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

Introduction to metabolomics research

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Targeted
Metabolomics &
Proteomics
Laboratory

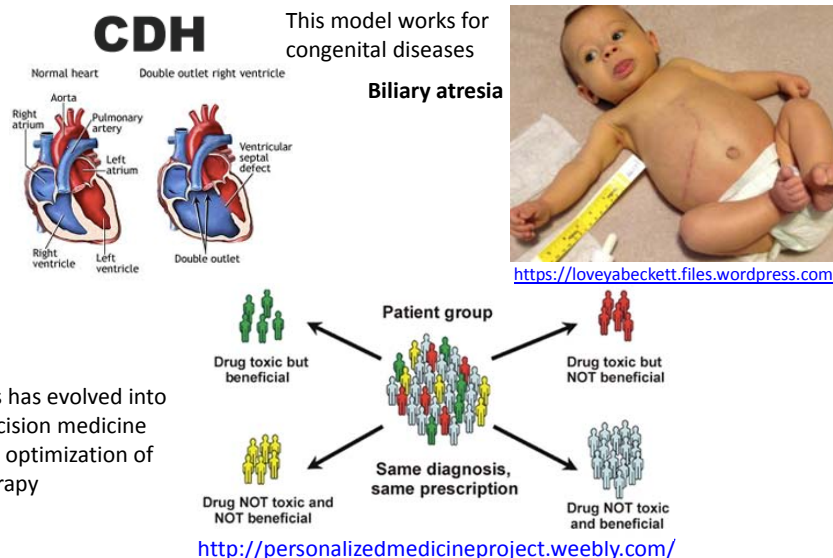
What is “Metabolomics”?

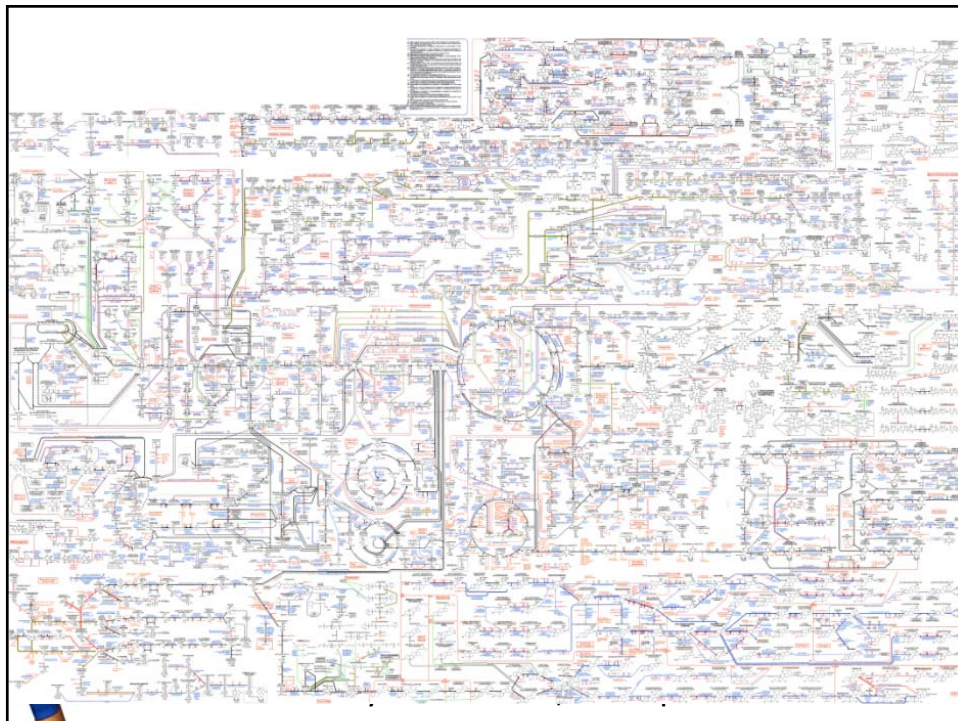
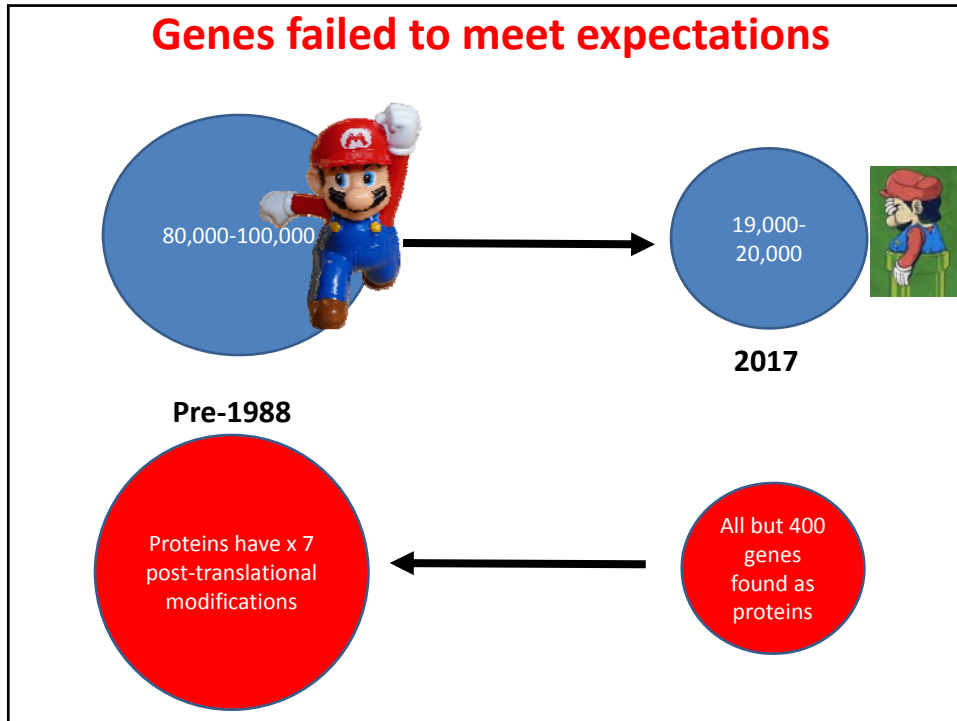
- Metabolomics is like other types of –omics analysis (microarray, RNA-Seq, proteomics, etc.)
 - Offers a “comprehensive” view of all detectable chemicals (not just metabolites)
 - Can be applied to body fluids
 - Plasma/sera, urine, saliva, tears, fecal water, etc.
 - Also to tissues
 - Liver, lung, heart, kidney, brain, eyes, etc.
 - And to single cells
 - Human, rodent, yeast, bacteria, etc.

