

From Profile to Mechanistic Understanding

Ian D Wilson

Imperial College London

i.wilson@imperial.ac.uk

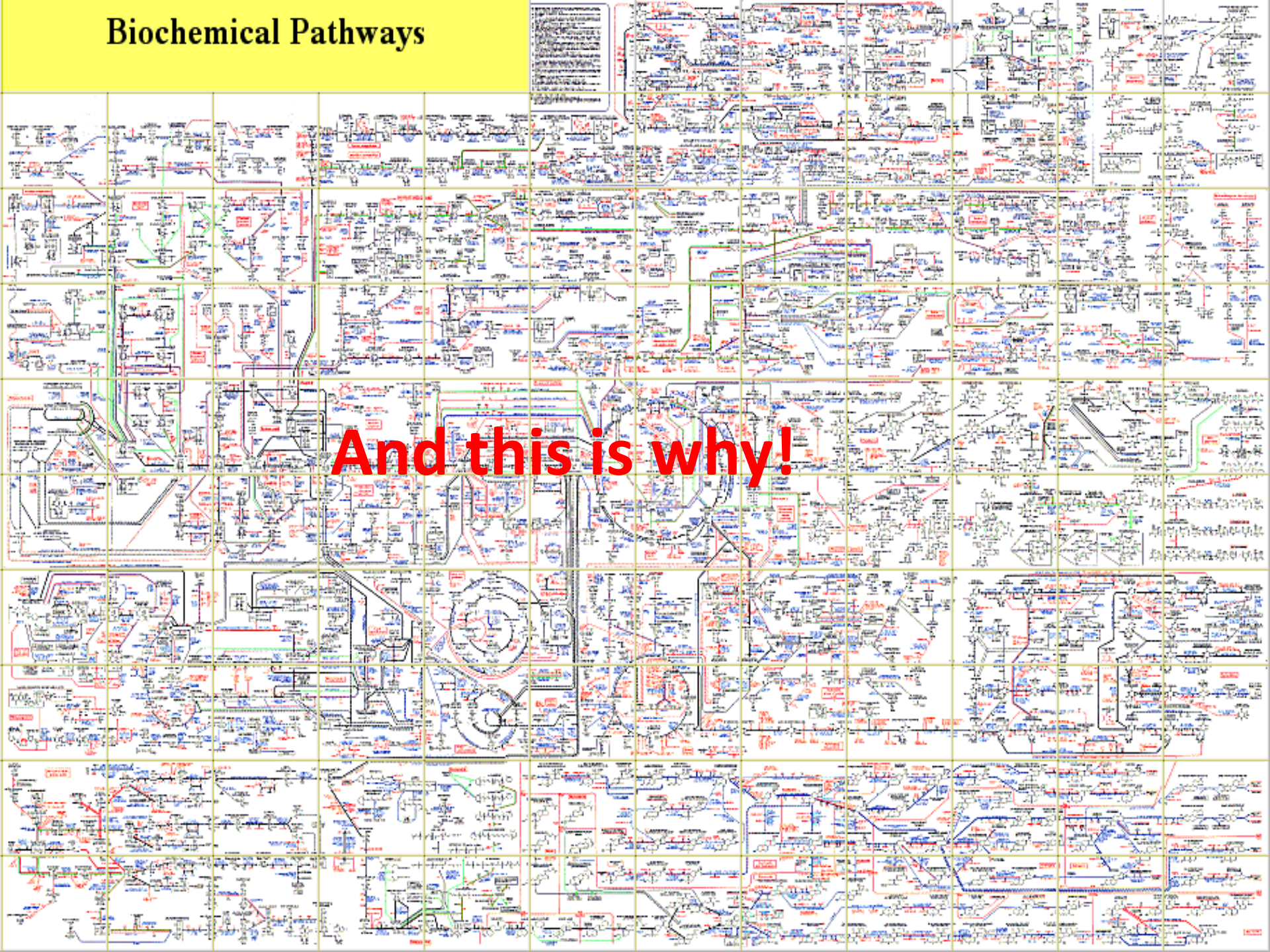


Metabolic Phenotyping

- Metabonomics/metabolomics – **untargeted** (and intended to be **unbiased**) metabolic profiling of biological samples
- The aim is to find “**biomarkers**” in areas such as basic biology, disease, toxicity etc.
- Ideally these should be **mechanistic** and **specific** for the condition under investigation
- Often they are not.....

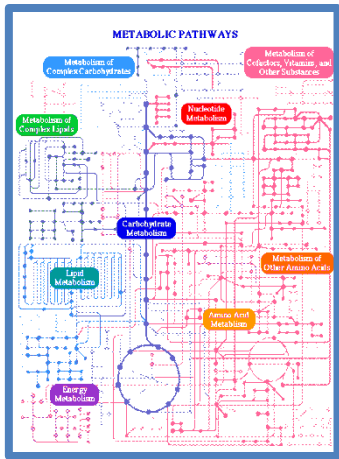
Biochemical Pathways

And this is why!

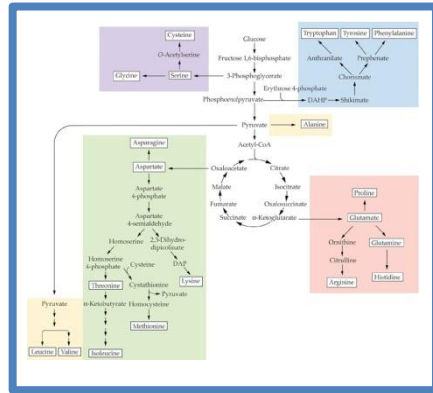


From Metabolic Phenotyping to Biomarkers

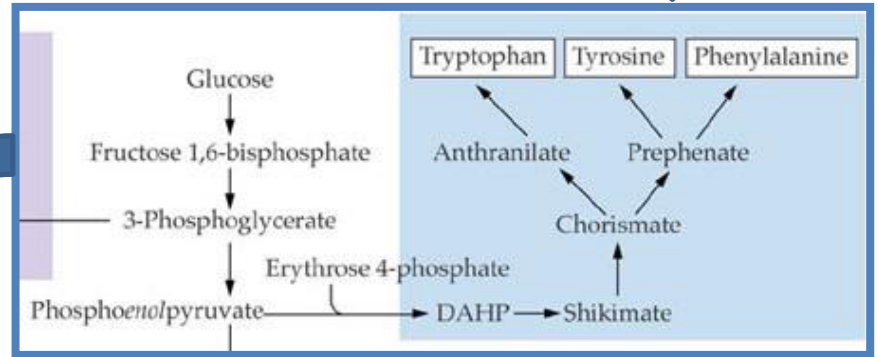
Global profiling – unbiased biomarker **Discovery**



Targeted classes of compounds - **Quantification**



Targeted pathways - **Confirmation**



Specific metabolites - **Application**

A Rodent Model of Alcoholism

- **A rodent model of alcoholism, mice/rats fed intra-gastrically with ethanol.**
- **Urine and liver extracts analysed by UPLC-MS with PCA for “biomarker” detection.**
- **What do we see? And what does it tell us?**

Alcoholic rodents

Output.M1 (PCA-X)
t[Comp. 1]/t[Comp. 2]

