

Metabolomics, a rapidly evolving contributor to precision medicine, and how to do it

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UAB

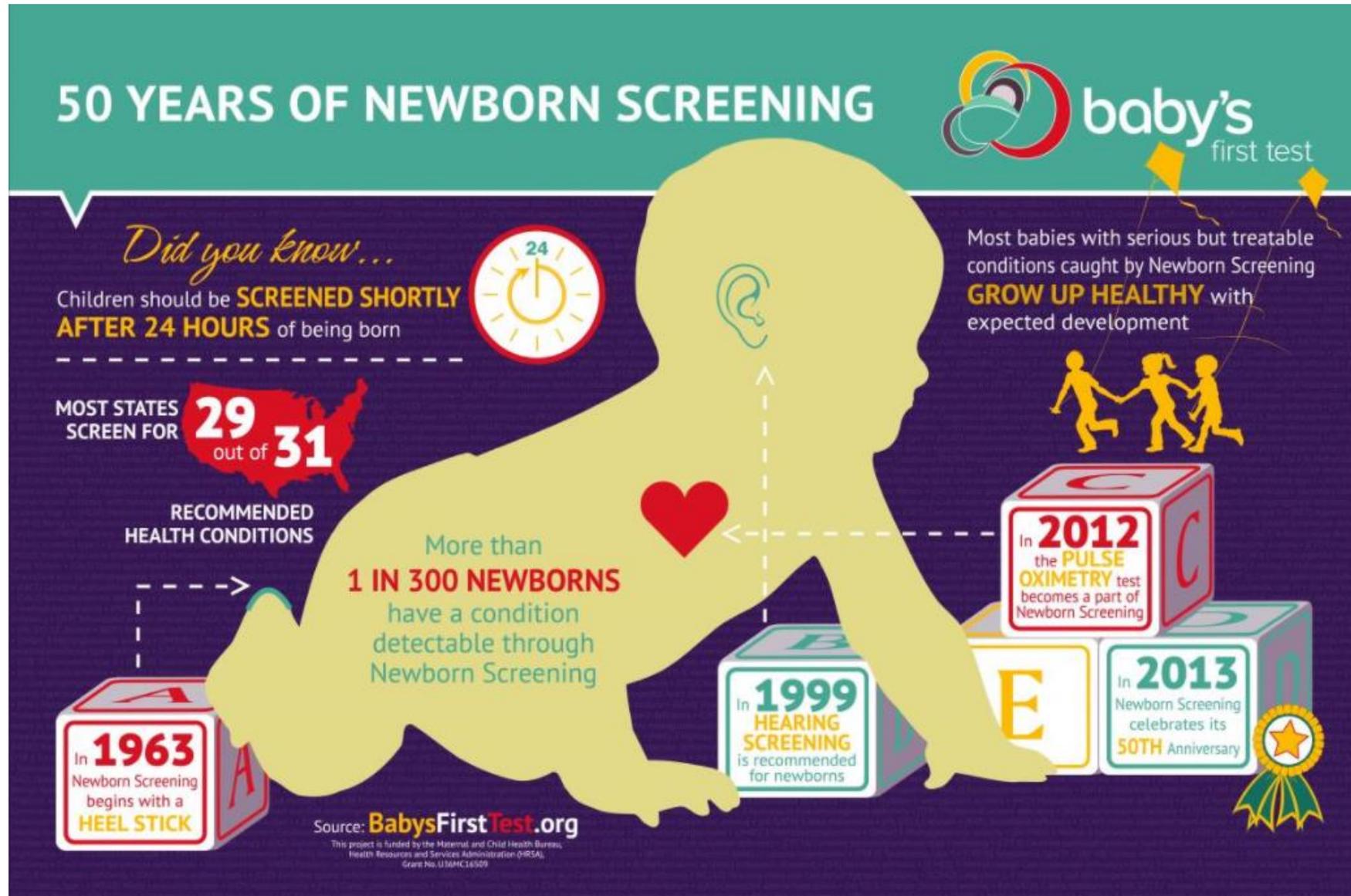
To be posted at <http://tmpl.uab.edu>

See also <http://www.uab.edu/proteomics/massspec/classes/schedule.php>

Synopsis

- **Why has the metabolome (and metabolomics) become so important?**
- **What is the metabolome?**
- **How do I do a metabolomics experiment?**
- **What platform can I use?**
- **How do I analyze the data?**
- **Can I integrate metabolomics data with other –omics data?**

Targeted metabolomics is not new



Metabolomics and NIH Research 1948-2018

Microbiome



Metabolomics



2014 – “deep” proteomics reveals the presence of 400+ intergenic proteins



2012 Human genome ENCODE project reveals the extent of DNA expression and roles for “junk” DNA, as well as intergenic proteins

1950s-60s emphasis on determining metabolic pathways – 20+ Nobel prizes



Bloch Lynen Krebs

The apparent circle is not a circle. Instead, it’s a spiral as higher and higher knowledge is gained.

1950s-early 1980s Identification and purification of proteins

1980-1988 Sequencing of genes – cDNA libraries – *orthogonal research*

1988-2000 Sequencing of the human genome – *period of non-orthogonal research* – where did all the genes go? junk DNA?

2004 Tiling arrays reveal that most of the genome is expressed

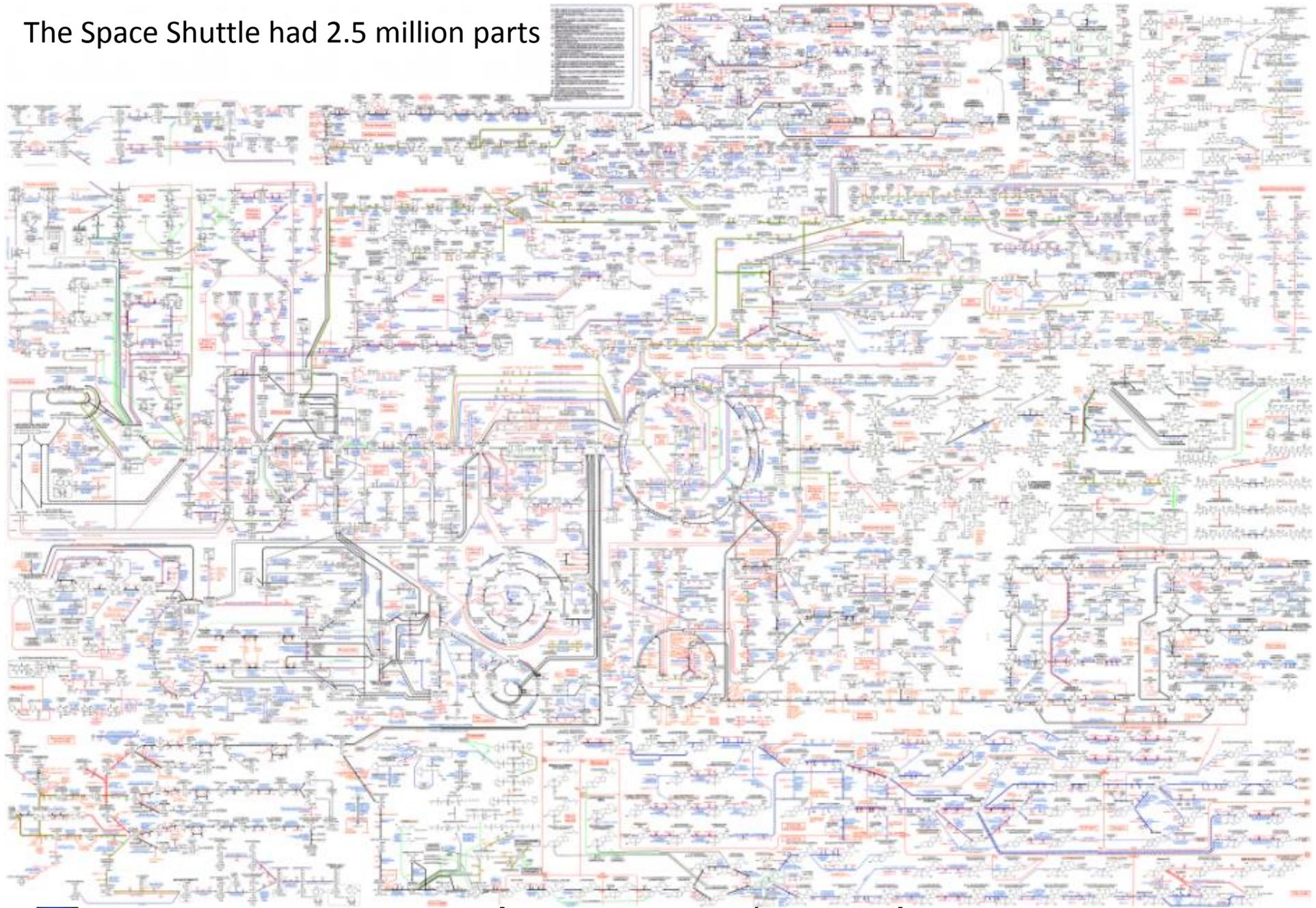
2006 First ENCODE project on 1% of the human genome reveals RNAs coming from more than one gene

So, what is the metabolome?

The metabolome is very complex!