Defining who we are chemically

- Are we “*Living in the Promisedland*” as per Willie Nelson’s song?
- Does an understanding of the functions of human genes define the chemical make up of our body fluids and tissues?
- How does metabolomics provide information on the circulating chemicals?
- Are the detected chemicals metabolites produced by human enzymes?
- So, what are we really exposed to? And does it make a difference?

A great deal of emphasis is being placed on the importance of DNA sequencing





Where does the metabolome come from?

- It starts with what fixes CO₂ and N₂



Trees convert
CO₂ to organic
compounds



Field of soybeans – they fix
N₂ because of nitrogen-
fixing bacteria in their root
nodules

Plants have more genes than humans

- Why? Plants can't run away!!
- Instead, they have to practice chemical warfare to prevent attack by aphids and microorganisms
- Many plants are poisonous to us
- Understanding which plants were safe to eat, or were so if cooked, represented the rise of agriculture and civilization



Compounds in plants and fruits

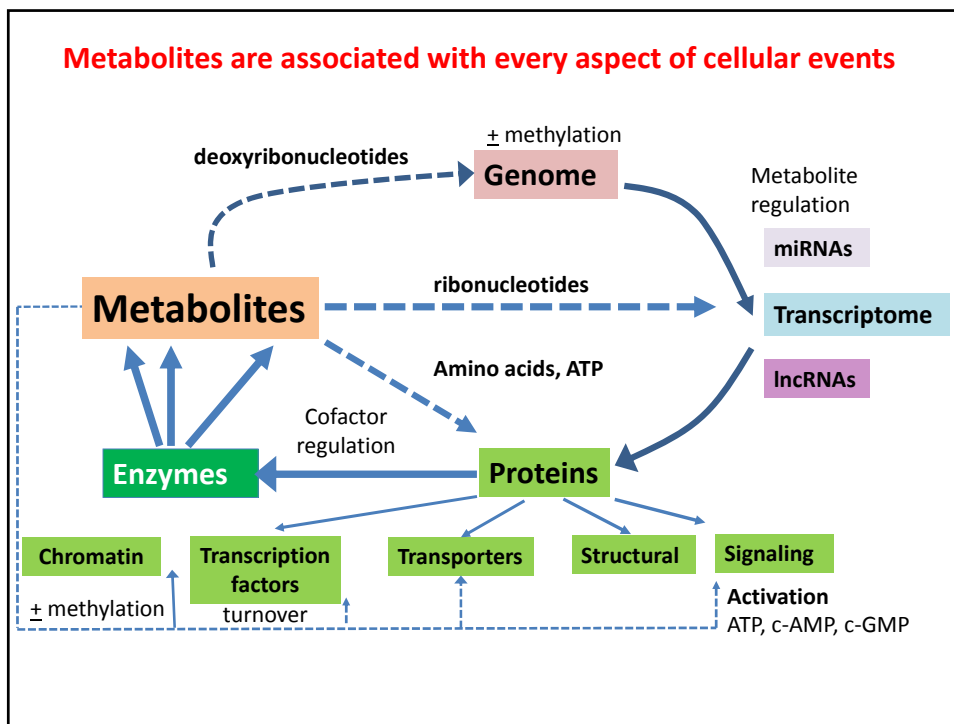
- Carotenoids
- Many vitamins
- Polyphenols and anthocyanins
- Not made by human cells



Other sources of body chemicals

- **The microbiomes**
 - Humans are not single organisms
 - Instead, we are super-organisms
 - The gut microbiome has 10 times the number of cells found in the rest of the (human) body
 - It makes novel compounds that are absorbed, enter the blood stream and tissues
- **Chemicals from the environment**
 - industrial contaminants, therapeutics, supplements
- **Interactions between the xenobiotics and the human enzyme systems**

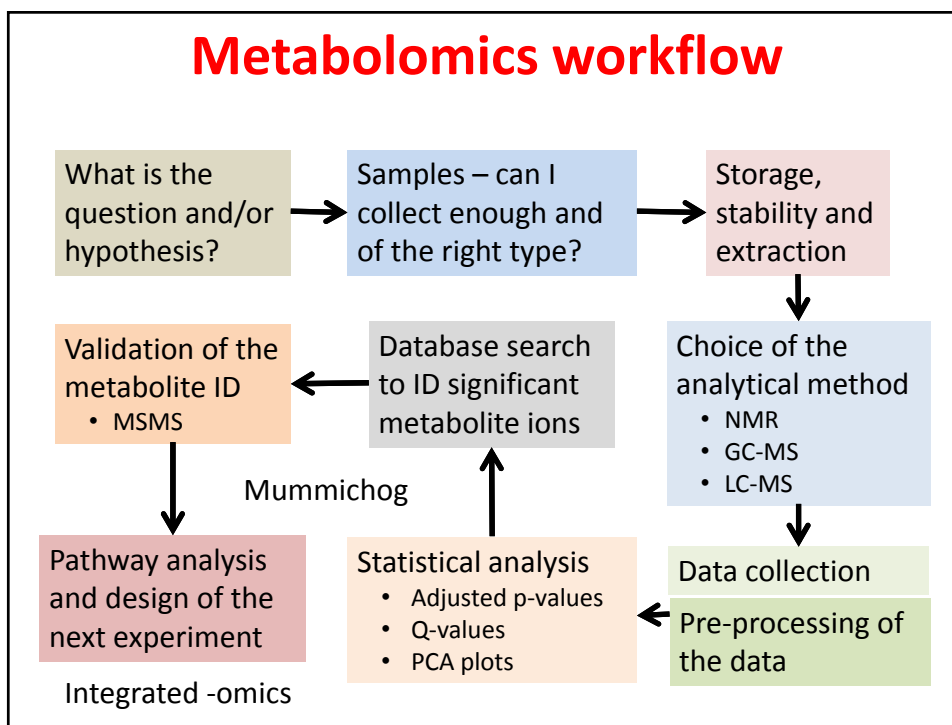




The metabolome is very complex!



