

Incorporating Improved Physiological Relevance in Tox21: An Evolution of Resolution

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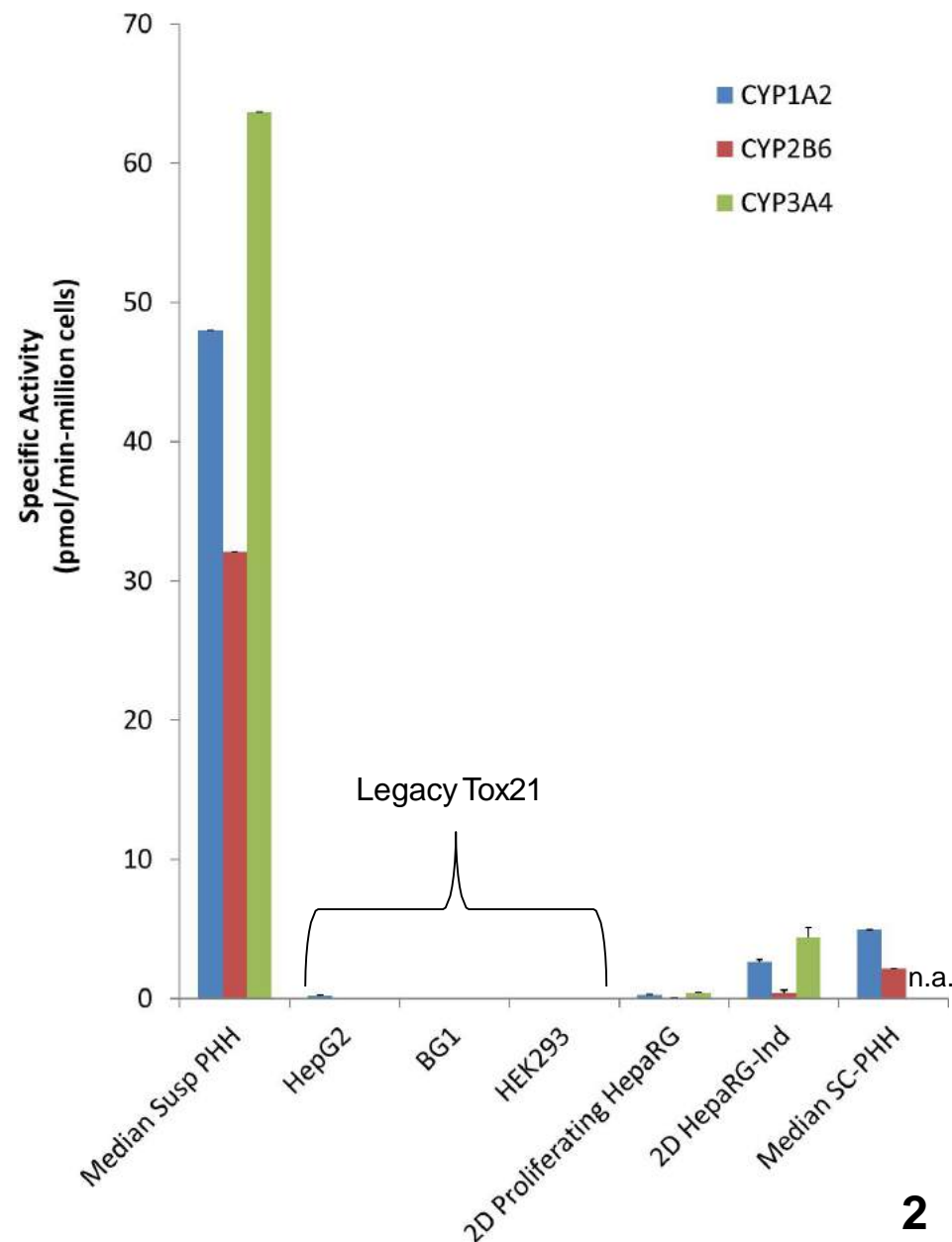
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Phase I & Phase II of Tox21 (NCATS, EPA, FDA & NTP)