URINE

Control mice

Alcoholic mice

Control rats

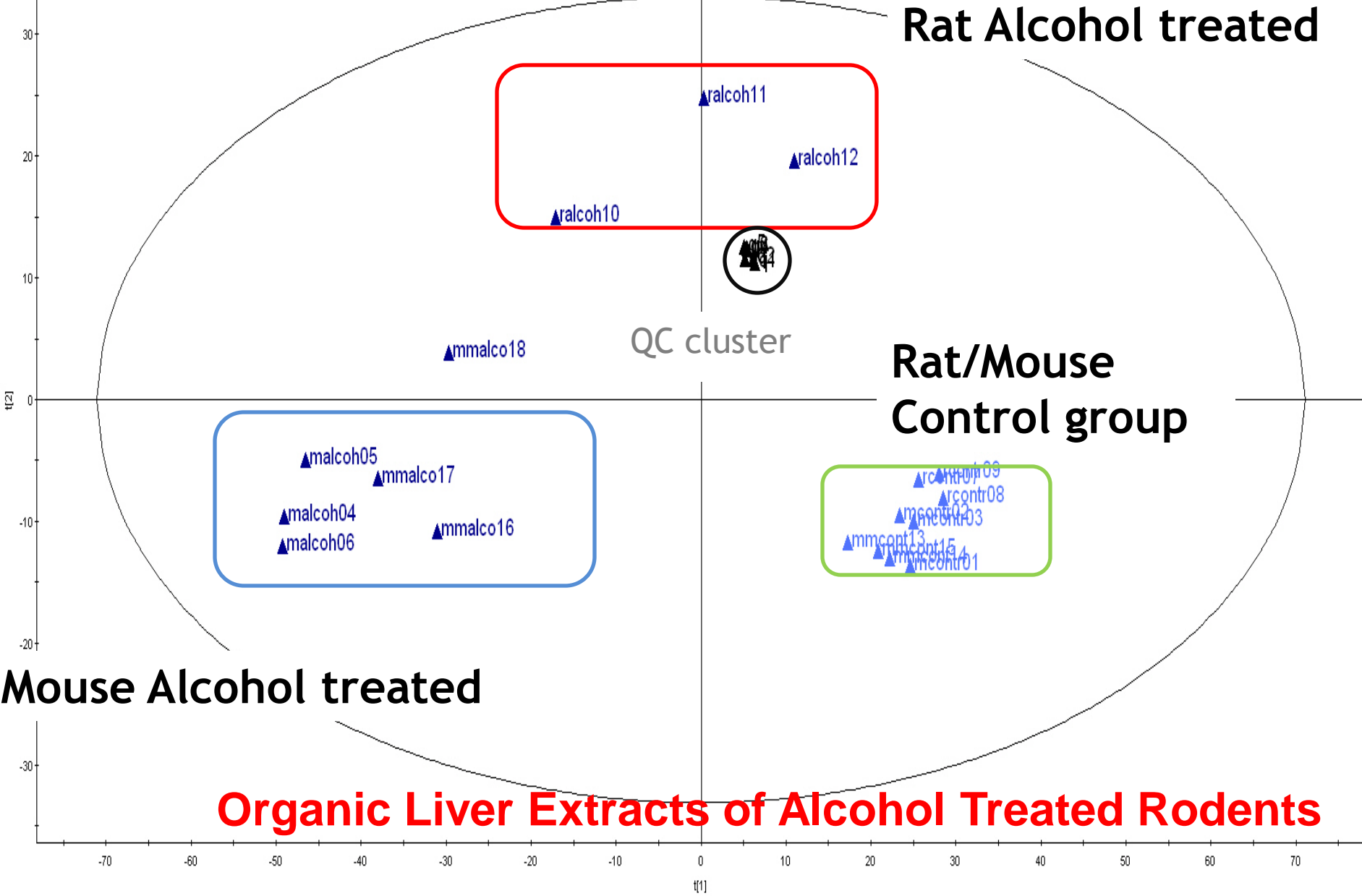
QCs

Alcoholic rats

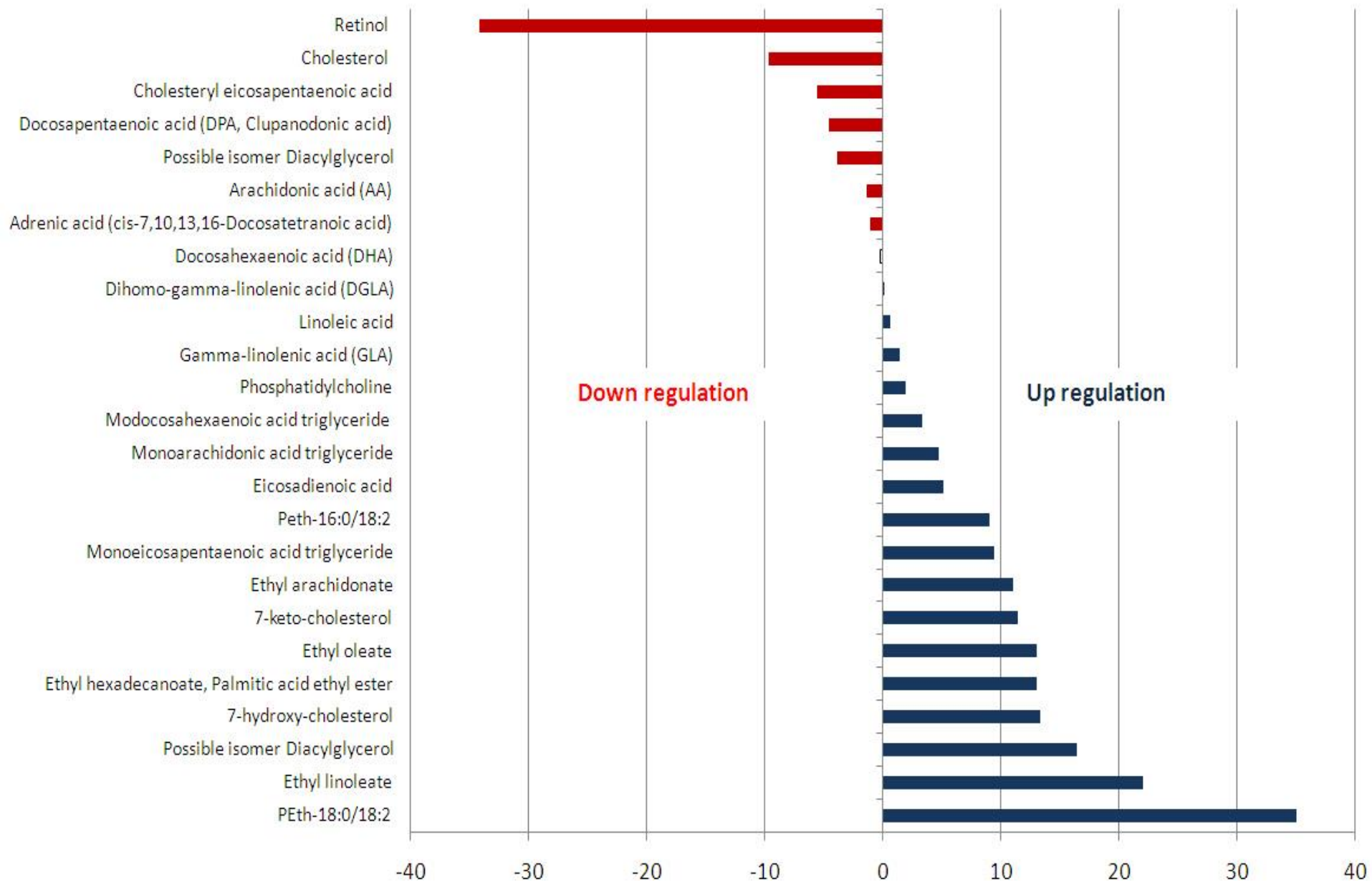
tryptophan metabolites 5-hydroxytryptophan & xanthurenic acid

R2X[1] = 0.54774 R2X[2] = 0.158417
Ellipse: Hotelling T2 (0.95)

Alcoholic rodents



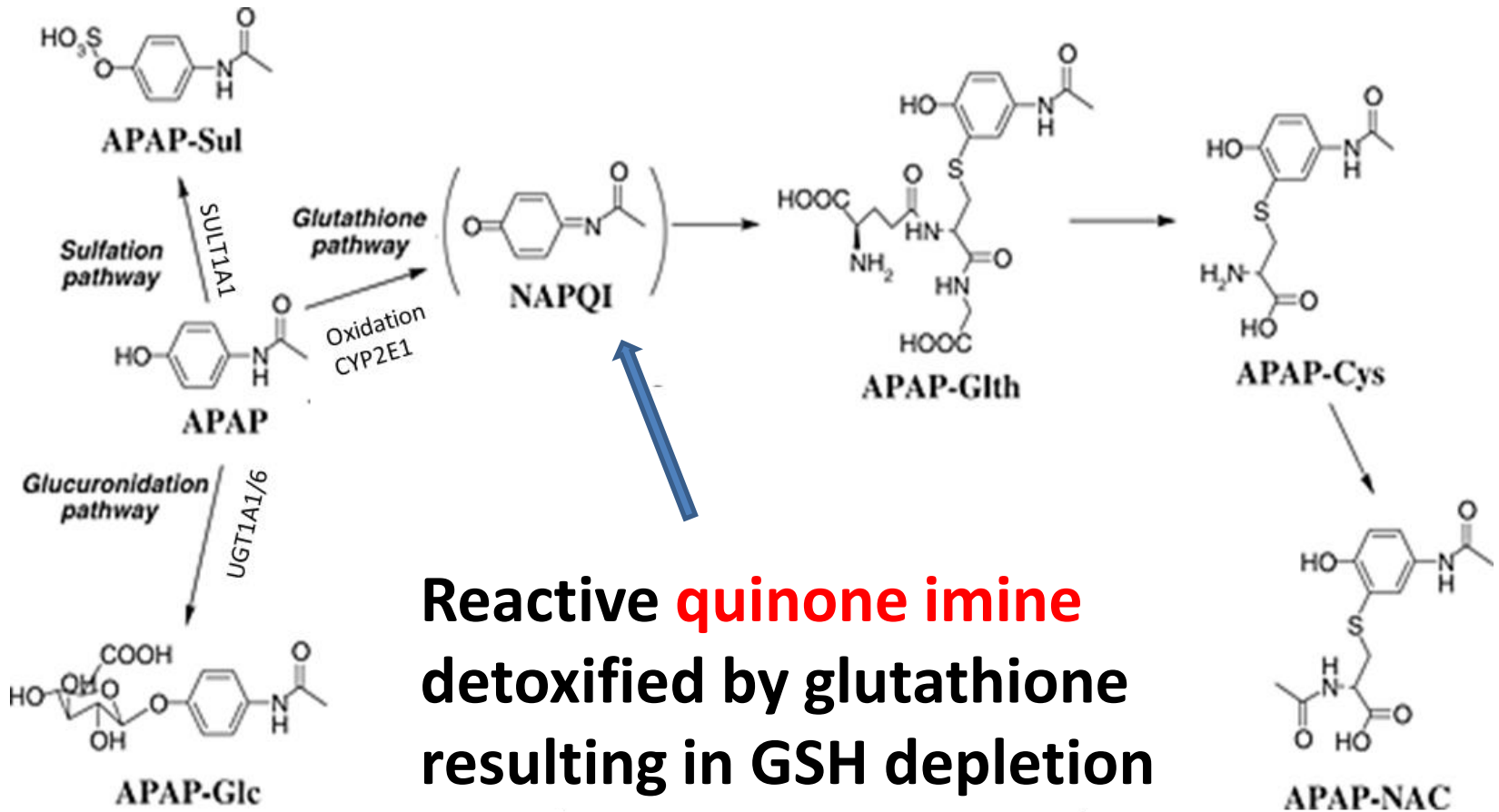
Liver “biomarkers” of alcohol treatment



Paracetamol Toxicity

- Paracetamol (acetaminophen) is a safe drug when taken at **therapeutic** doses but a major human **hepatotoxin** in overdose
- Results in a very large number of liver transplants
- Toxicity is the result of the formation of **reactive** metabolites