The Space Shuttle had 2.5 million parts



Amino acids

Essential

Non-essential

Arg*

Ala

His

Asn

Ile

Asp

Leu

Cys

Lys

Gln

Met

Glu

Phe

Gly

Thr

Pro

Trp

Ser

Val

Tyr



Have to eat foods rich in these



Meat eater

Berry eater

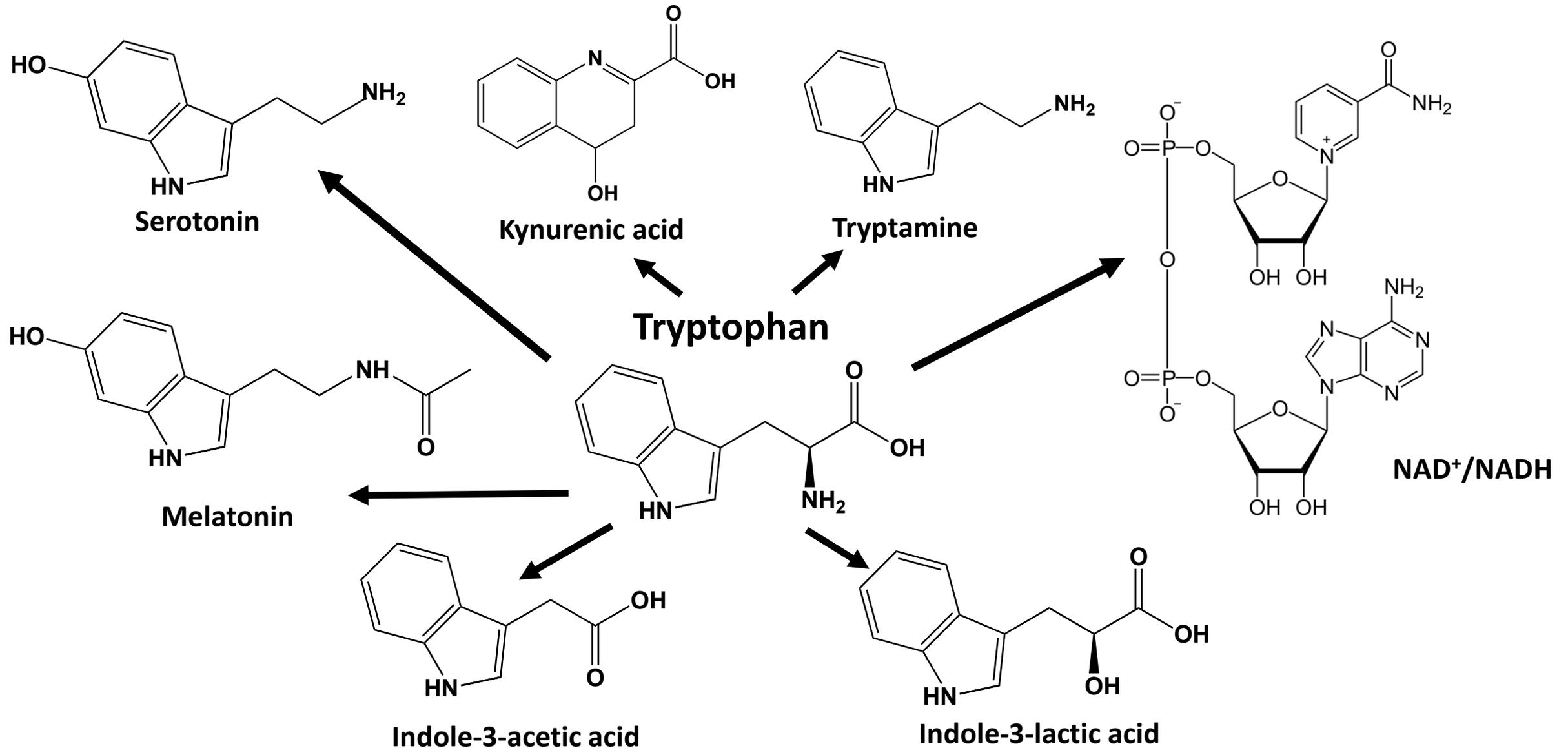


Vegetarian

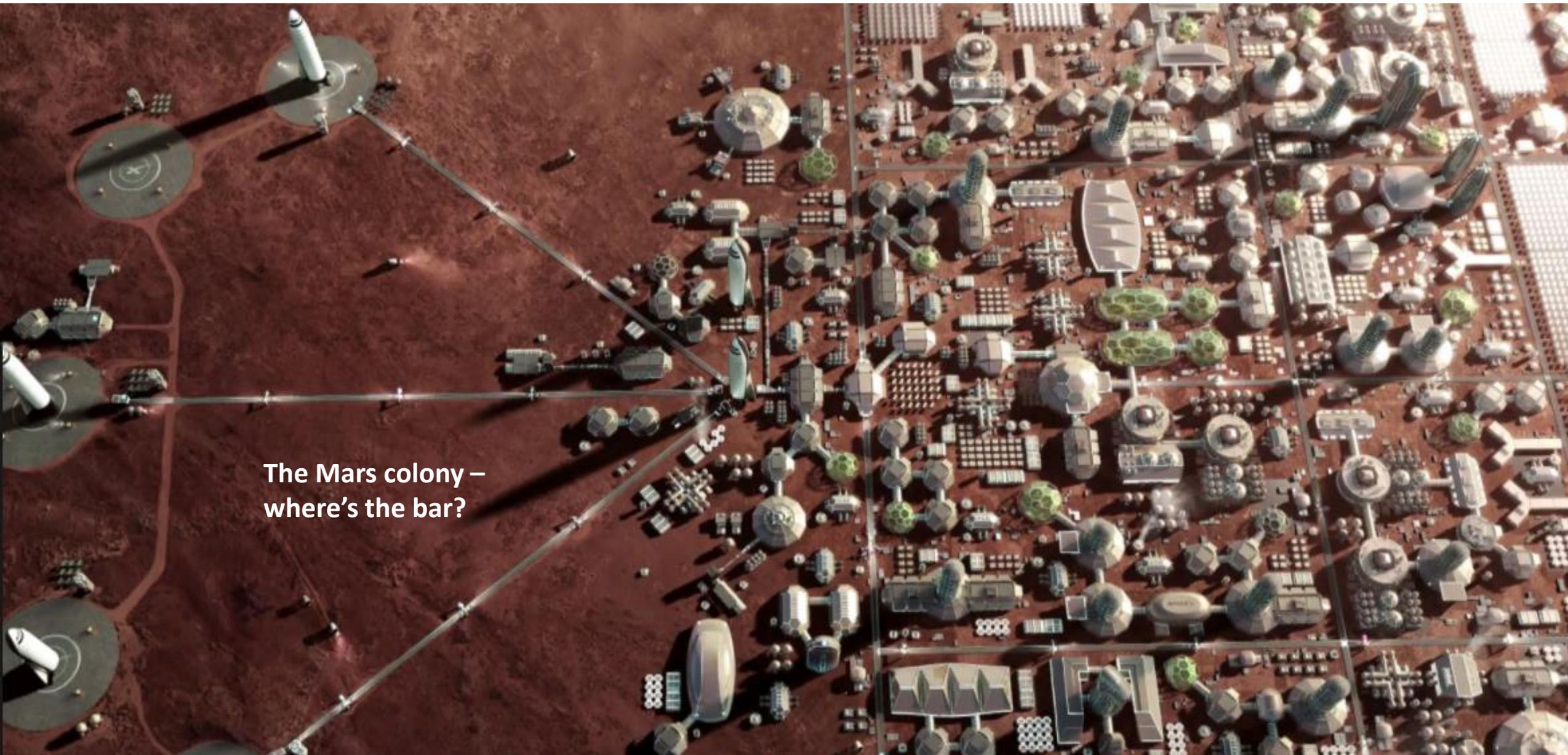
Fruitarian



Amino acids are not just used for making proteins



Important points to understand



The Mars colony –
where's the bar?



Dark beer – B vitamins



Rich in polyphenols



Yoghurt +polyphenols



Tempeh - soy



Kimchi - cabbage



Bitter beer – male hop UK



Alcohol/calories

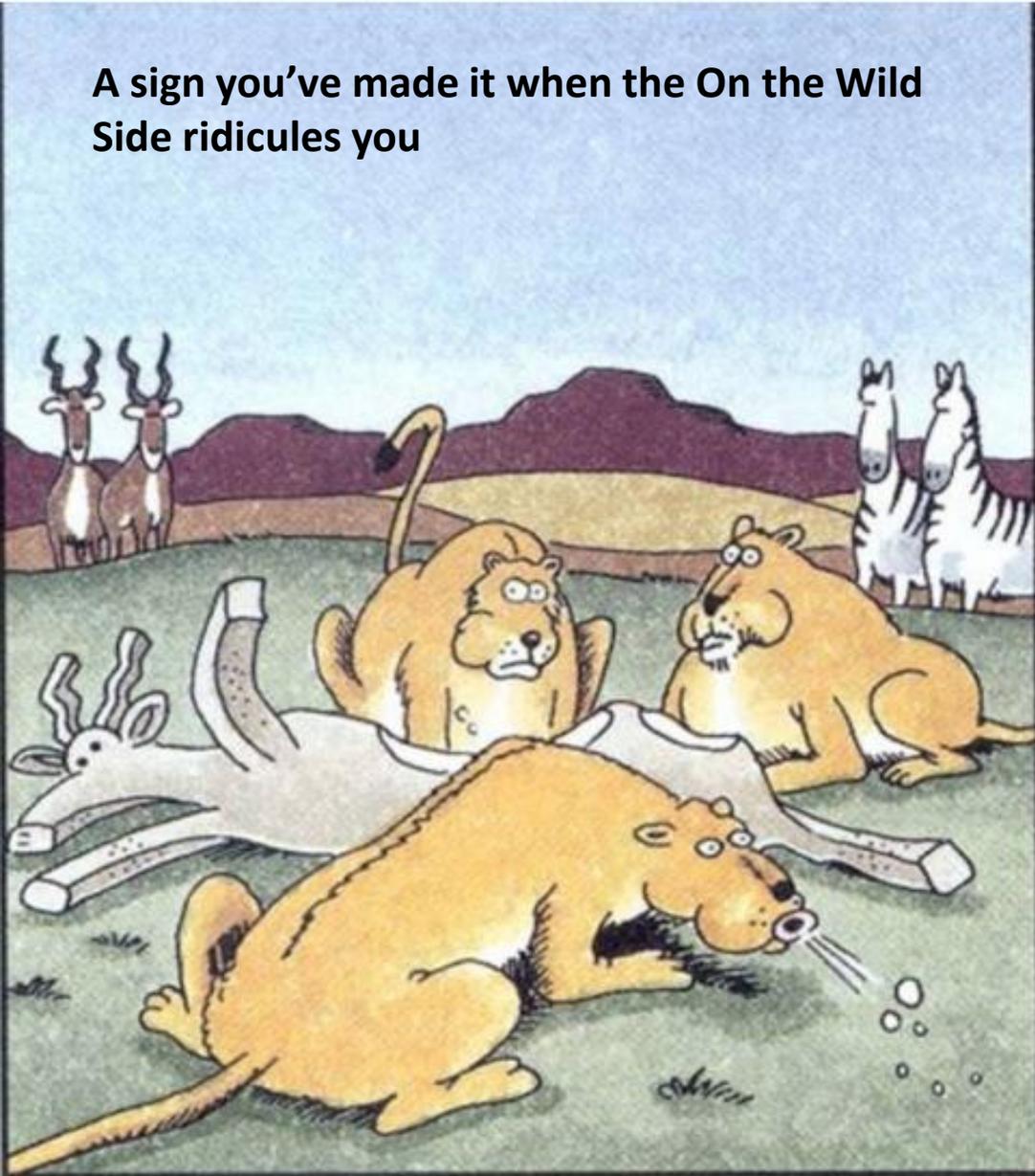


Plain, but fermented



Ogi + hibiscus flower syrup

A sign you've made it when the On the Wild Side ridicules you



In sudden disgust, the three lionesses realized they had killed a tofudebeest—one of the Serengeti's obnoxious health antelopes.