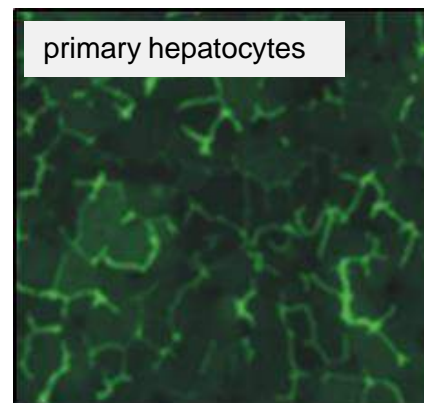
- Phase I (2005-2010)
 - Proof of concept
- Phase II (2011-present)
 - ToxCast Phase II: ~700 compounds, ~700 assays
 - NCGC/NCATS: 10k compound library
 - nuclear receptor
 - cellular stress pathways



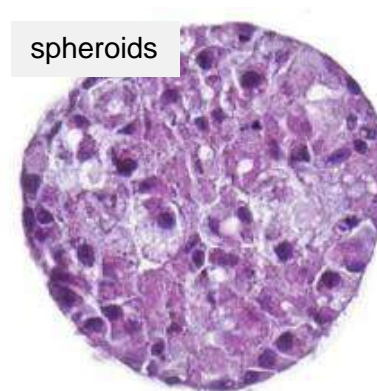


Tox21 Phase III: Predictive Toxicology Screening

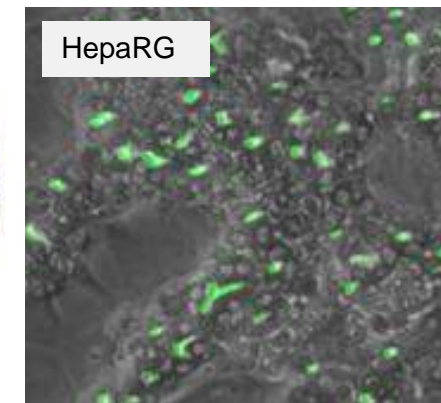
- Physiologically-relevant in vitro screening models
 - **xenobiotic metabolism** ‘competence’
 - longevity to address longer-term exposures (weeks/months)
- Data-rich, high dimensional assay platforms as windows into chemical-induced perturbation cascades
 - **high throughput transcriptomics** (time & concentration)
 - high content imaging (i.e., stress pathways)
 - metabolomics (xenobiotic & endogenous)
- Quantitatively link chemicals & metabolites to biological response pathways
 - C_{max}/EC_{50} ratios
 - AUC correlations
 - IVIVE
- Extend ‘normal’ models to susceptibility models & extrahepatic
 - inflamed liver, fatty liver, cholestatic liver
 - extrahepatic tissues (i.e., kidney, intestine)
 - developmental tissue models (i.e., neonatal, stem cells)