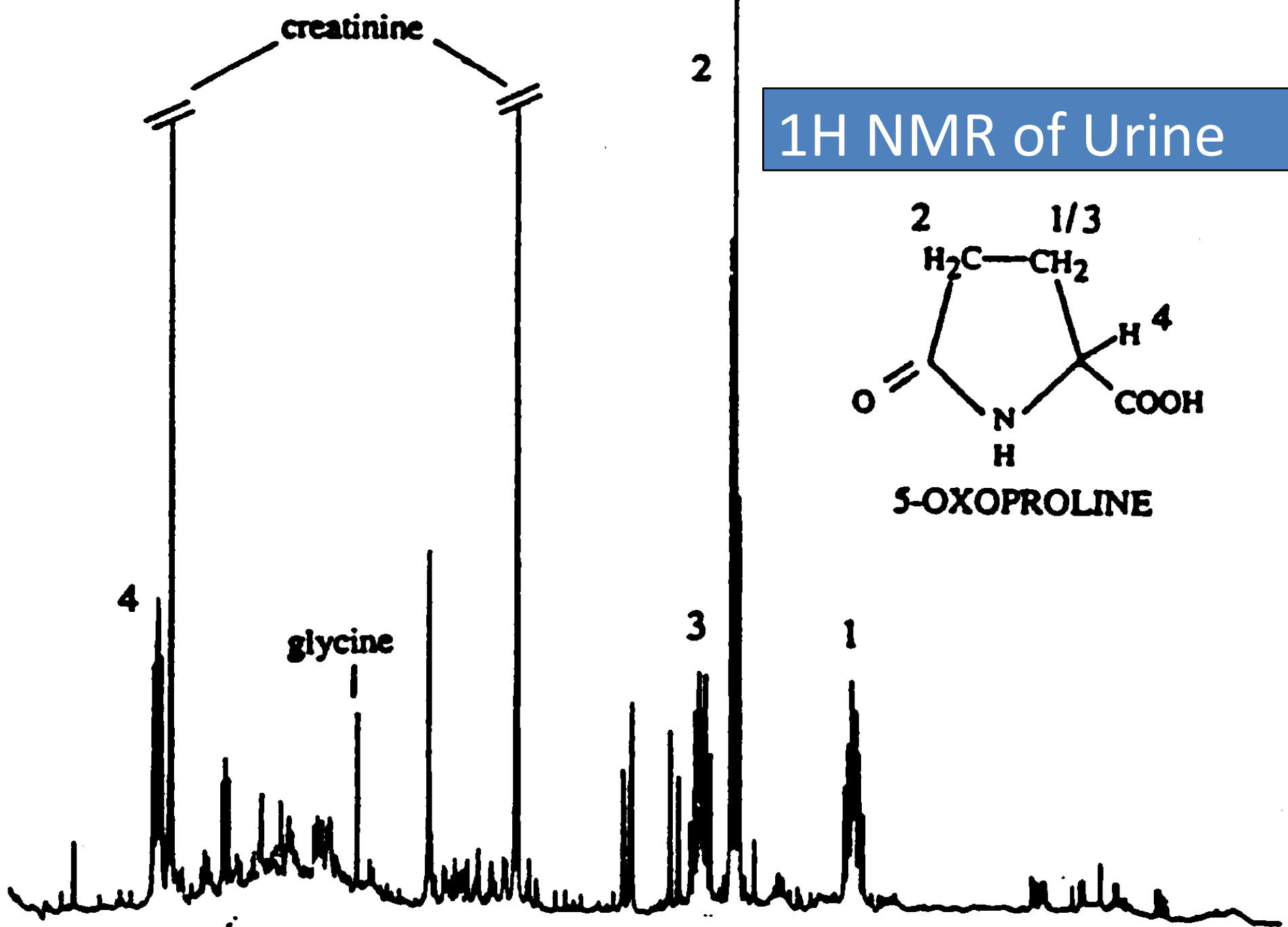
Paracetamol Metabolism



Matabonomics & Paracetamol Toxicity

- **F.Y. Ghauri *et al*** Induction of **5-oxoprolinuria** in the rat following chronic feeding with N-acetyl 4 aminophenol (paracetamol), *Biochem. Pharmacol.* 46 (**1993**) 953–957.
- ***An NMR-based study examining rat urine***
- **T. Soga *et al*** Differential metabolomics reveals **ophthalmic acid** as an oxidative stress biomarker indicating hepatic glutathione consumption, *J. Biol. Chem.* 281 (**2006**) 16768–16776.
- ***A CE-MS-based study in mice***

1H NMR of Urine



Elevated 5-Oxoproline reported in human overdose - translatable to humans.....

JAOA

CASE REPORT

5-Oxoproline-Induced Anion Gap Metabolic Acidosis After an Acute Acetaminophen Overdose

David T. Lawrence, DO; Laura K. Bechtel, PhD; Nathan P. Charlton, MD; and Christopher P. Holstege, MD

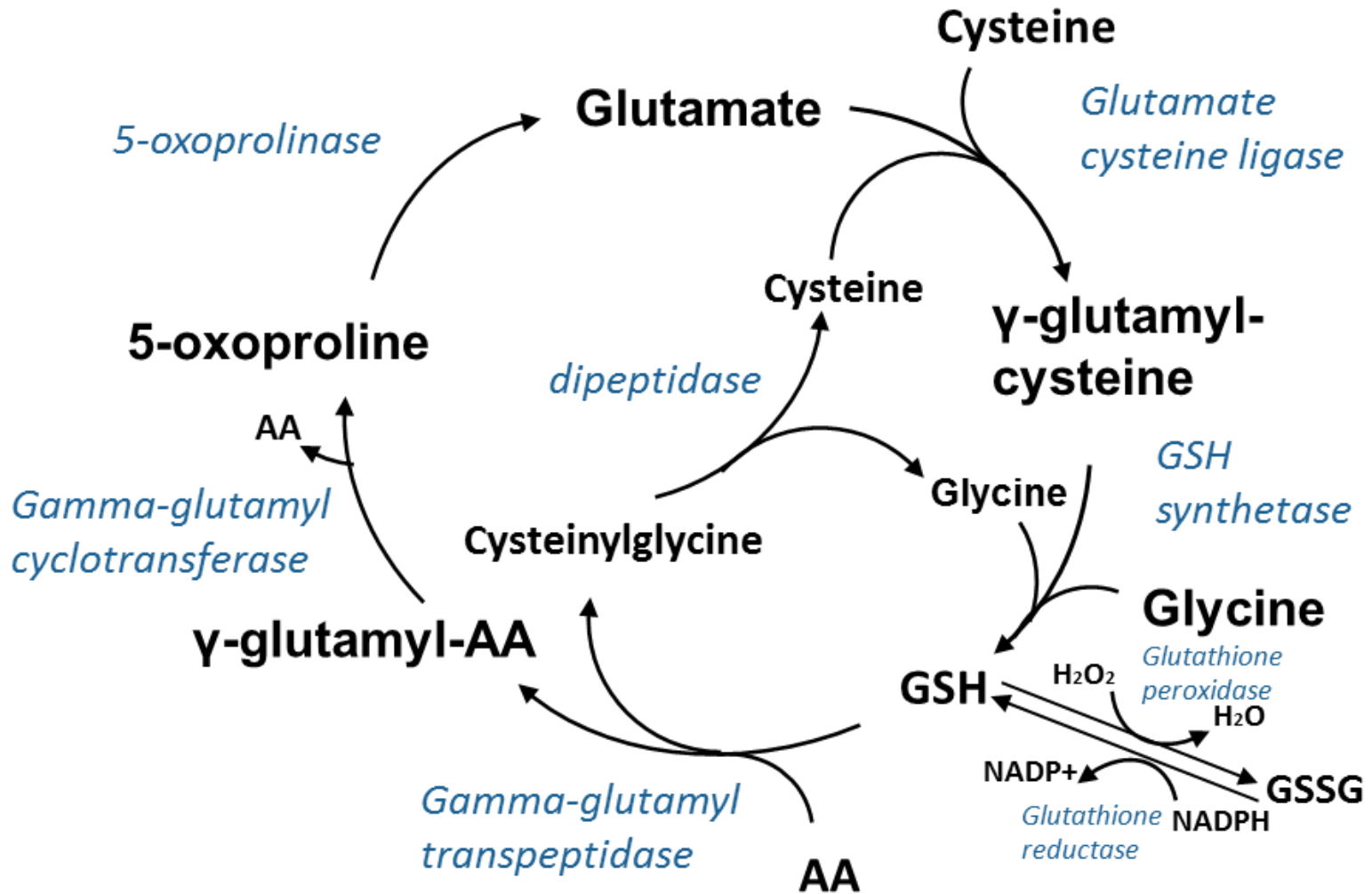
5-OXOPROLINEMIA CAUSING ELEVATED ANION GAP METABOLIC ACIDOSIS IN THE SETTING OF ACETAMINOPHEN USE