Be kind to your “cat”

Vet. Pathol. 25:48–57 (1988)

Veno-occlusive Disease of the Liver in Captive Cheetah



The main hepatic lesion was seen in 60% of the sexually mature cheetah (out of 126 captive animals). Observed in 1 year olds, but got worse with age and led to liver failure. Came from supplementation of the horsemeat diet with soy protein and the phytoestrogens therein.

Cats are exquisitely sensitive to aspirin and tylenol

- The defect is in UGT1A6 which has become a pseudogene – the WT form glucuronidates phenols (a mechanism to excrete them)
 - Cats are hypercarnivores
 - Not exposed to modern drugs or plants in which there are substantial amounts of phenols
 - Victims of “Use it or lose it”
 - Diet-driven evolution
- Mutations in exon 1
 - Stop codons at bp 274-276 and 379-381 (>10 MYA)
- UGT1A1 that glucuronidates bilirubin is unaffected

Measuring the metabolome



radiodetector

GC in oven

Paper tape
recording

Radio-GC analysis metabolomics in its infancy

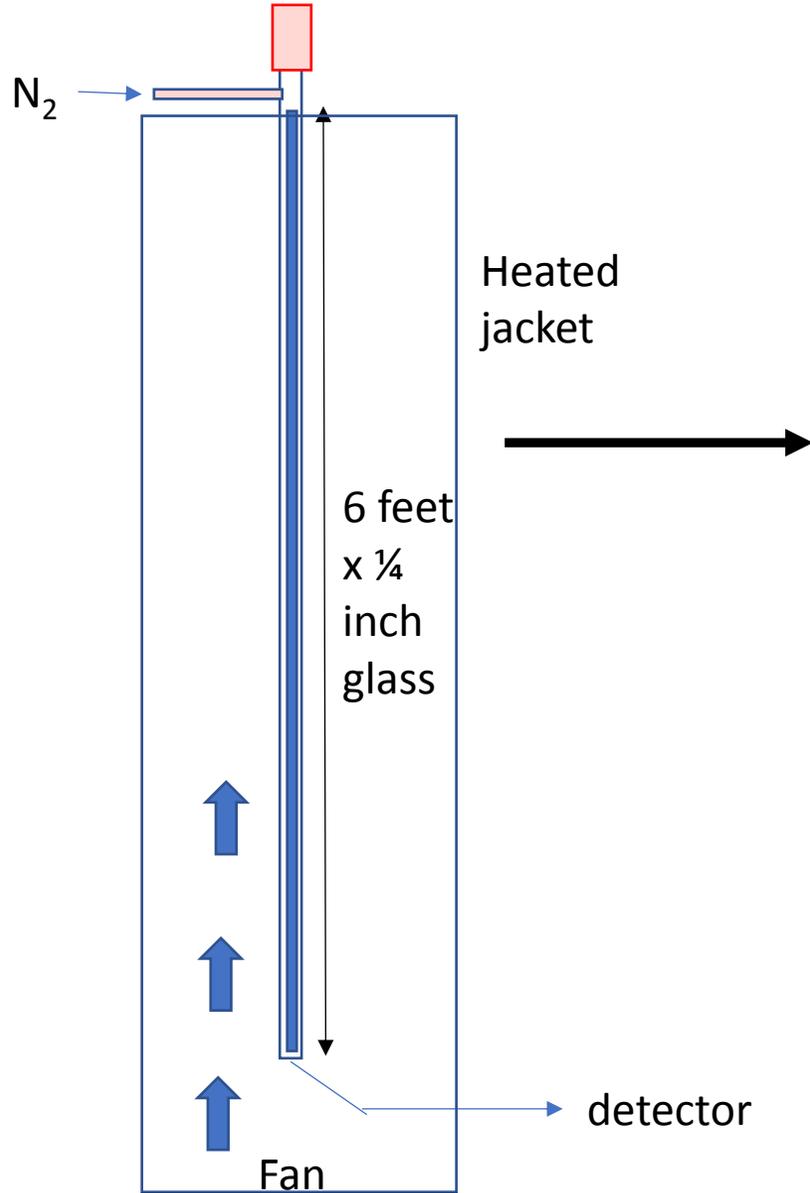
Radio gas-liquid chromatography with
digitization of collected data

Developed this for my PhD work (1967-
1970) to study glucose metabolism in
acellular slime mold, *Physarum
polycephalum*

Modern metabolomics

Evolution of GC columns

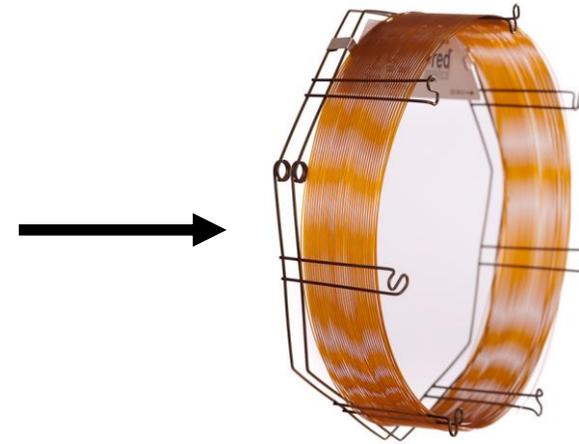
Spring-loaded clip for sample loading



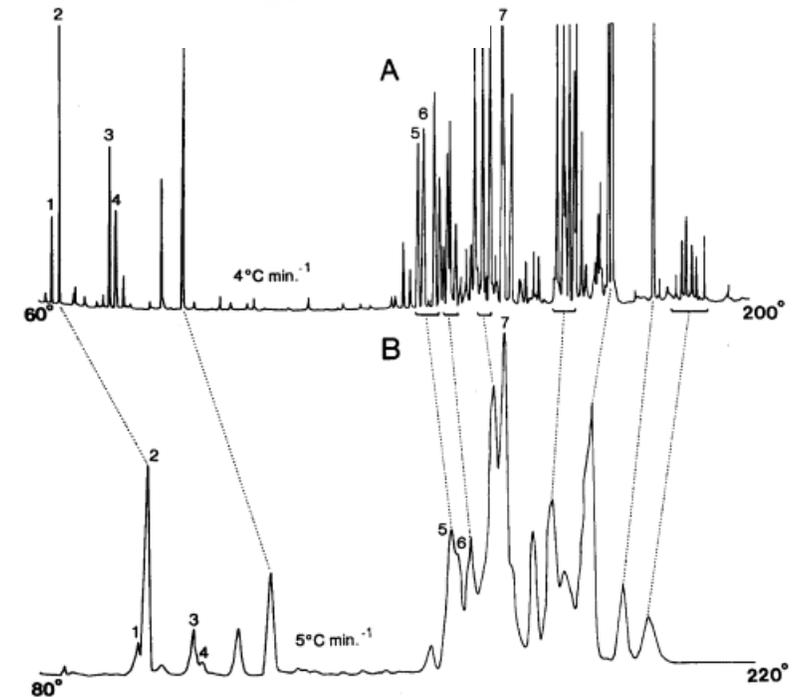
Coiled or U-shaped columns

Back pressure was a problem due to gas compressibility

Open tubular capillary (Golay) was patented in 1956. Not developed until the 1970s.



Quartz capillary GC column



HPLC

- **Its principle**

- Martin and Synge (1941).. *“the smallest HETP (height equivalent to a theoretical plate) should be obtainable by using very small particles and a high pressure difference across the length of the column.”*

- **It has several advantages over GC**

- Not necessary for the biochemical to go into the gas phase prior to separation
- The stationary phase can be modified to many different chemistries
- **The mobile phase (a liquid) is essentially non-compressible**
 - Linear flow velocity is the same at the top and bottom of the column

- **One big disadvantage**

- Smaller particles => smaller HETP & better efficiency, but => greater back pressure

- **Open tubular nanoLC?**

- Can engineering coat the walls of an extended nano-fluidics network (reproducibly!!)

$$\Delta P = \frac{\eta FL}{K^0 \pi r^2 d_p^2}$$

The diagram shows the Hagen-Poiseuille equation for pressure drop in a tube. The numerator is ηFL , where η is labeled 'viscosity', F is labeled 'flow', and L is labeled 'length'. The denominator is $K^0 \pi r^2 d_p^2$, where K^0 is labeled 'specific permeability', r is labeled 'column radius', and d_p is labeled 'particle diameter'. Green arrows point from the labels to their respective variables in the equation.