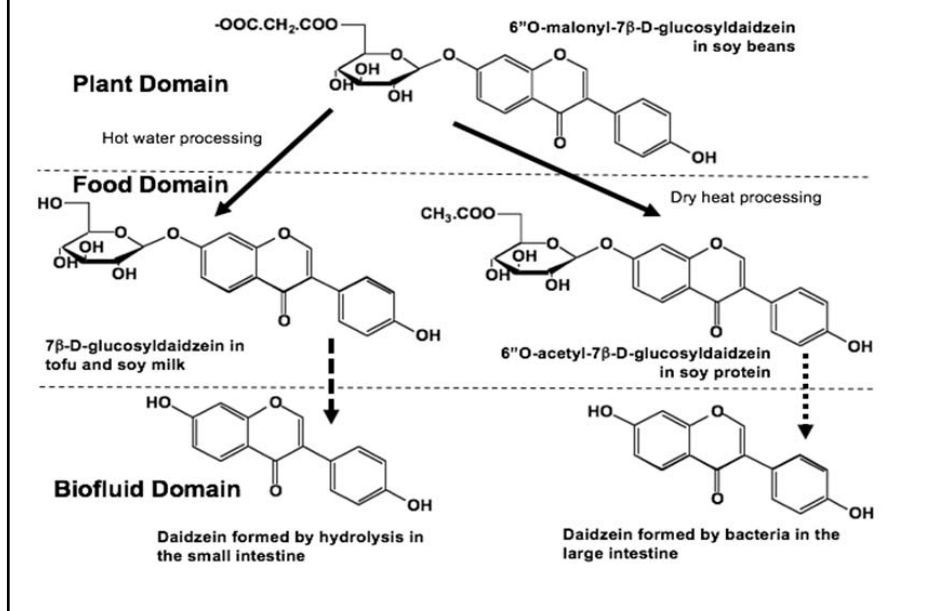
Metabolomics workflow



Course goals

- To understand
 - The **vital** roles of metabolites
 - The **origins** of metabolites
 - That metabolomics is **high dimensional**
 - The best methods for **extracting metabolites**
 - How to select the **analytical approach**
 - **Qualitative** and **statistical analysis** of the data
 - How to **identify** the “interesting” metabolites
 - How to map to (or define) **pathways**
 - The value of **stable isotopes**

Complexity in metabolism beyond cell culture Inter-genome events



Where did metabolomics come from?