Patil Armenian, MD,* Roy R. Gerona, PhD,† Paul D. Blanc, MD,*‡
Alan H. B. Wu, PhD,* and Somnath Mookherjee, MD‡

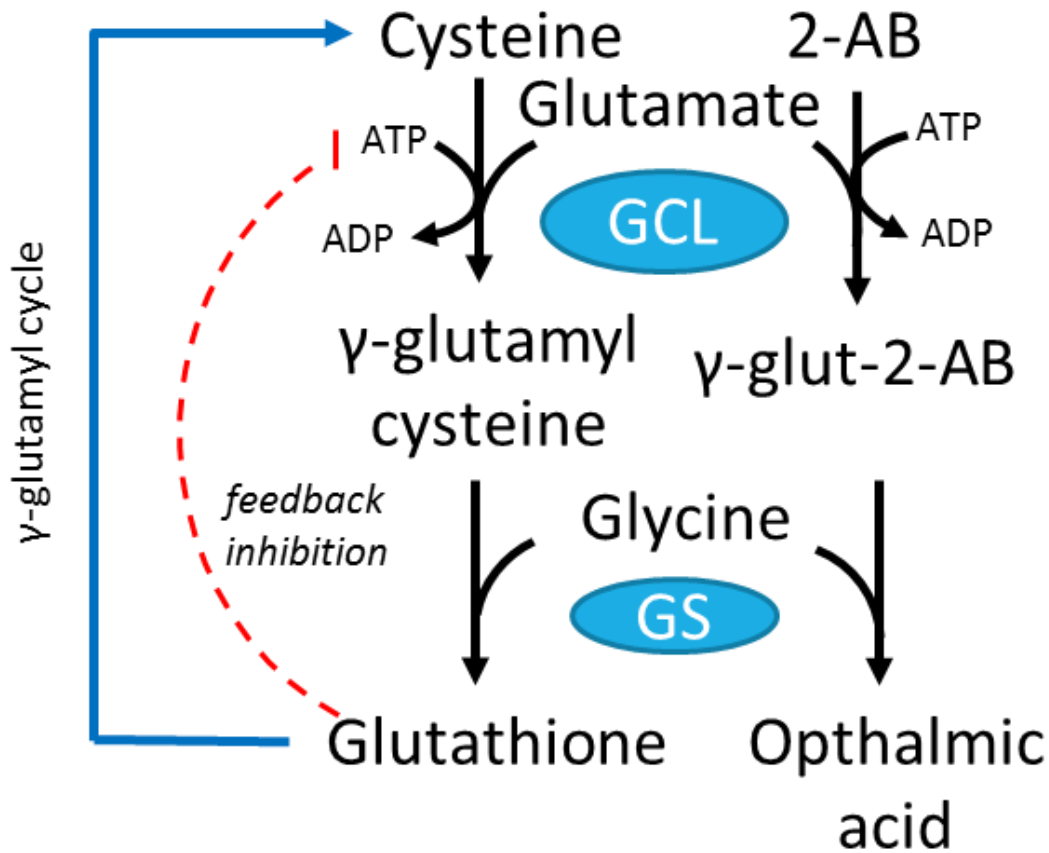
*Division of Clinical Pharmacology, California Poison Control System, San Francisco Division, †Department of Laboratory Medicine, San Francisco General Hospital, and ‡Department of Medicine, University of California-San Francisco, San Francisco, California
Reprint Address: Patil Armenian, MD, California Poison Control System, San Francisco Division, UCSF Box 1369, San Francisco, CA 94143-1369

So what is the role of 5-oxoproline?

5-Oxoproline & Glutathione Biosynthesis

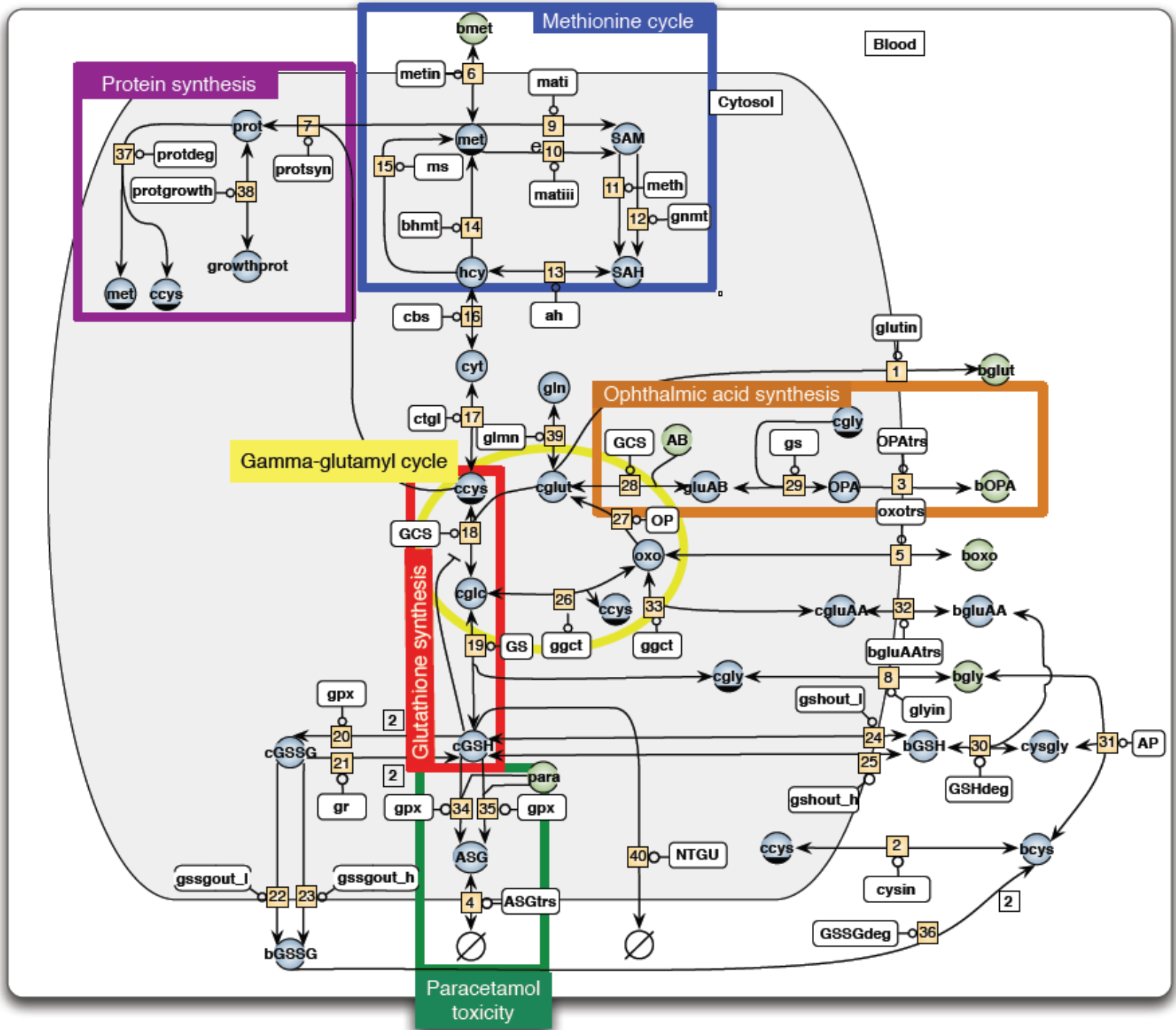


Ophthalmic acid Biosynthesis



Why are these metabolites biomarkers?

- Paracetamol is toxic via reactive metabolites formed by cytochrome P450 2E1
- The reactive quinoneimine metabolite is detoxified by conjugation with glutathione
- Depletion of glutathione is followed by damage to cellular macromolecules and cell death
- 5-oxoproline (pyroglutamate) and ophthalmic acid are involved in the biosynthetic pathways that relate to glutathione
- Why both metabolites? What are these biomarkers telling us? which is one is best? Or do you need both?
- Can we model it using systems biology?