UPLC operates at 15,000 psi

Mass analyzer of choice for untargeted metabolomics

- Quadrupole-orthogonal time-of-flight (Q-TOF)



Agilent 6500



Waters Synapt
G2/HMDS



Bruker

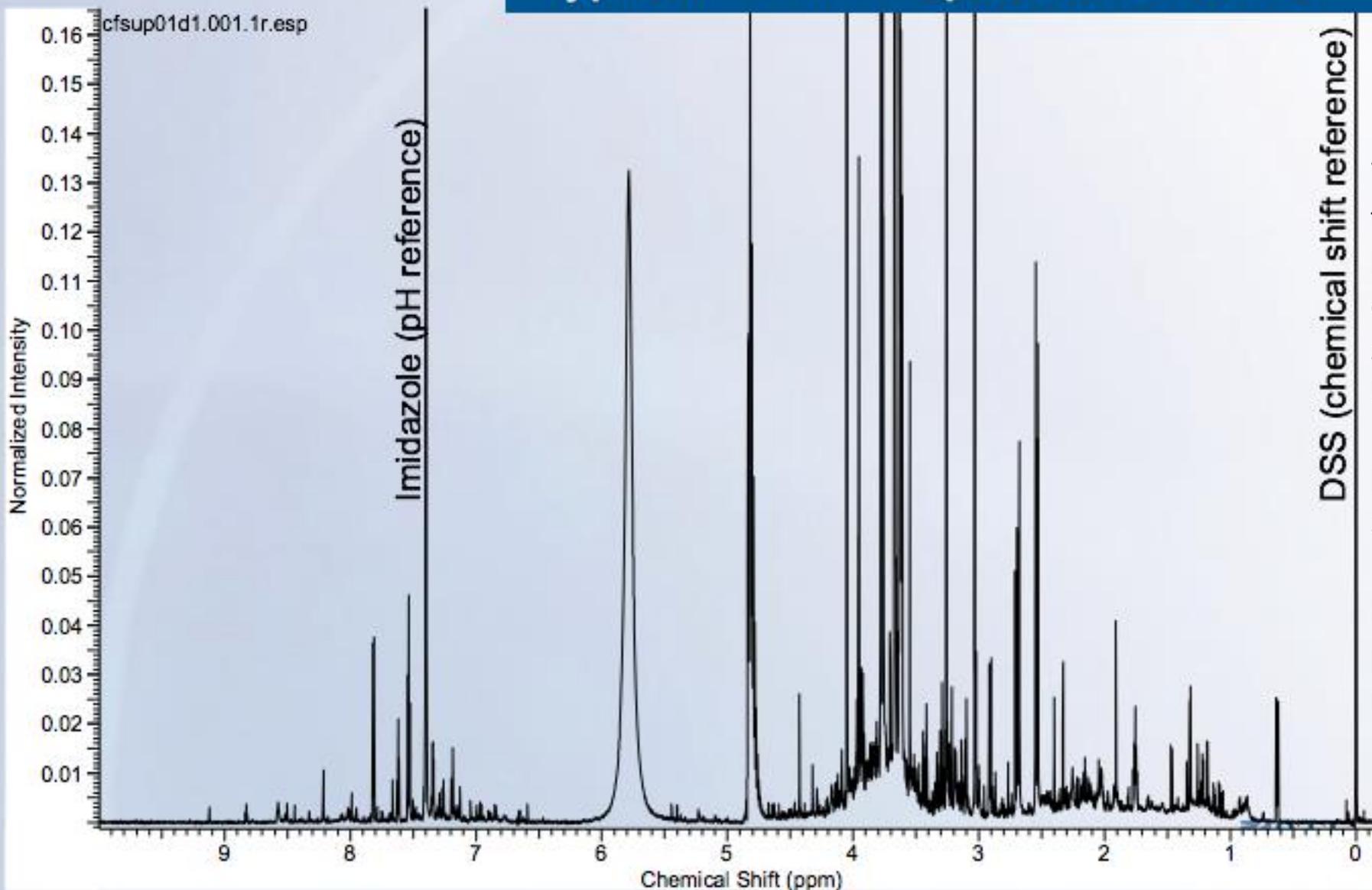


Sciex TripleTOF 6600

Current models have 40-80,000 mass resolution and 1-3 ppm mass accuracy

Nuclear Magnetic Resonance

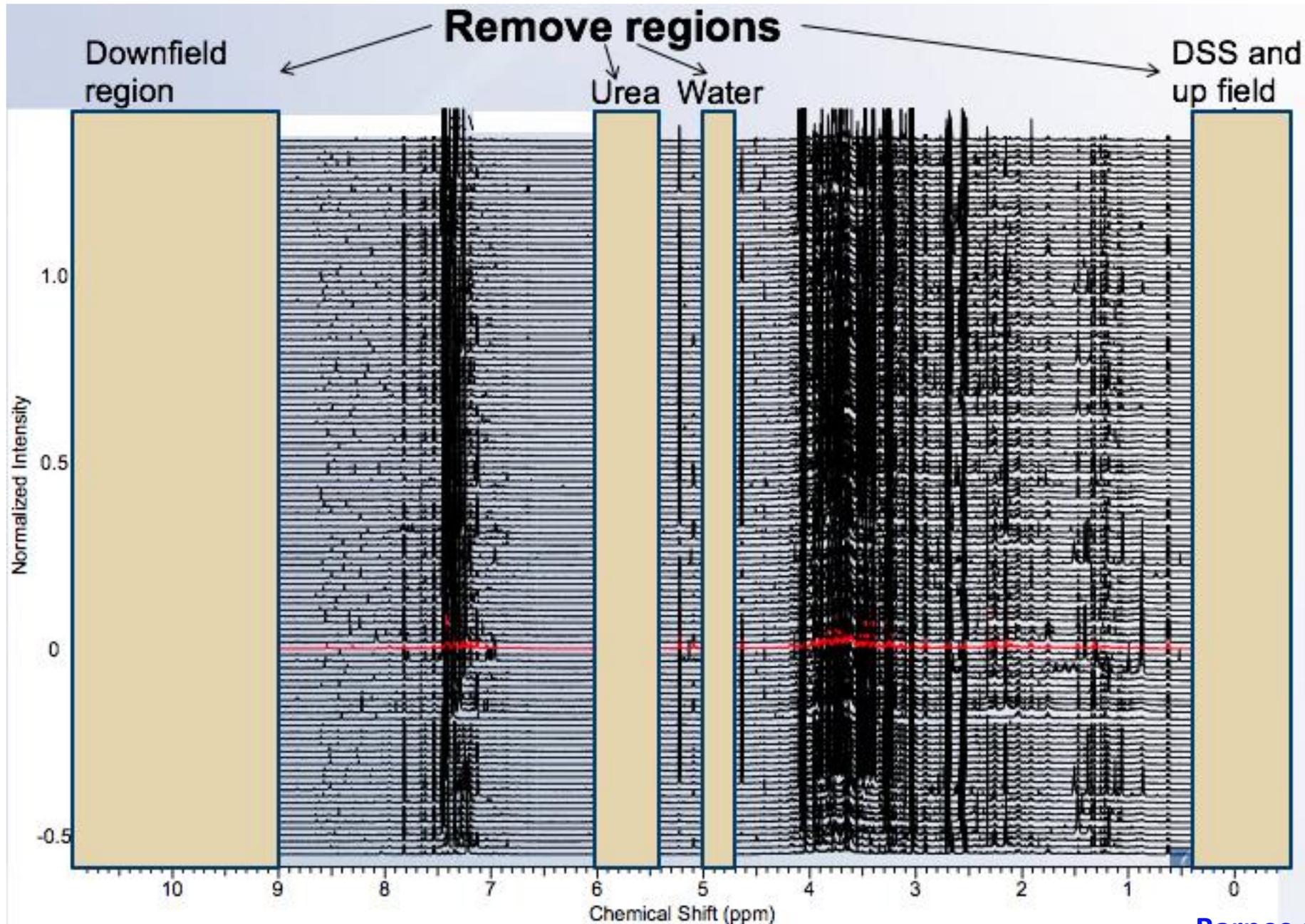
Typical ^1H NMR Spectrum of urine



Whereas I had to get rid of H_2O , these days it is not necessary. Instead, D_2O is added (10%) to calibrate the instrument. Pulse sequences are used to suppress the 55 M H_2O signal.

DSS is used as the reference signal and is set to 0.00 ppm.

Imidazole is used to determine the pH. This allows correction of proton resonances that are pH-sensitive.



NMR is **highly reproducible** and **quantitative**, but has limited sensitivity. Measurements made at one institution or machine can be reproduced (after proper calibration) at another another institution or machine type.

The NMR signal is spectroscopic and does not change.

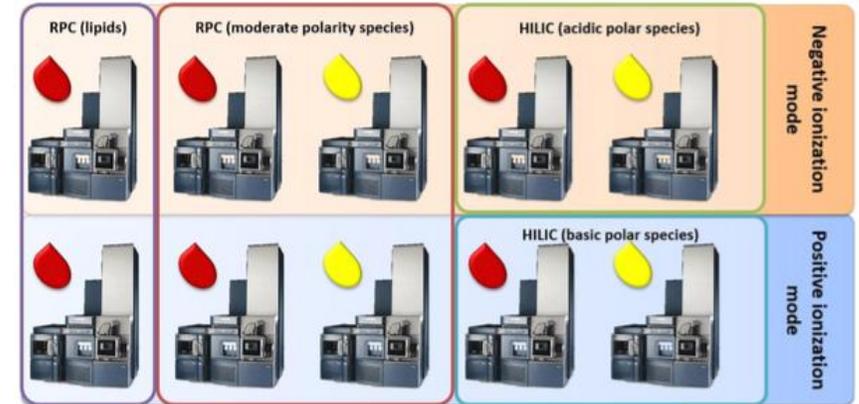
MRC-NIHR National Phenome Centre

Imperial College, London

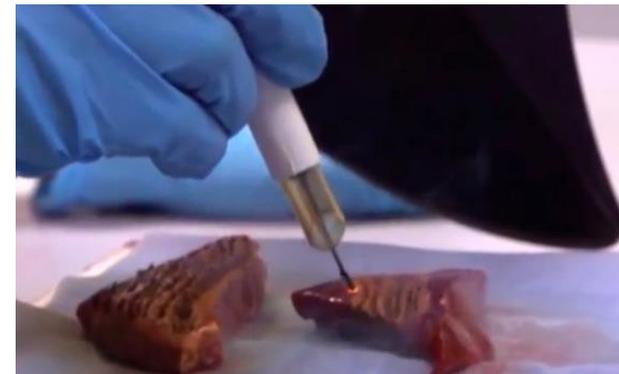


600 MHz NMR instruments
in surgical suite

This is Next-GEN
precise medicine
for everyone



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



Iknife - revolutionizing surgery

The UK National Phenome Center, LC-MS labs



UAB capabilities in metabolomics

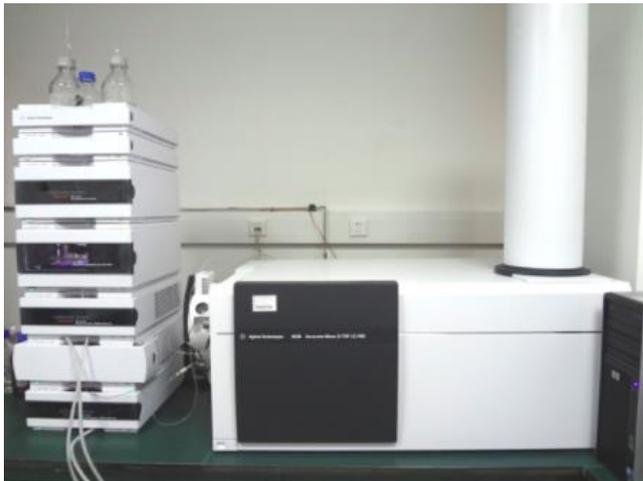


**SCIEX 5600 TripleTOF
with Eksigent nanoLC
Research**

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



SCIEX 6500 Qtrap with SelexION



**Agilent 6530 QTOF???
For Clinical Metaboomics**



**Central Alabama NMR facility
Chemistry Bdg
William Placzek, PhD, Director
Chad Petit, PhD
205-934-2465**

NIH Common Fund Metabolomics Program



METABOLOMICS WORKBENCH

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Welcome to the UCSD Metabolomics Workbench, a resource sponsored by the Common Fund of the National Institutes of Health.