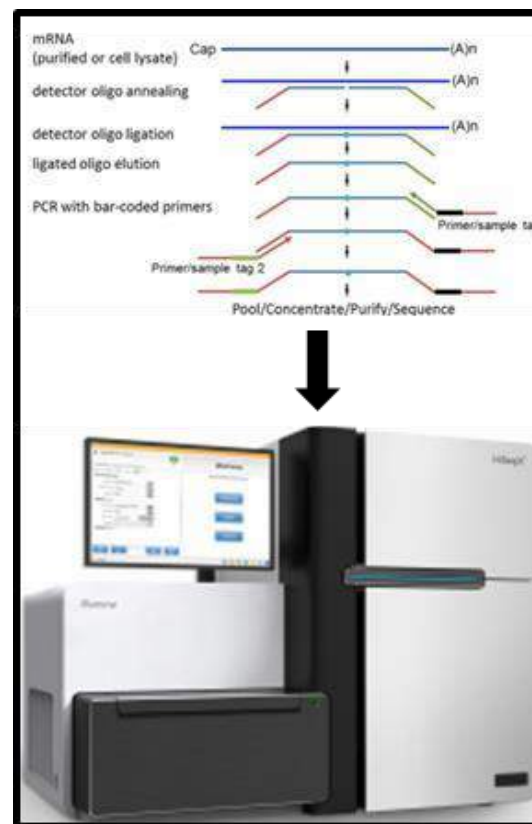
primary hepatocytes



spheroids



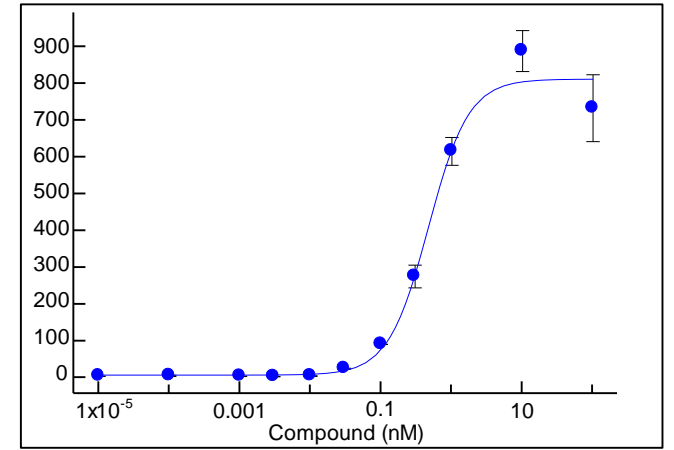
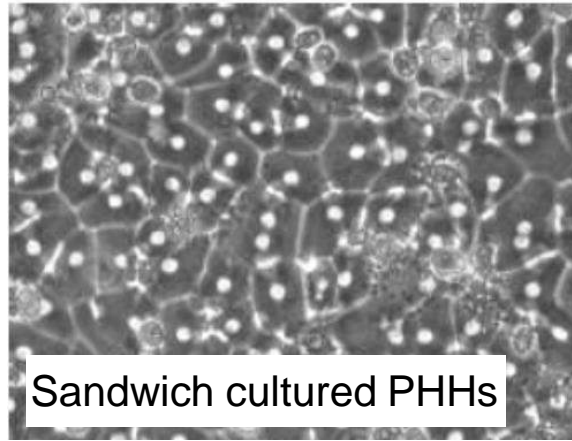
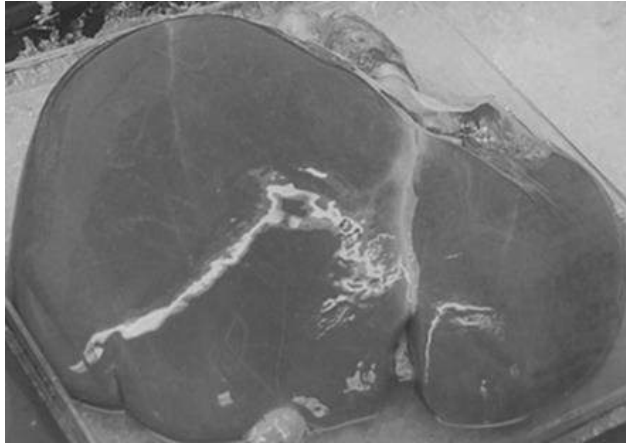
HepaRG





- **Sandwich cultures of primary human hepatocytes to explore:**
 - Ginkgo biloba extract (GBE) biological response equivalence across 29 lots
 - Predict their potential for liver enzyme induction in humans
 - Link GBE constituents with response phenotypes: liver enzyme induction/receptor pathway activation, cytotoxicity & oxidative stress
- **3D organotypic spheroid cultures of HepaRG cells to:**
 - Screen chemicals with physiologically-relevant xenobiotic metabolism
 - Support functional liver enzyme induction pathways
 - Identify liver injury potential with chemicals
- **High throughput transcriptomics paired with in vitro liver models to:**
 - Explore tissue model functionality, chemical-biological interaction gene & pathway dynamics
 - Characterize the impact of cell culture configuration
 - Dose-response modeling of benchmark dose & biological pathway perturbations