The model

- Kinetic model - reversible Michaelis-Menten equations

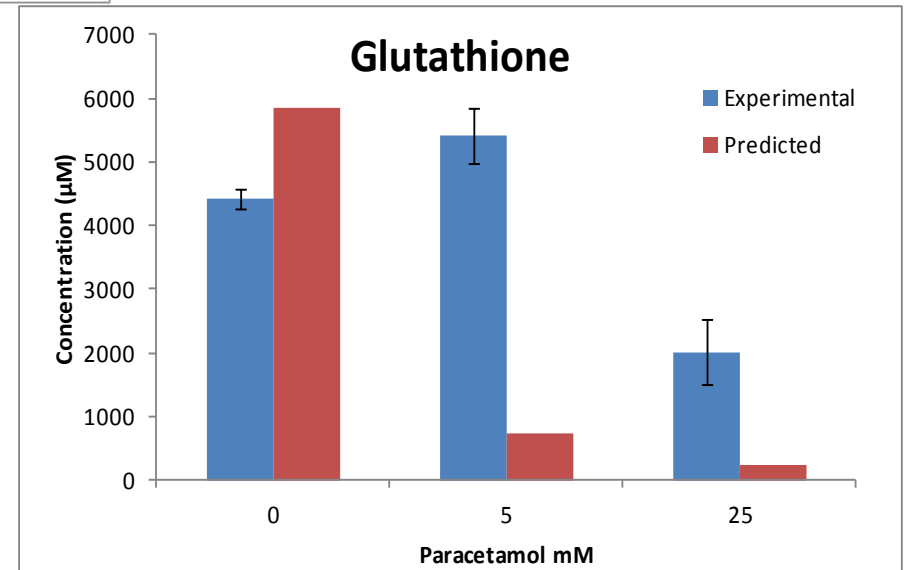
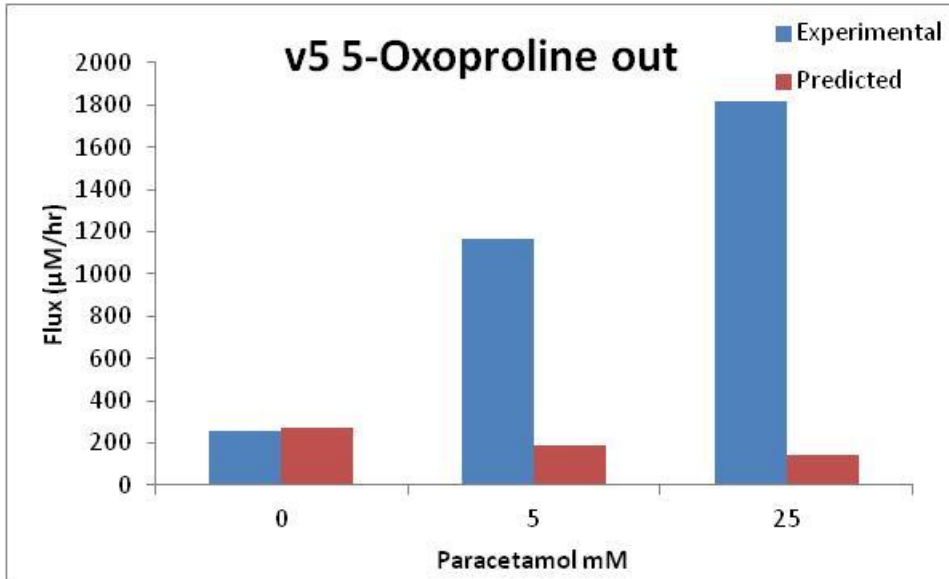
$$\begin{aligned} \frac{d([cGSH] \cdot V_{cell})}{dt} = & \frac{vGSf1 \cdot [cglc] \cdot [cgly] \cdot \left(1 - \frac{[cGSH]}{keqGS1 \cdot [cglc] \cdot [cgly]}\right)}{kGSeglc \cdot kGSegly \cdot \left(1 + \frac{[cgly]}{kGSegly} + \frac{[gluAB]}{kGSgluAB}\right) \cdot \left(1 + \frac{[cglc]}{kGSeglc} + \frac{[cGSH]}{kGSecGSH} + \frac{[OPA]}{kGSOPA}\right)} \\ & - 2 \cdot \frac{H2O2 \cdot vmGPX \cdot [cGSH]^2}{(H2O2 + 9 \cdot kmGPXH2O2) \cdot (kmGPXgsh + [cGSH])^2} \\ & + 2 \cdot \frac{eNADPH \cdot vmgr \cdot [cGSSG]}{kmgrGSSG \cdot kmgrNADPH \cdot \left(1 + \frac{eNADPH}{kmgrNADPH} + \frac{[cGSSG]}{kmgrGSSG} + \frac{eNADPH \cdot [cGSSG]}{kmgrGSSG \cdot kmgrNADPH}\right)} \\ & - \frac{vmgshoutl \cdot [cGSH]^3}{kmgshoutl^3 + [cGSH]^3} - \frac{vmgshouth \cdot [cGSH]}{kmgshouth + [cGSH]} \end{aligned}$$

- Used mostly literature values
- V_{max} and unknown parameters were fitted to experimental data

Fitting to experimental data

- Use an in vitro cell model (THLE-2E1cells)
- Incubate cells at three paracetamol concentrations: 0, 5mM and 25mM
- Measure (LC-MS) the concentration of metabolites in the glutathione pathway (intracellular and extracellular)
 - Measured 15 fluxes and concentrations using quantitative targeted metabolite determination by LC-MS

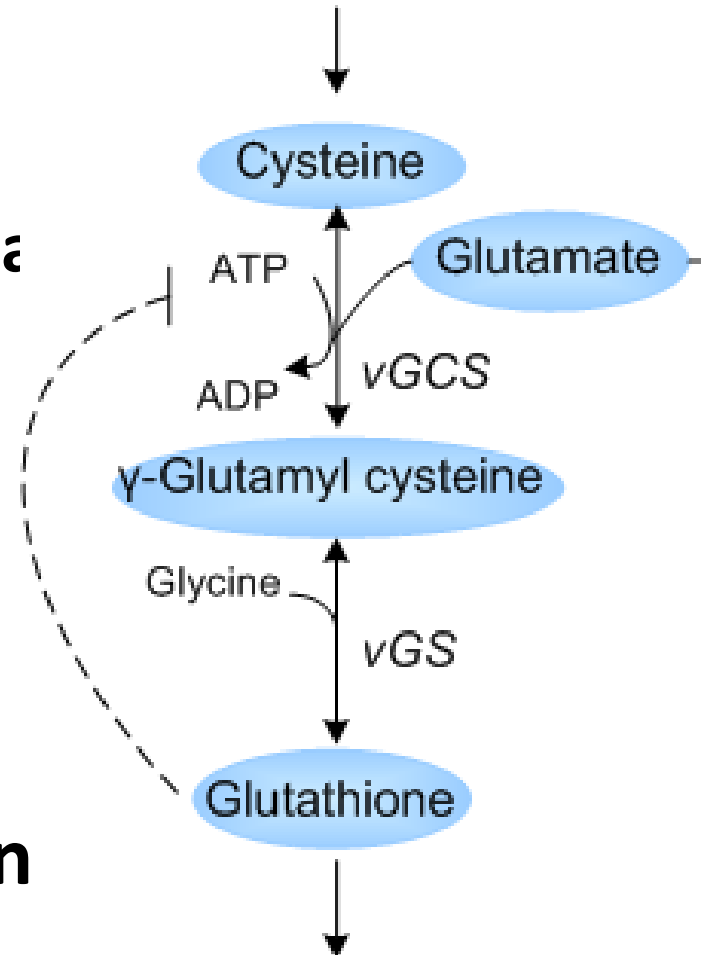
Experiment vs Model



The model predicts the drug free conditions well, but not what occurs with increasing paracetamol exposure