NIH Metabolomics Data Repository

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Quick Links - Key Resources

EVENTS CALENDAR

- updated 02/25/2018

Metabolomics News

02-11-2018 - **Research Associate for Metabolomics Research** - The Sumner Lab at the University of North Carolina at Chapel Hill Nutrition Research Institute is seeking an individual who is highly skilled and experienced in mass spectrometry methods to serve as a Research Associate for metabolomics research. The Research Associate will work as an integral part of the Sumner Lab to conduct metabolomics investigations in nutrition, environmental exposures, and metabolic variation. The Research Associate will acquire, analyze, and interpret findings using mass spectrometry, software for statistical and multivariate analysis, and pathway mapping. Additionally, the individual should have a good understanding of quality control procedures using in mass spectrometry. The Sumner Lab is located at the University of North Carolina at Chapel Hill Nutrition Research Institute on the North Carolina Research Campus - just 20 minutes from the Charlotte, NC.

Additional details regarding the position are located at <https://unc.peopleadmin.com/postings/134864>. To apply for the position, please click on "Apply for this job" link that is located near the top of the web page.

Regional Comprehensive Metabolomics Resource Cores (RCMRC)s

[Michigan Regional Comprehensive Metabolomics Resource Core \(MRC\)²](#)

[NIH West Coast Metabolomics Center at UC Davis](#)

[Eastern Regional Comprehensive Metabolomics Resource Core \(ERCMRC\)](#)

[Southeast Center for Integrated Metabolomics \(SECIM\)](#)

[Resource Center for Stable Isotope-Resolved Metabolomics \(RC-SIRM\)](#)

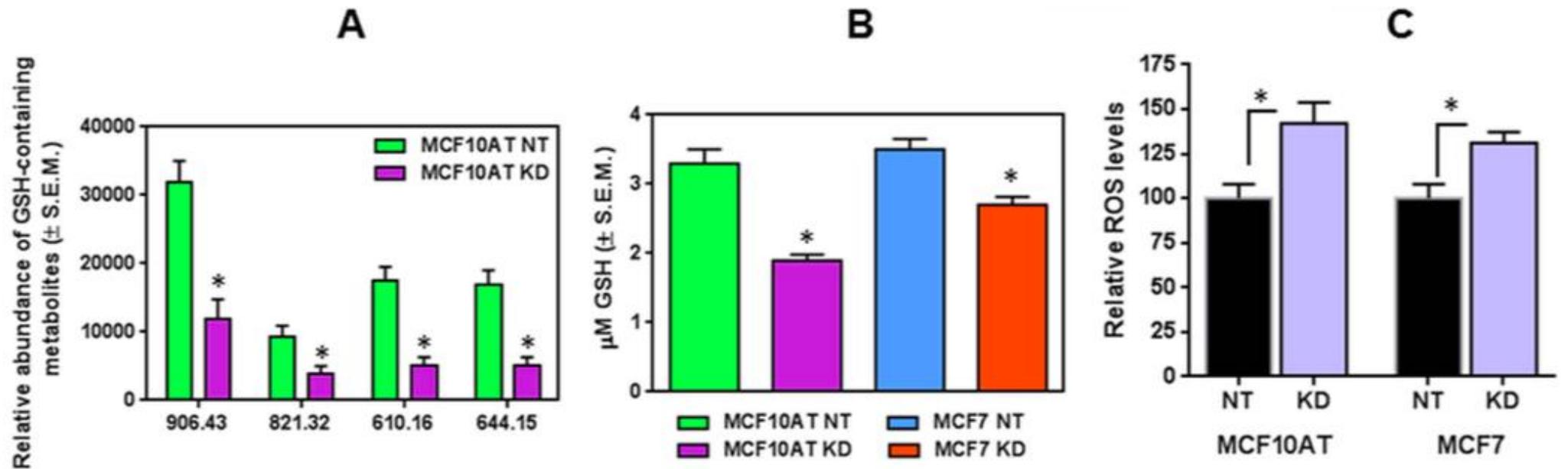
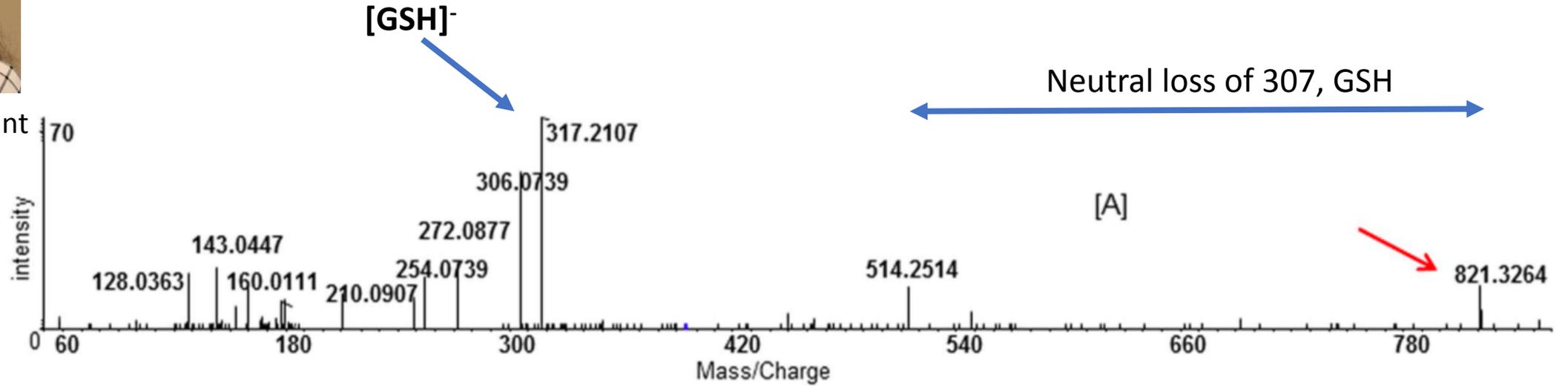
<http://www.metabolomicsworkbench.org/>

