ES026560, P50 ES026071, R01 DK107900, R01 AI121252, R01 GM124061, and EPA (83615301)