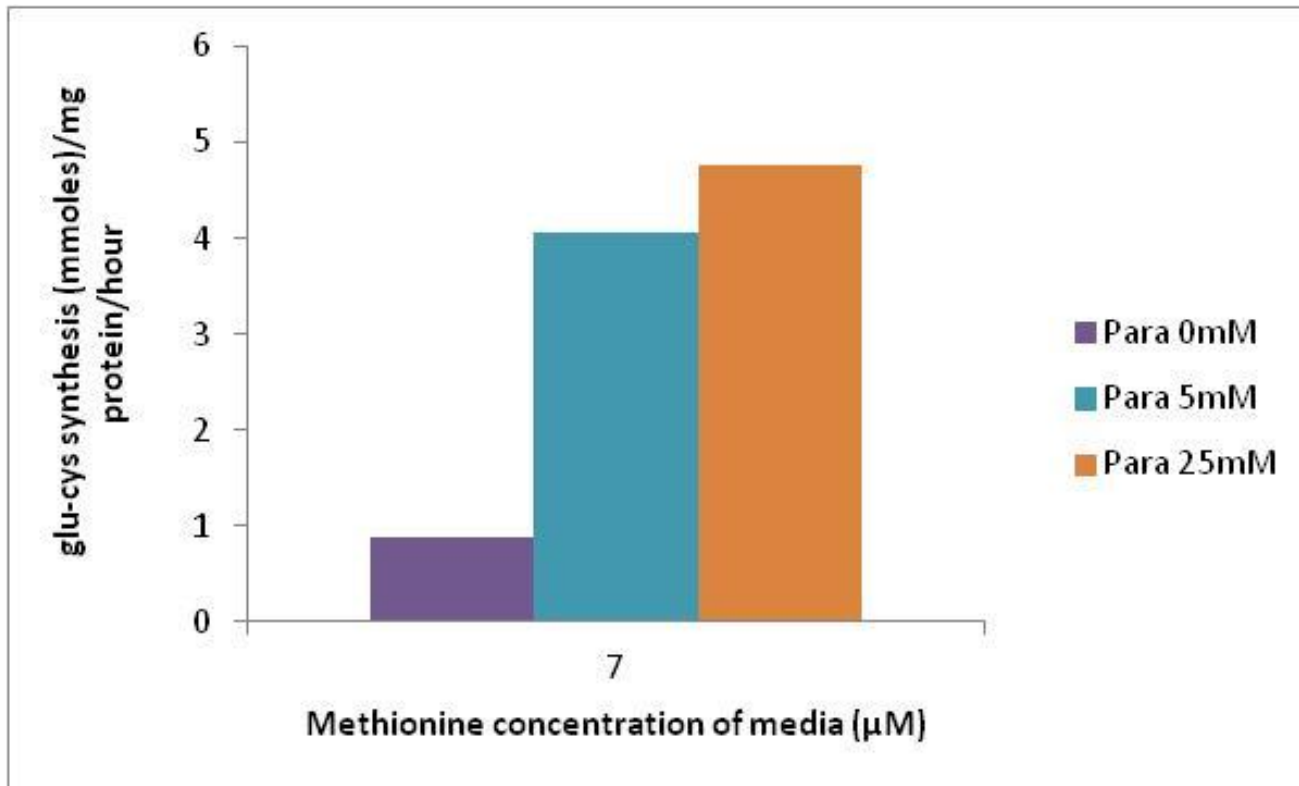
Model predicts control in glutamyl-cysteine synthetase (GCS)

- Metabolic control analysis
- At 25mM paracetamol V_{GCS} had a control of:
 - 1 on 5-oxoproline flux
 - 0.9 on GSSG synthesis
 - 2 on GSH export
- Increases in GCS reported in the literature



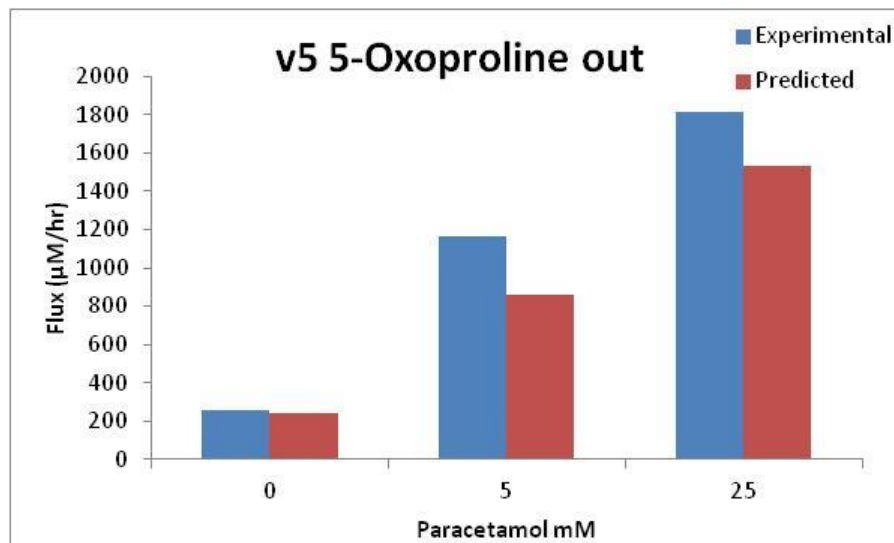
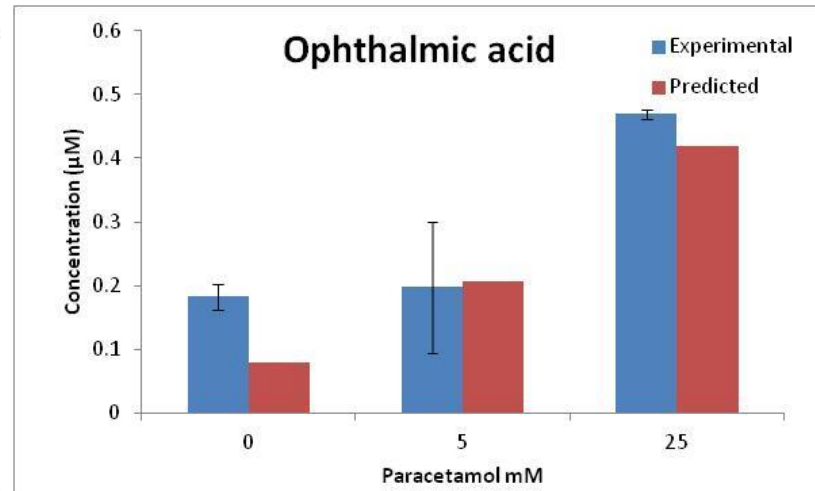
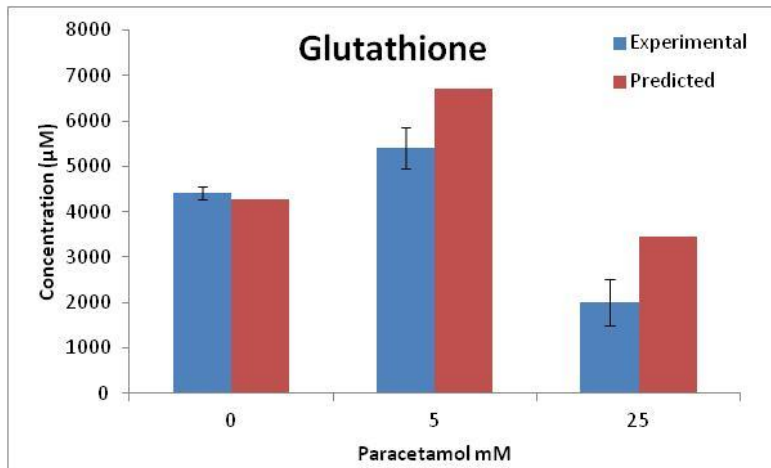
GCS activity vs paracetamol exposure

- Enzyme activity assay



Add adaption of GCS to model

- Good fits for GSH, OPA & 5-Oxo concentrations and fluxes



Biomarking two stages of toxicology

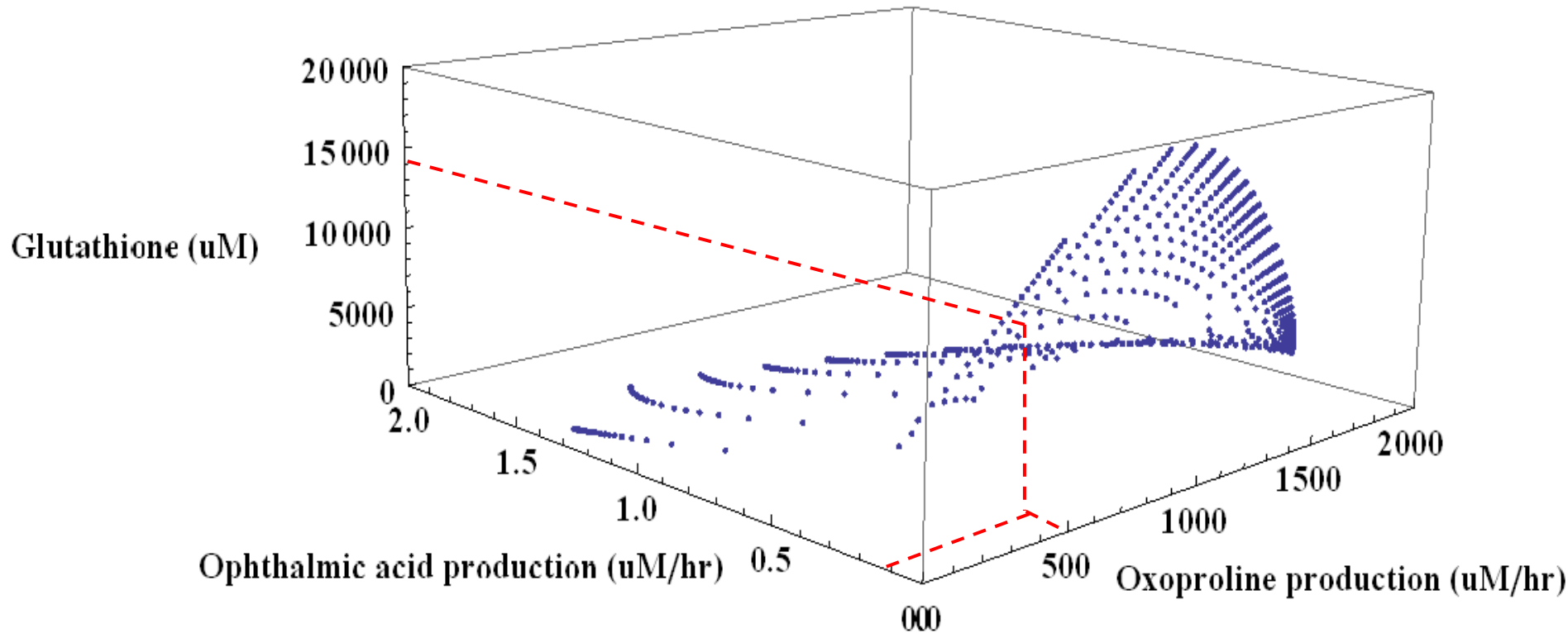
- Stage 1:
 - Paracetamol-conjugate formation
 - **Glutathione synthesis increases**
 - No depletion in cellular glutathione
- Stage 2:
 - Paracetamol-conjugate formation
 - Methionine and cysteine depleted
 - Glutathione synthesis decreases
 - **Depletion of cellular glutathione**