Transition of mass spectrometry to biology

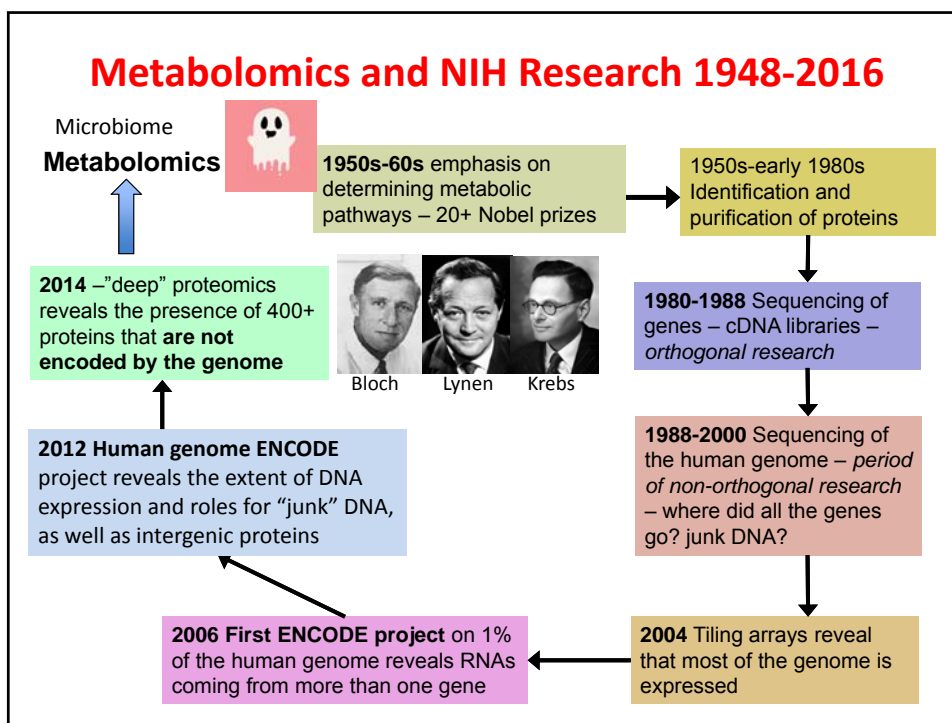


Ralf Schoenheimer



David Rittenberg

- While the politicians, tyrants, dictators and despots were salivating at the thought of developing nuclear weapons from unstable isotopes in the early part of the 20th Century, two scientists began the pursuit of the peaceful use of stable isotopes, initially deuterium (^2H), and later carbon (^{13}C) and nitrogen (^{15}N), to study biochemical pathways
- Understanding the pathways of metabolism was born



Progress in LC-MS

- Commercial HPLC appeared in the early 1970s to separate thermally stable and unstable molecules
- The challenge remained to find a way to get the unstable compounds into the gas phase
 - Applied to macromolecules (peptides, proteins) as well as metabolites
- Thermospray had some initial success
- **Electrospray ionization** and **chemical ionization** radically changed analysis, allowing compounds to go into the gas phase at atmospheric pressure and room temperature

LC-MS

- Suddenly, there were what appeared to be no limits (or very few) to what could be analyzed
- Unheard of, robust mass spectrometers came into play
 - “A reliable mass spectrometer” was considered in 1990 to be an oxymoron

Types of LC-MS analysis

Single quadrupole
LC-MS analysis

LC-time-of-flight
(TOF)-MS

FT-ICR MS

Orbi-trap

Triple quadrupole
LC-MS analysis

Multiple reaction
monitoring (MRM)

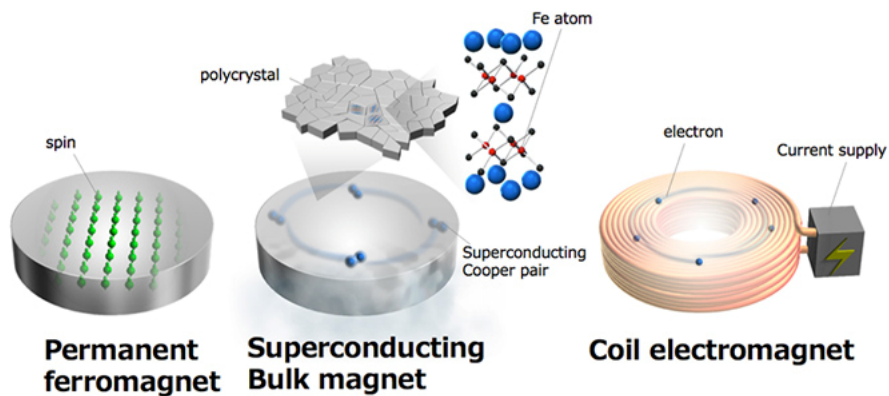
Q-TOF

TripleTOF



Ion Mobility

NMR spectroscopy and metabolomics

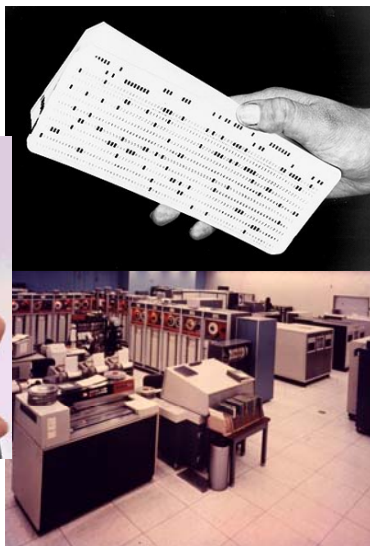
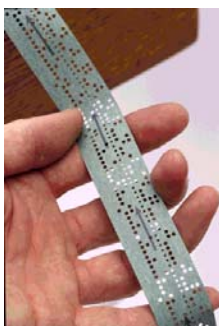


https://nationalmaglab.org/images/news_events/news/2015/october/pnictide_magnetism_1oct2015.jpg

NMR has had several critical development steps – Fourier Transform analysis of collected data, increase in field strength with superconducting magnets, micro-coil, cryogenic analysis, hyperpolarization.