Examples of metabolomics applications

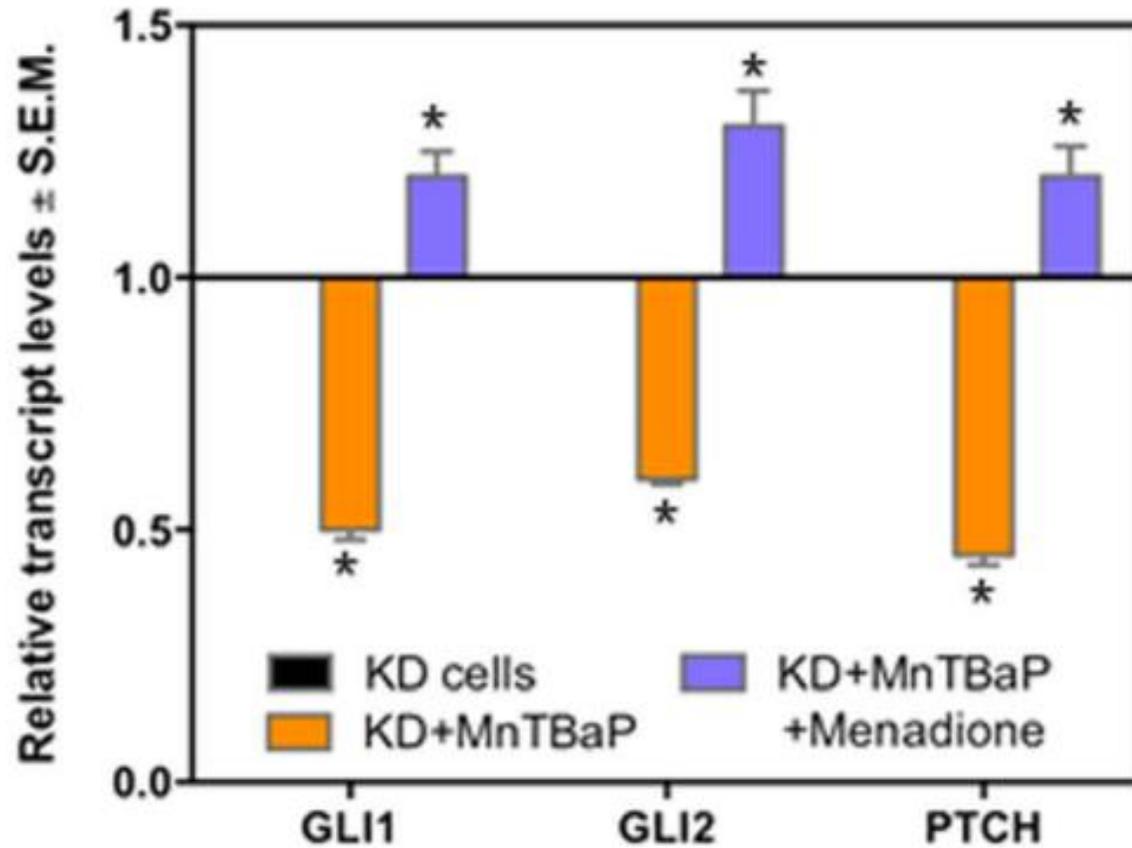


Effect of loss of Merlin in MCF10 cells

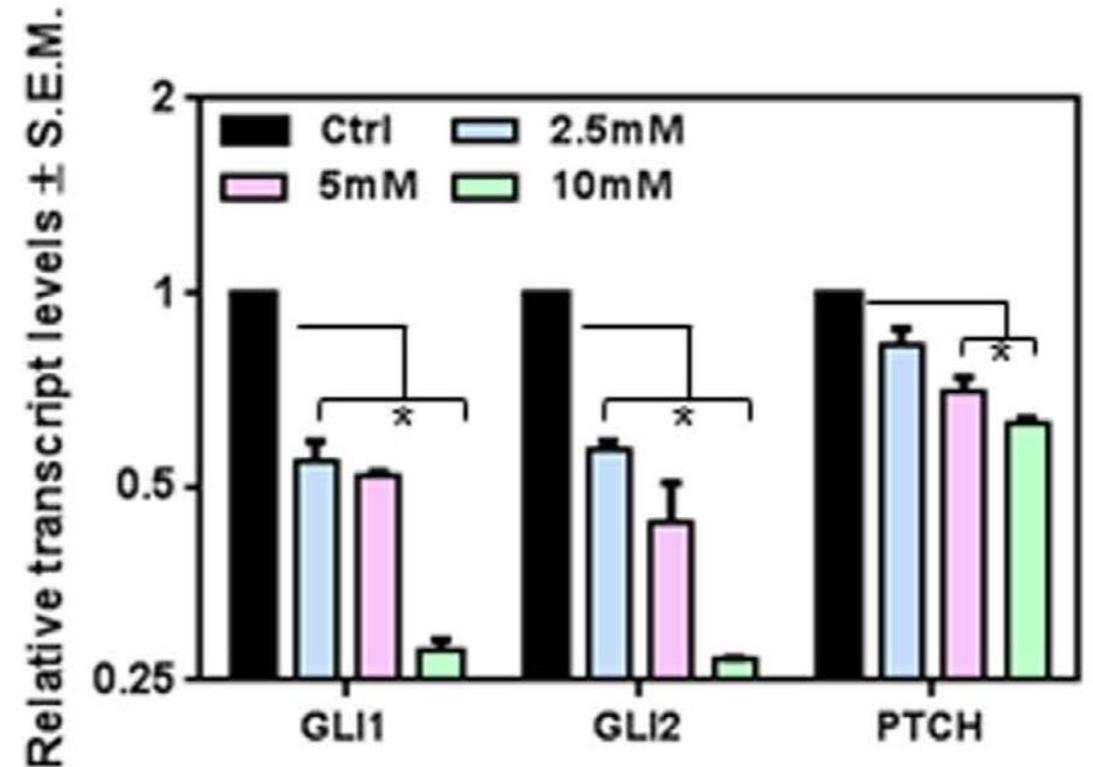
Lalita Shevde-Samant



Loss of GSH leads to activation of sonic hedgehog signaling



Menadione induces ROS



GSH restores the suppression of Hgh signaling



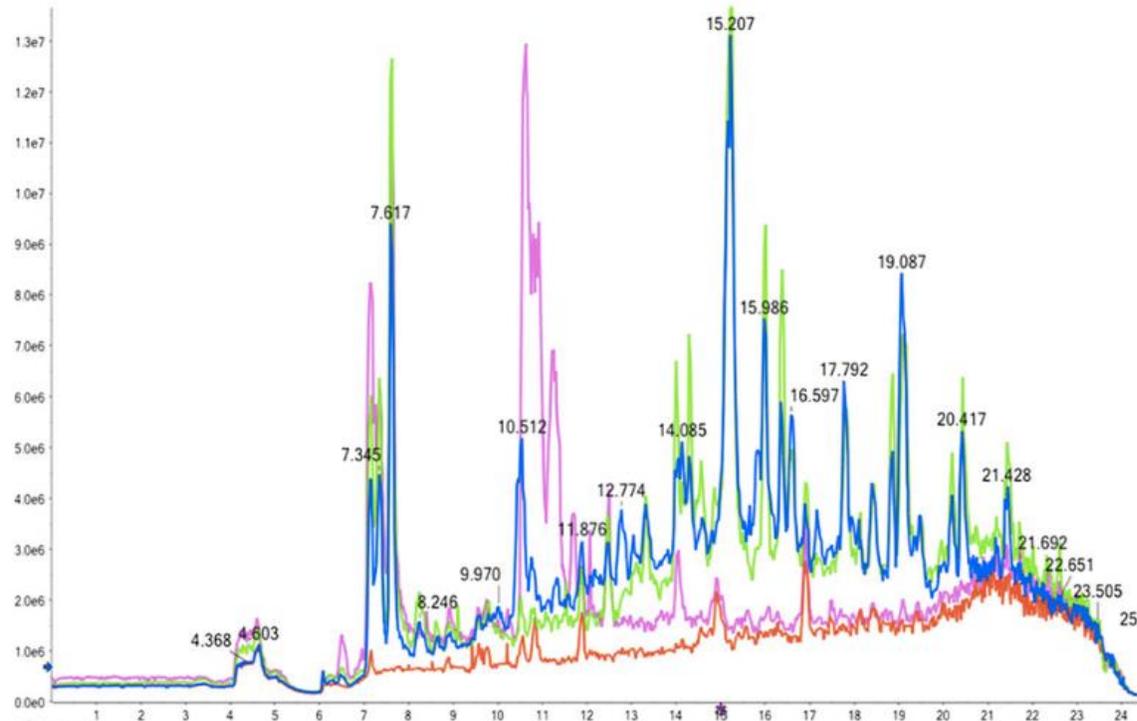
Fecal metabolites (!) in children with spondyloarthritis ± immunosuppression

- **Challenge**

- Transferring metabolites in feces in 20% glycerol to a microfluidic network ending in a 5 μ m diameter spray needle

- **Method**

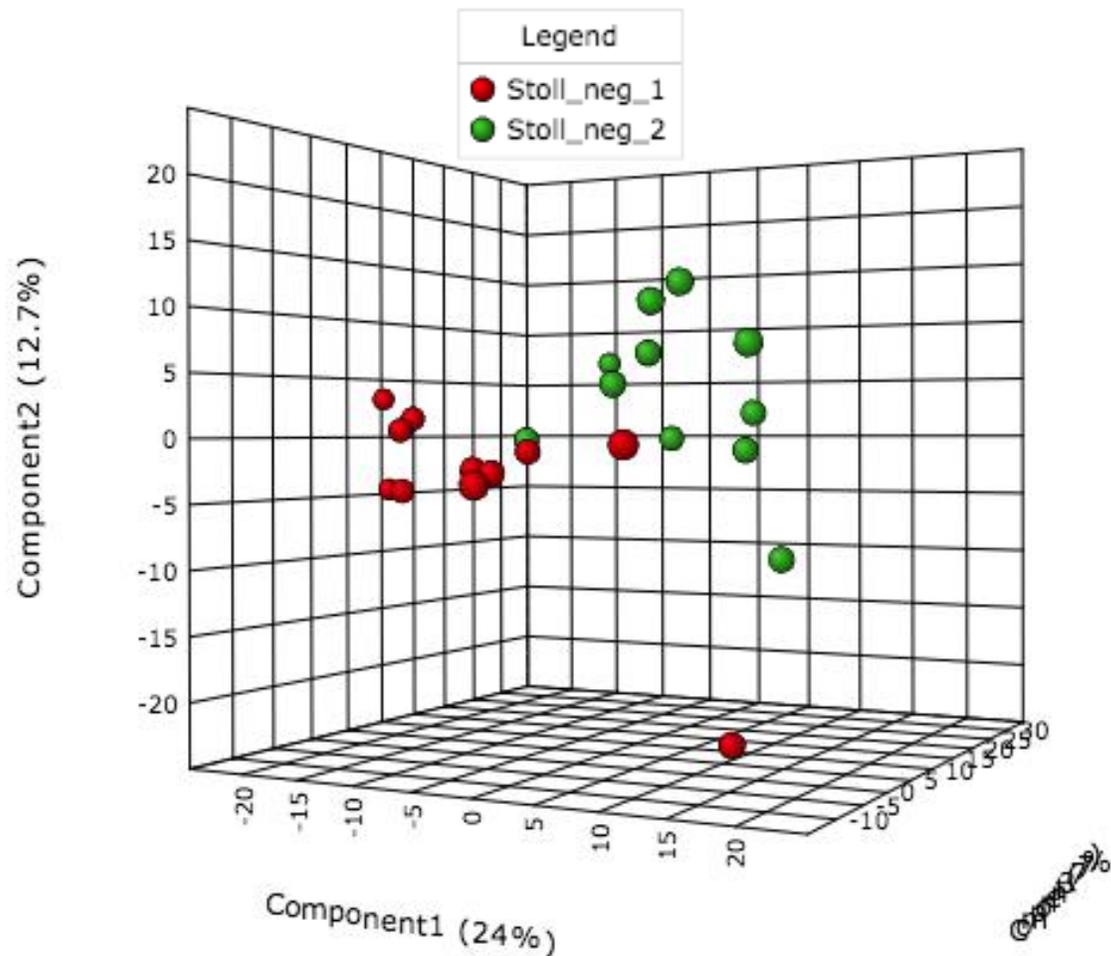
- Disperse 100 mg feces in 0.1% formic acid
- Centrifuge at 15,000 g to get “fecal water”
- Extract the fecal water with ethyl acetate



- Purple trace – 0.1% formic acid
- Green trace – neutral pH (water)
- Blue trace – 150 mM NaOH
- Red trace - Blank

It worked!!