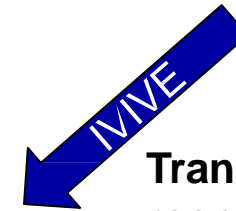
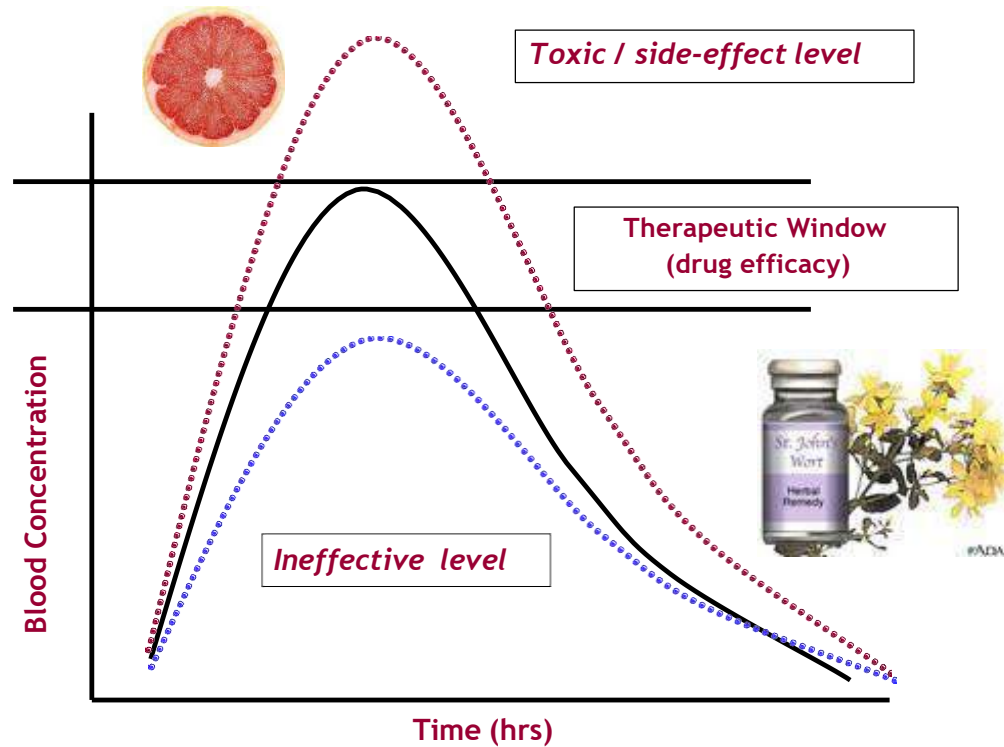
Prediction of Drug Metabolism & Drug-Drug Interactions with In Vitro Models



Drug Labelling

negative

positive

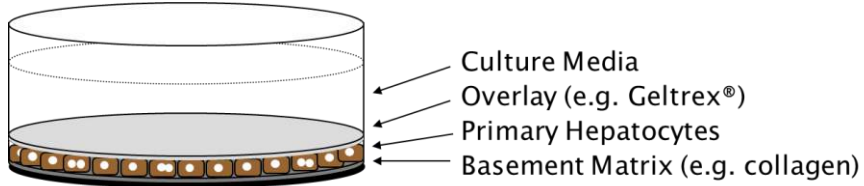


Translation of in vitro responses to clinical effects



Ginkgo Biloba Extract Study Design Summary

- Sandwich cultures of PHHs (SC-PHHs)



- Endpoint Assays:

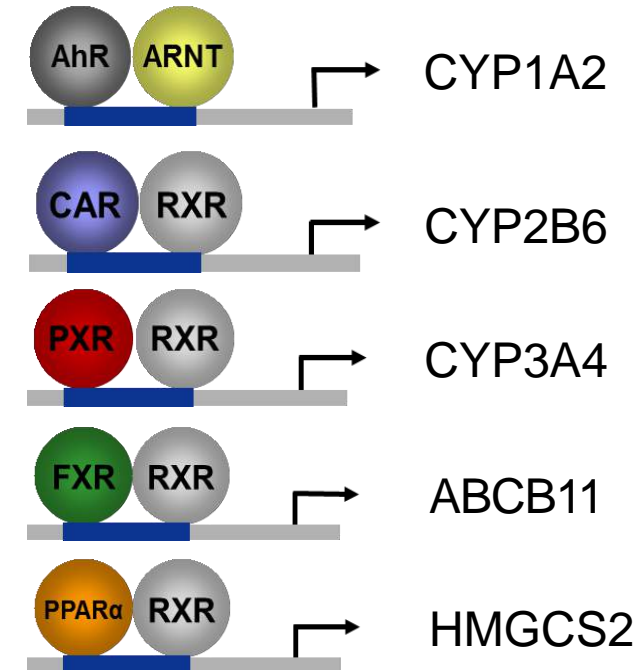
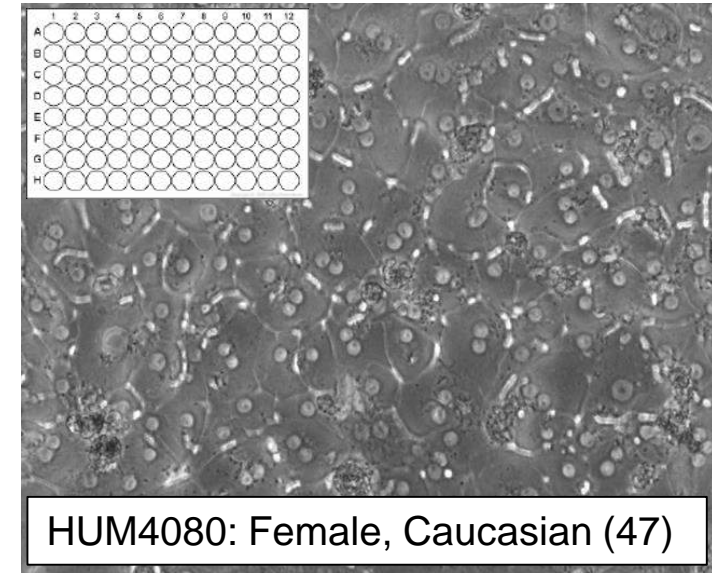
- Cell Health:

- Photomicrographs of each culture (Incucyte)
- ATP Depletion (CellTiter-Glo)
- Reactive Oxygen Species (H₂O₂, ROS-Glo)

- Liver enzyme induction/receptor activation

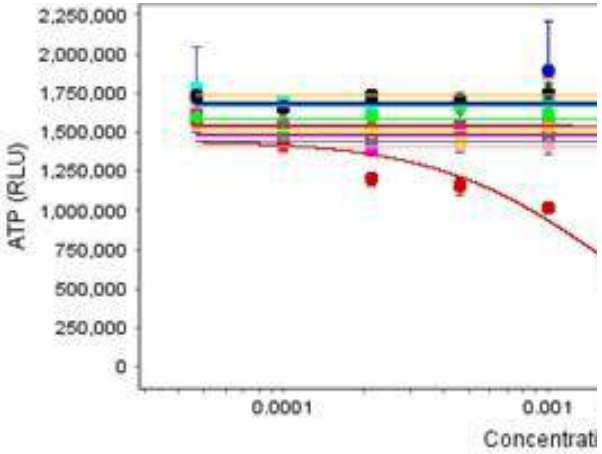
- mRNA (TaqMan)
 - 5 target genes, sentinels to hepatic receptor pathways
 - 3 Endogenous Ctrls (PSMB6, GAPDH, β-actin)
- Liver enzymatic activity
 - CYP1A2 (phenacetin to acetaminophen)
 - CYP2B6 (bupropion to hydroxybupropion)
 - CYP3A4/5 (midazolam to 1-hydroxymidazolam)