Changing times in Computing

- 1950 The Cambridge colleagues of Watson and Crick calculated the structure of DNA by putting data onto punched cards and taking them by train to London for analysis – and to the fog – the “cloud” in 1950s
- 1964 Seymour Cray develops the CDC 6600 (1 Mflops)
- 1967 I used paper tape to collect data from a radio gas chromatograph and then submitted them via a terminal reader to the CDC 6600 at the University of London



Today in Computing



On my desk in 2017

- The Apple MacBook Air with 2 quad core Intel i7 processors
 - Operates at 2.0 GHz
 - Memory of 8 GB
 - Access 1.333 GHz
 - 512 GB Flash memory storage
 - 10 Gbs Thunderbolt I/O
- Also cost ~\$2,000



IBM Blue-Gene

- Parallel processing with 2,048 700 MHz computers operating at 4.733 Tflops
- Replaced by Cheaha, in its current configuration it has 48 compute nodes with two 2.66GHz 6-core Intel CPUs per node (576 cores total)
- It operates at 6.125 Tflops

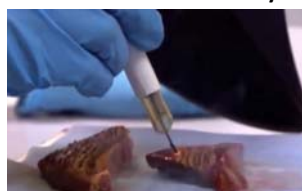
MRC-NIHR National Phenome Centre



600 MHz NMR instruments
in surgical suite



Mass spectrometers (10 Q-TOFs) each
dedicated to one assay format



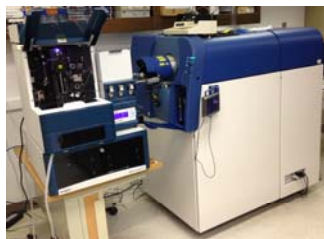
Iknife - revolutionizing surgery

This is Next-GEN precise medicine

The UK National Phenome Center, LC-MS labs



UAB capabilities in metabolomics

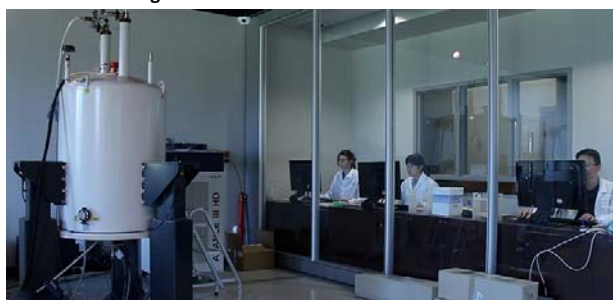


SCIEX 5600 TripleTOF
with Eksigent nanoLC

TMPL mass spec lab
MCLM 459/427
Stephen Barnes, Director
205-934-7117/3462



SCIEX 6500 Qtrap with SelexION



Central Alabama NMR facility
Chemistry Bdg
William Placzek, Director
205-934-2465

Great challenges in metabolomics

- **The extent of the metabolome**
 - From gaseous hydrogen to earwax
- **Having complete databases**
 - METLIN has 220,000+ metabolite records, but your problem always creates a need to have more
 - Improvement in the size of a MSMS database
- **Storing and processing TBs of data**
- **Standards and standard operating procedures**
- **Being able to do the analyses in real time**

NIH Common Fund Metabolomics Program

- **Metabolomics Workbench:**
<http://www.metabolomicsworkbench.org/>
- **Regional Comprehensive Metabolomics Research Centers**
 - University of Michigan: <http://mrc2.umich.edu/index.php>
 - UC Davis Metabolomics Center: <http://metabolomics.ucdavis.edu/>
 - UNC-CH: <http://www.uncnri.org/wp-content/uploads/2016/12/NIHERCMRC.pdf>
 - SE Center for Integrated Metabolomics: <http://secim.ufl.edu/>
 - Resource Center for Stable Isotope Metabolomics:
<http://bioinformatics.cesb.uky.edu/bin/view/RCSIRM/>
 - Mayo Clinic Metabolomics Resource: <http://www.mayo.edu/research/core-resources/metabolomics-resource-core/overview>