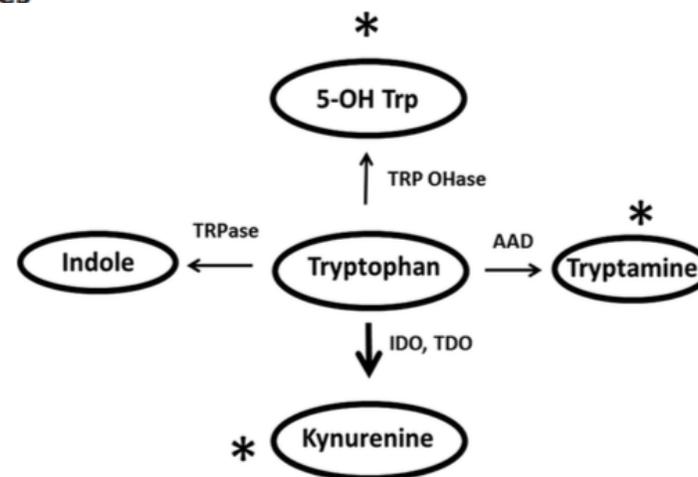
Results of the study



3D-Partial Least Squares Discriminant Analysis

Table 2. Pathways under-represented among patients in cohort 1

Pathway	Overlap size	Pathway size	P-value
<i>Negatively charged ions</i>			
Glycosphingolipid biosynthesis—ganglioseries	5	7	0.00091
Tryptophan metabolism	13	46	0.00106
Glycosphingolipid biosynthesis—globoseries	3	3	0.00122
Glycosphingolipid biosynthesis—hexosamines	3	3	0.00125
<i>Positively charged ions</i>			
Tryptophan metabolism	13	46	0.0038
Xenobiotics	3	3	0.00544

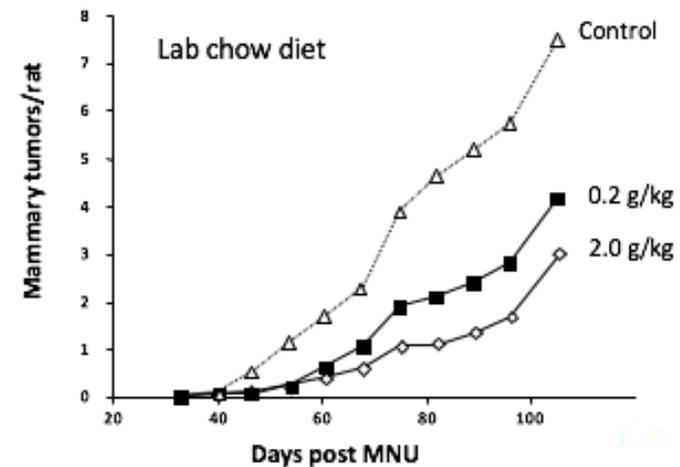


Tryptophan is the source of ligands for the aryl hydrocarbon receptor



Rat model of mammary carcinogenesis – originally animals fed **chow** diets – tumors induced by a single injection of MNU (Clinton J. Grubbs).

Chemopreventive agent evaluation – pre-1985 to now. **Very reproducible**. Barnes & Grubbs (1990) showed that removal of soy from the diet substantially increased tumors. Genistein added to the diet reversed tumor induction, but only if rats received soy early in life.



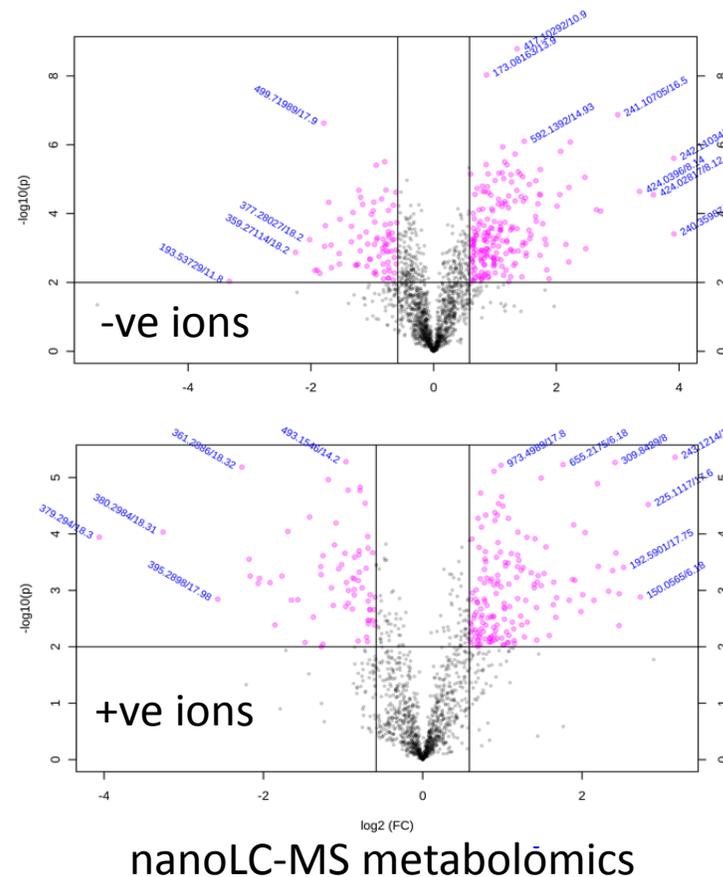
2013 – the diet (NIH-7001) was treated to control for microbial growth with Co-60 γ -radiation. **Unexpectedly, tumor number was reduced by 40%!!**

Experiment repeated with one batch of NIH-7001 that was split into two, one of which was exposed to the Co-60 γ -radiation. **Again, tumor number was reduced (by 38%).**

Questions:

1. Is the fecal microbiome altered? **YES**
2. Is the urinary metabolome altered? **YES**

More fundamental, did the Co-60 γ -radiation alter the diet?
ANSWER, yes, big time



possible identity *
oxo-2-nonenal
hydroxy-oxo-nonenal
:2-oxo
:2-monoOH (HODE)
:2-diOH
:0-diOH ester of 8:1-diOH
:1-diOH ester of 8:1-diOH-oxo
:1 ester of 18:2-diOH

identified by MSMS

17 Aug;52(8):507-516

Future: the metabolome of a patient

- **Metabolomics on urine/plasma/serum can assess:**
 - From the pattern of human (and perhaps microbial) metabolites, how does the patient's metabolome change during different stages of their disease process (acute and recovery)
- **Medications**
 - Are they taking their medication and is it the correct medication?
 - What other medications are they taking (prescribed, antibiotics, OTC, *other*)?
 - What beverage did they drink last?
 - Are they consuming unusual foods/dietary supplements?
- **What is the subject's metabolic age?**

Advanced metabolomics

Single cell analysis – Peter Nemes, PhD (U. Maryland)

http://www.uab.edu/proteomics/metabolomics/workshop/2017/videos/nemes1_day2.html



**The iKnife – precision surgery (metabolomics) on the operating table –
Mr. James Kinross, PhD, FRCS (Imperial College, London)**

http://www.uab.edu/proteomics/metabolomics/workshop/2016/videos/kinross_day2.html

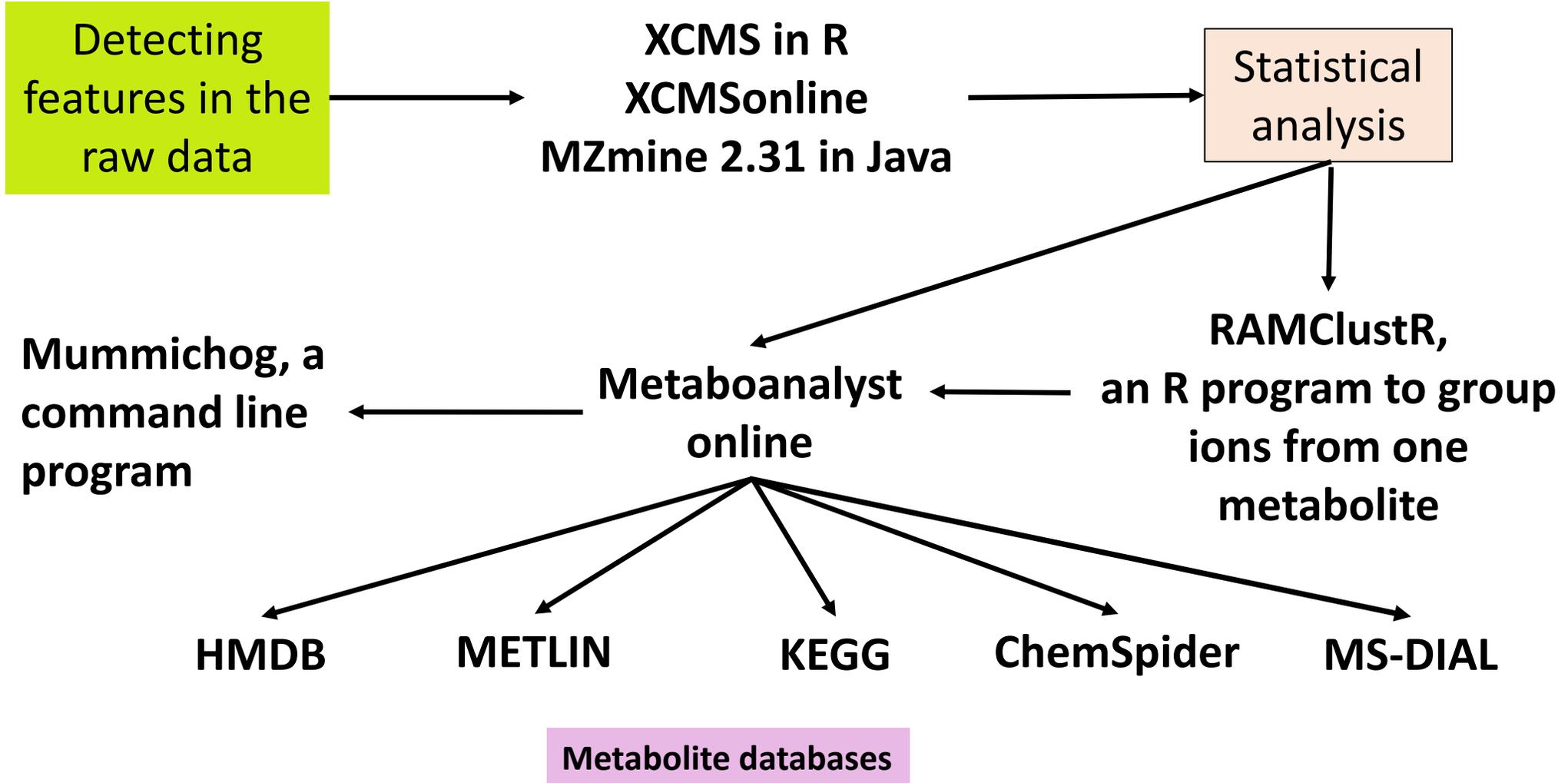


Tissue imaging metabolomics – Janusz Kabarowski, PhD (UAB)

http://www.uab.edu/proteomics/metabolomics/workshop/2017/videos/kabarowski_day4.html