5-oxoproline

**Ophthalmic
acid**

Biomarkers complement each other

- *In vitro* measuring biomarkers simultaneously will give unique glutathione concentration

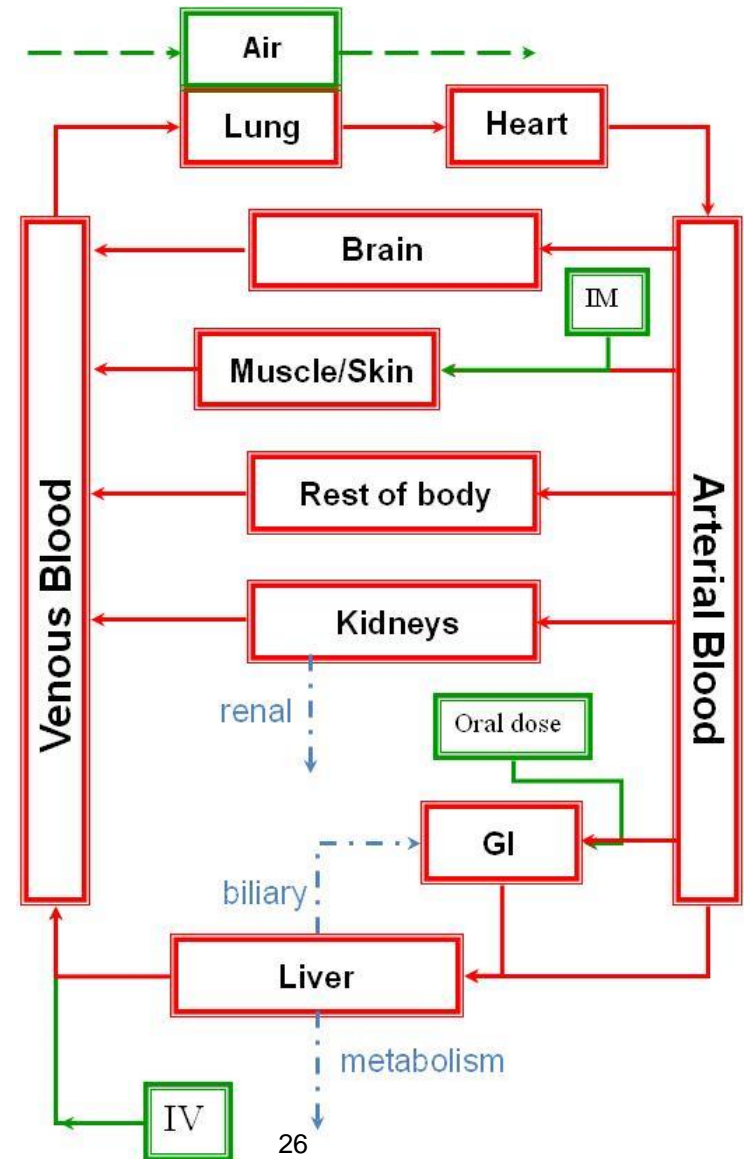


In vitro to in vivo –

Physiologically Based PK Modelling (PBPK)

Organism mathematically described as a series of compartments that represent tissues and organs

- Arranged to reflect anatomical layout
- Connected by the arterial and venous pathways
- Defined using literature tissue volumes and blood flow rates
- Progress of drug through the body can be followed by defining the discrete ADME properties



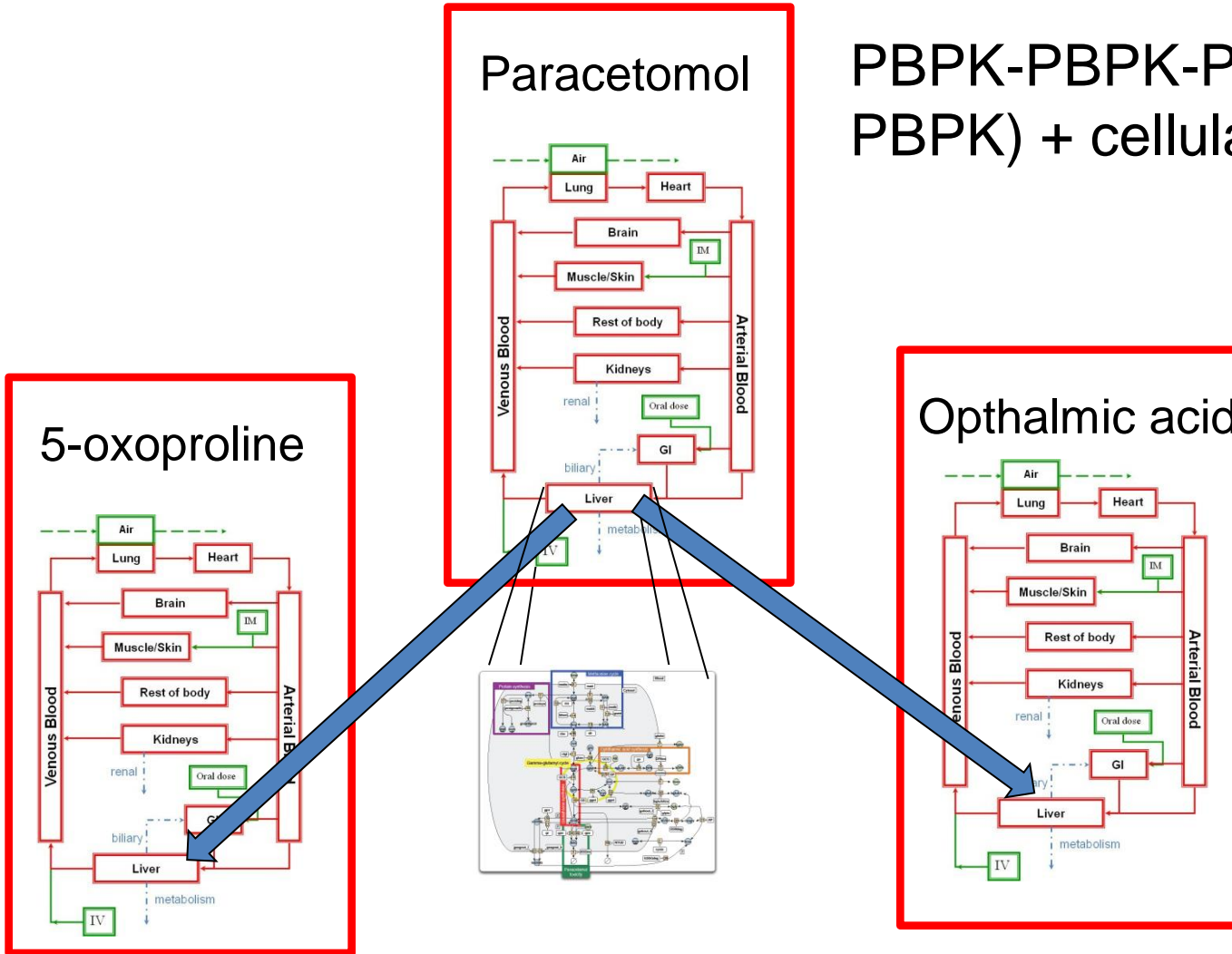
Model for Both Drug and Biomarkers

Paracetamol

PBPK-PBPK-PBPK (Triple PBPK) + cellular model.