- Samples frozen for future transcriptomics/metabolomics

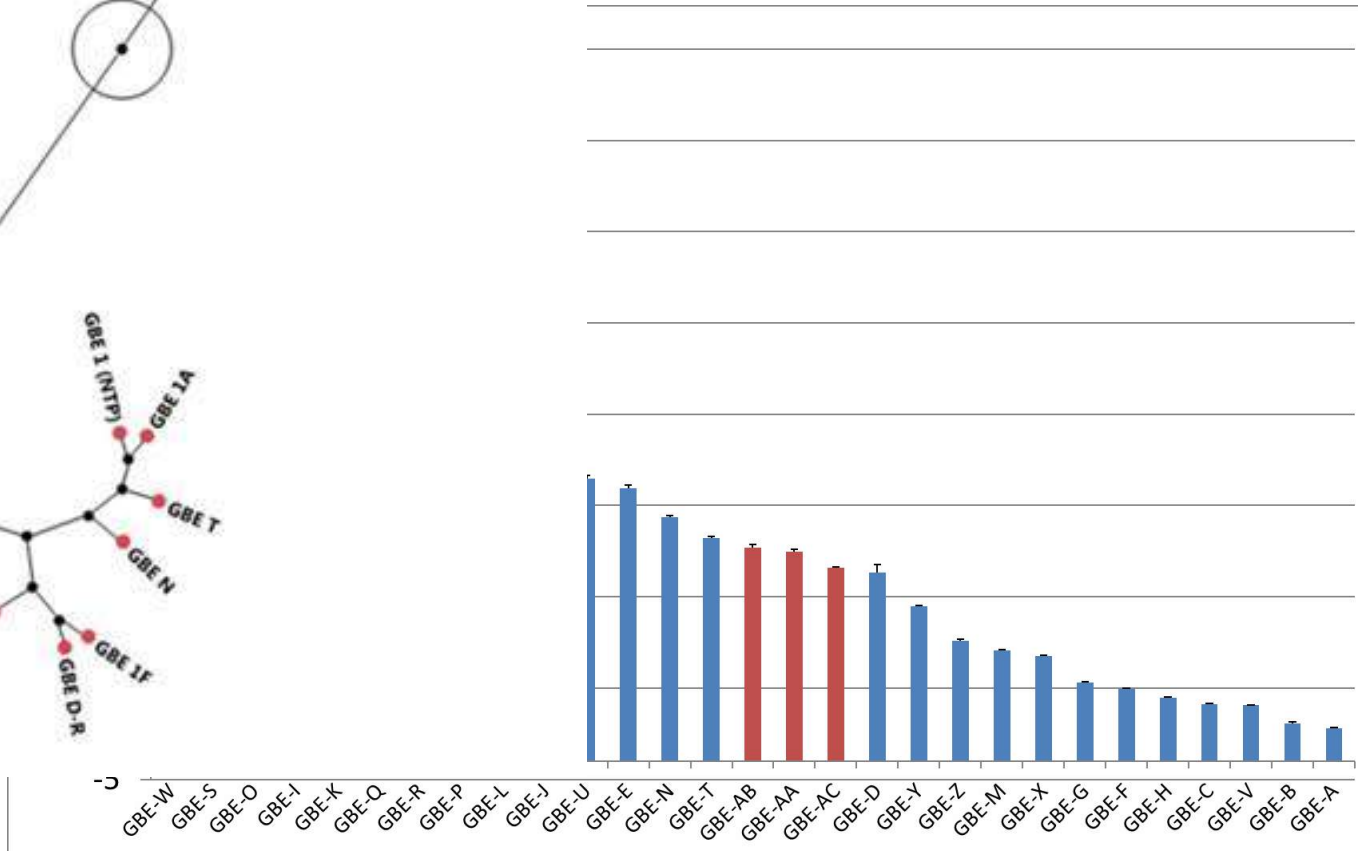
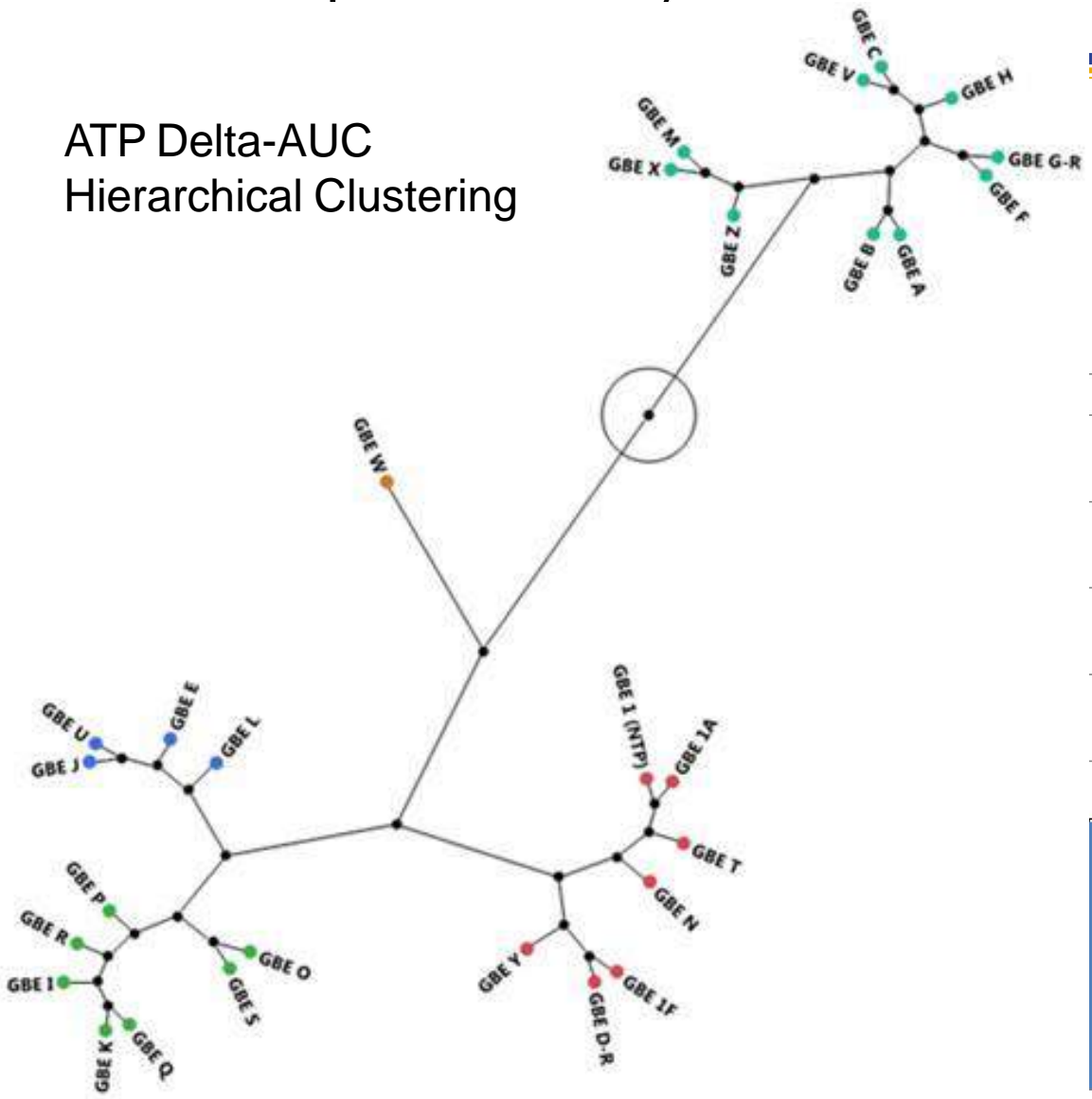