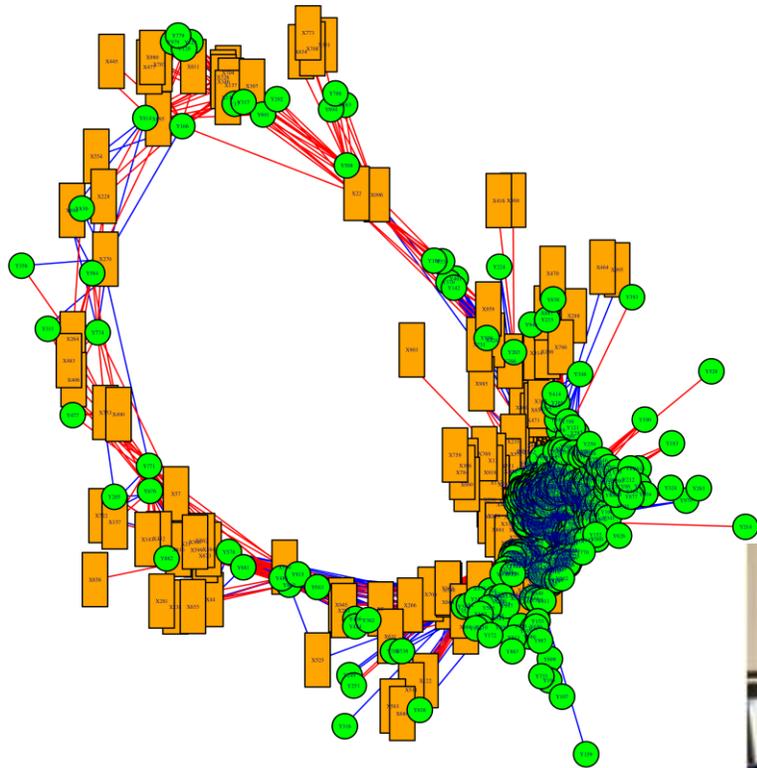


Data analysis



Network Analysis using xMWAS

Using all samples



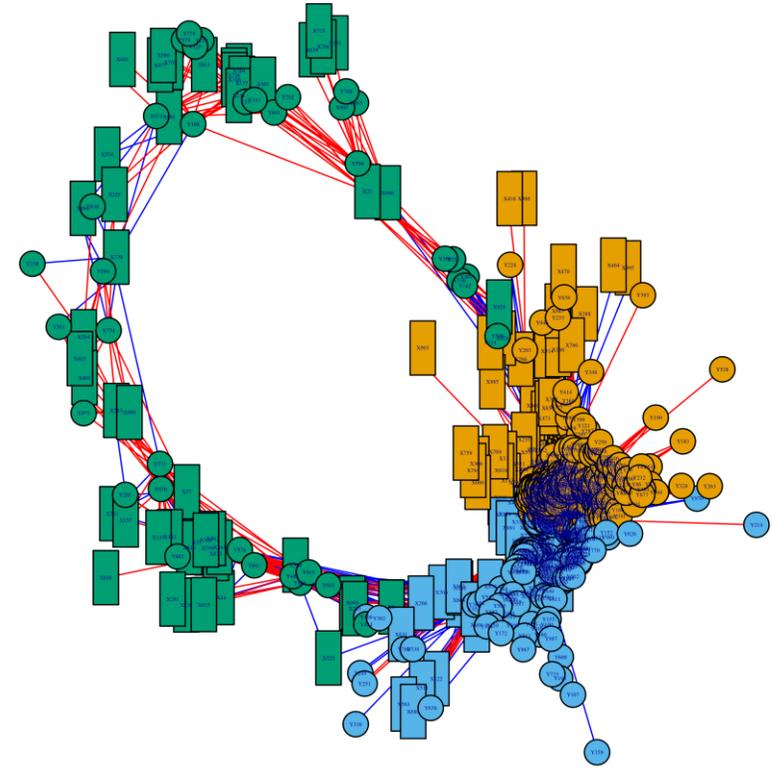
(Edges) Red: +ve correlation; Blue: -ve correlation
(Nodes) Rectangle: Positive; Circle: Negative

Multidata_Network_threshold0.9.png

(Edges) Red: +ve correlation; Blue: -ve correlation

(Nodes) Rectangle: Positive; Circle: Negative

Using all samples



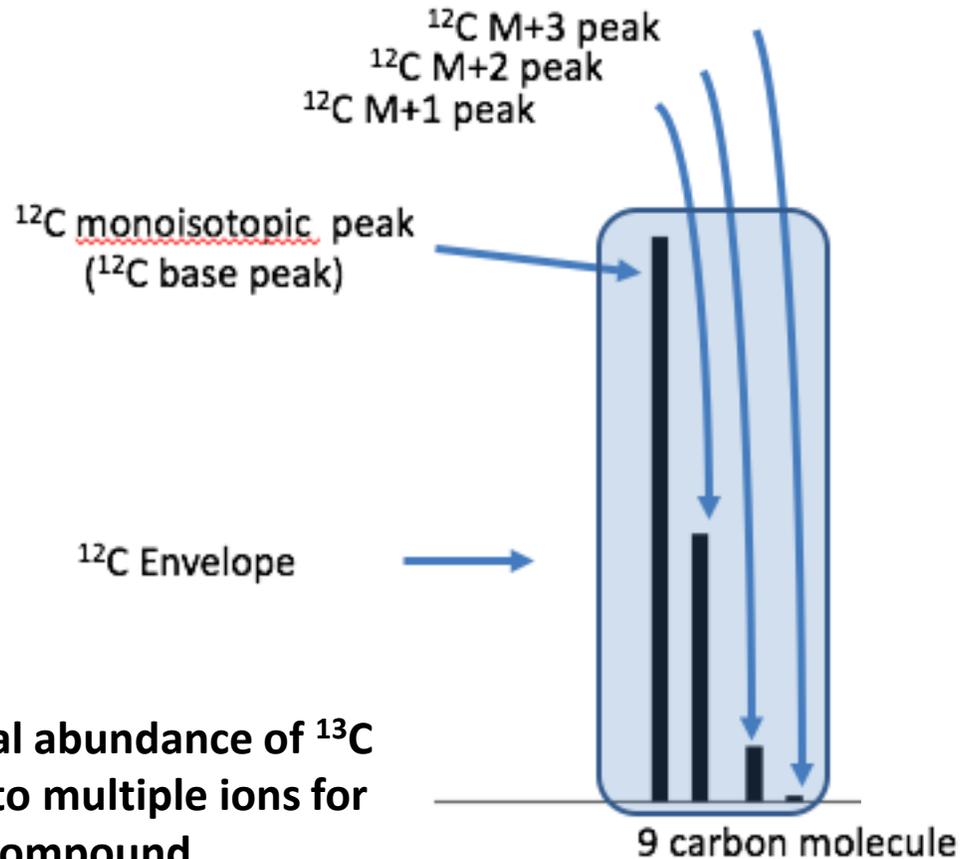
Each community (C) is represented by a different color:
C1; C2; C3;

Multidata_Network_threshold0.9_communities.png



Uppal K, Ma C, Go YM, Jones DP, Wren J.
Bioinformatics. 2018 Feb 15;34(4):701-702.

Isotope ratio outlier analysis to facilitate identification and reproducibility in metabolomics



Natural abundance of ^{13}C leads to multiple ions for each compound

This reagent is added to all samples
– allows high quality QA/QC

Growing yeast on 95% ^{13}C -glucose leads to all metabolites with this distinct isotope signature and no ^{12}C signal

Key issues in metabolomics

- **Design the experiment well**
 - Discuss with a statistician before starting
- **Collect samples as best you can – better fresh**
 - Ideally, have no hemolysis in blood samples
 - Tissue samples should be frozen at -80°C ASAP
 - In animal experiments, flush the tissue to be excised with ice-cold PBS, then freeze clamp
 - For cells, decant medium, flush dish with ice-cold PBS to remove extracellular components (10 s), and then add methanol cooled in dry ice
- **Some samples will have already been collected**
 - Learn as much as you can about how the sample was handled and stored

Key issues in metabolomics-2

- **Numbers of samples per group** (to develop hypotheses)
 - Cells 3-5
 - Mice/rats 10-12
 - Patients 20-25 (controlled study)
 - Patients 100-500 (Epidemiologic or uncontrolled study)
- **The numbers needed to test hypotheses depend on the variance observed in the preliminary study** (work with a statistician to evaluate this)
 - Stan Hazen, discovering trimethylamine N-oxide in patients with adverse cardiovascular risk, chose a wide range of risk and carefully matched the risk patients with healthy controls (avoid antibiotics and other medications)
 - His initial study had **50** patients and controls – a validation study had **25** per group

Key issues in metabolomics-3

- **How much will it cost?**

- A standard approach is to (1) extract the biological material, (2) carry out nanoLC-MSMS (negative and positive ions) and (3) process the data

Extraction

\$12.50 per sample

Nano-LC-MSMS

\$200 per sample

Data analysis

\$400 per study

Examples

- 2 groups of cells (n=5)
 - $2 \times 5 \times \$12.50 + 10 \times \$200 + \$400 = \$2,525$
- 2 groups of animals (n=10)
 - $2 \times 10 \times \$12.50 + 20 \times \$200 + \$400 = \$4,650$
- 2 groups of patients (n=25)
 - $2 \times 20 \times \$12.50 + 50 \times \$200 + \$400 = \$10,525$

Summary

- **Metabolomics (integrated metabolism and chemistry of living cells) has had a long history and depends on the ability to separate, recognize and quantify individual components**
 - Its development has depended on engineering and micro/nano system innovations as well as computational development
- **Metabolomics is an important aspect of the overall research on the functions that control life (along with other –omics research) and is an important adjunct to current precision medicine**
- **UAB is building a considerable experience in metabolomics – analytical resources have moderate capacity – need to be expanded**

Acknowledgements

- AT (Tony) James
- Sir Ernst Boris Chain
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S10 RR027822



Landon Wilson



Taylor Berryhill



Mikako Kawai

Thank you - Questions?