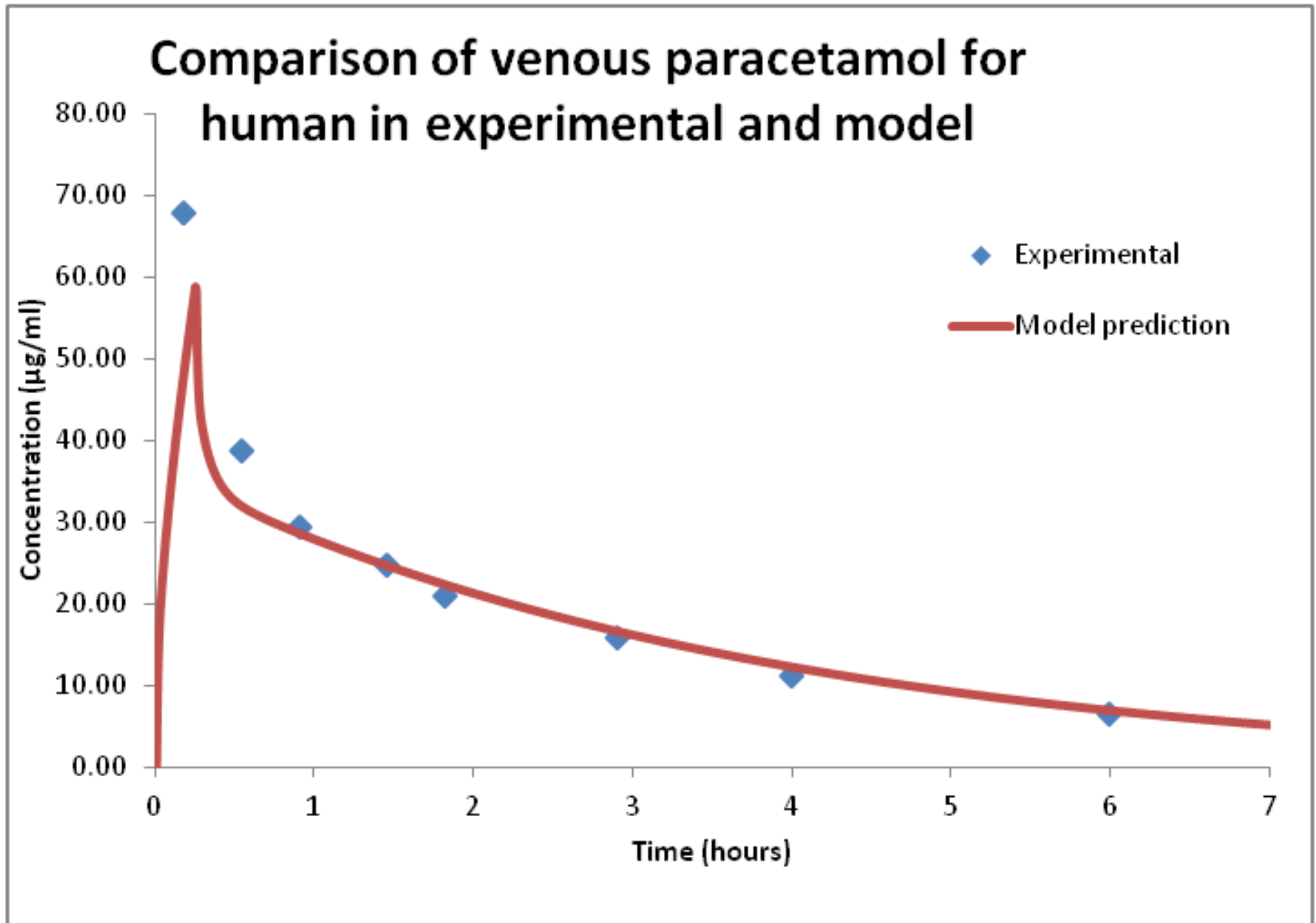
5-oxoproline

Ophthalmic acid

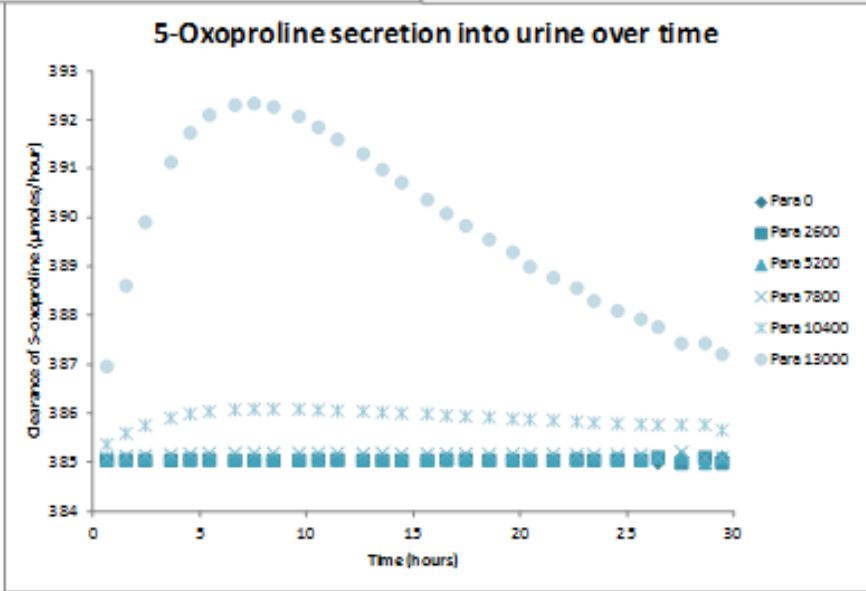
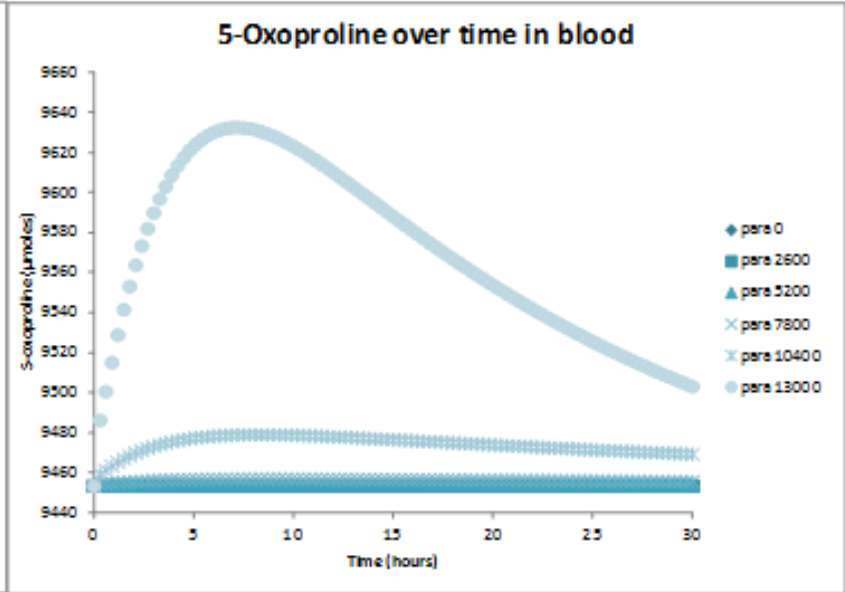
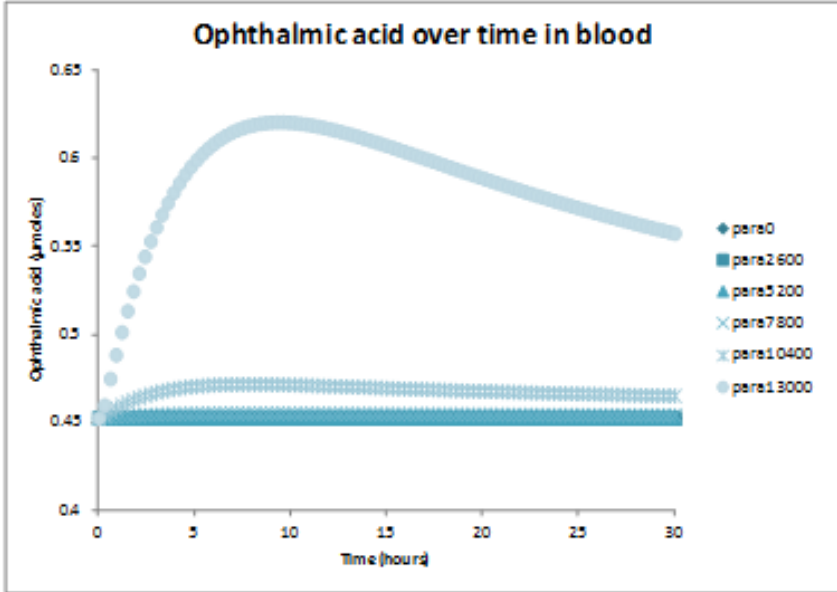


Before simulating the effects of paracetamol on the biomarkers we need to know that the PBPK models reproduces the literature results.

Paracetamol in Humans: Actual vs Predicted



Simulating the effects on 5-oxo and OPA after dosing paracetamol



**Biomarker
s raised in
blood and
urine**

Conclusions

- **Biomarkers – discovering metabolites as *potential* biomarkers is easy**
- **But they only become real biomarkers when they have been validated**
- **And are only useful when you understand what they are telling you**
- **Biomarker validation requires targeted methods and bespoke studies**
- **Systems Biology should be a key to deeper understanding**

Acknowledgements

- **Robert S. Plumb, Elaine Holmes, Jeremy K. Nicholson**
- **Eleni Gika, Filippos Michopoulos, Georgios Theodoridis**
- **Suzanne Geenen, Hans Westerhof**
- **Neil Loftus, Neil Kaplowitz, Cheng Ji**