ATP Depletion Assays with SC-PHHs & GBEs

ATP Delta-AUC
Hierarchical Clustering

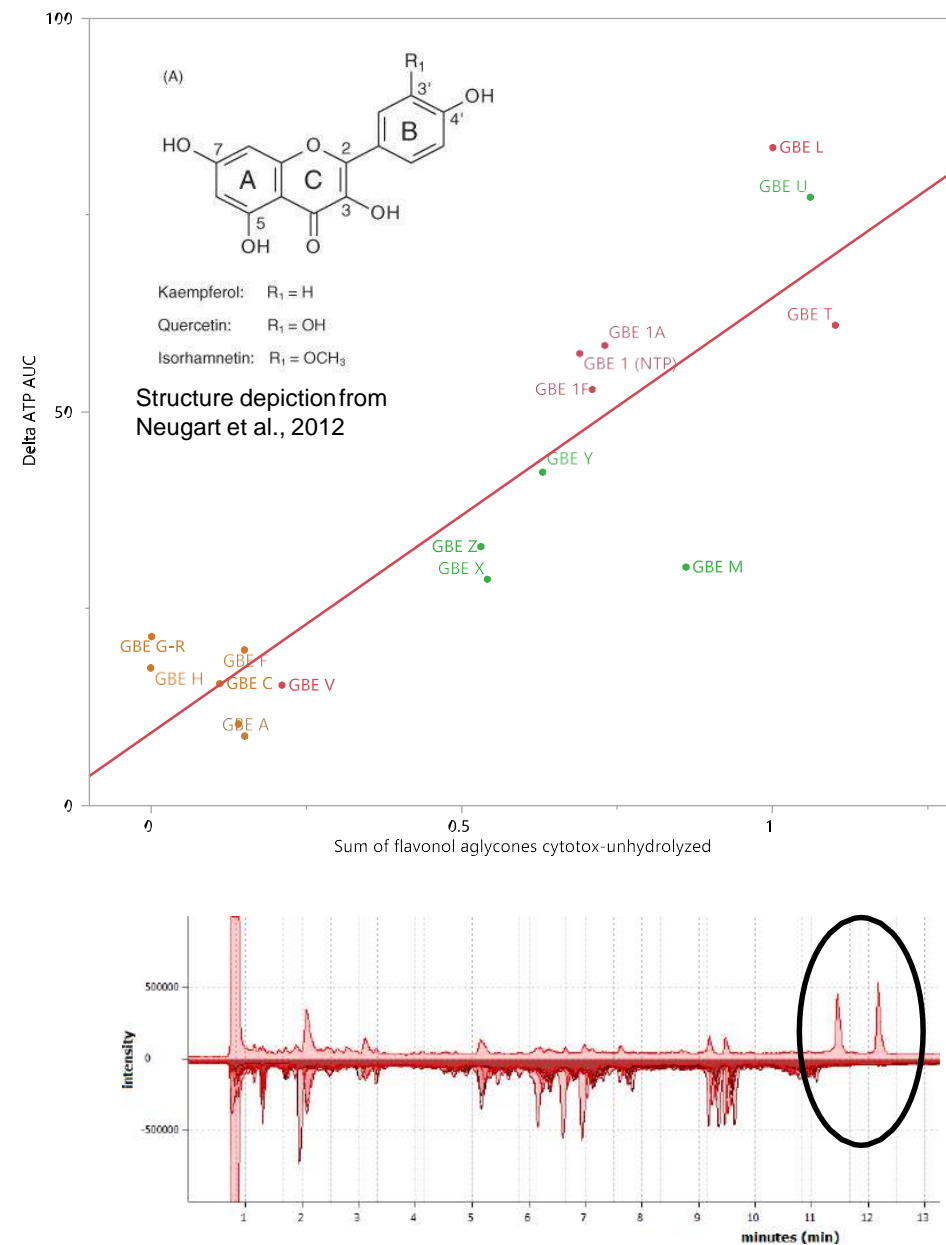
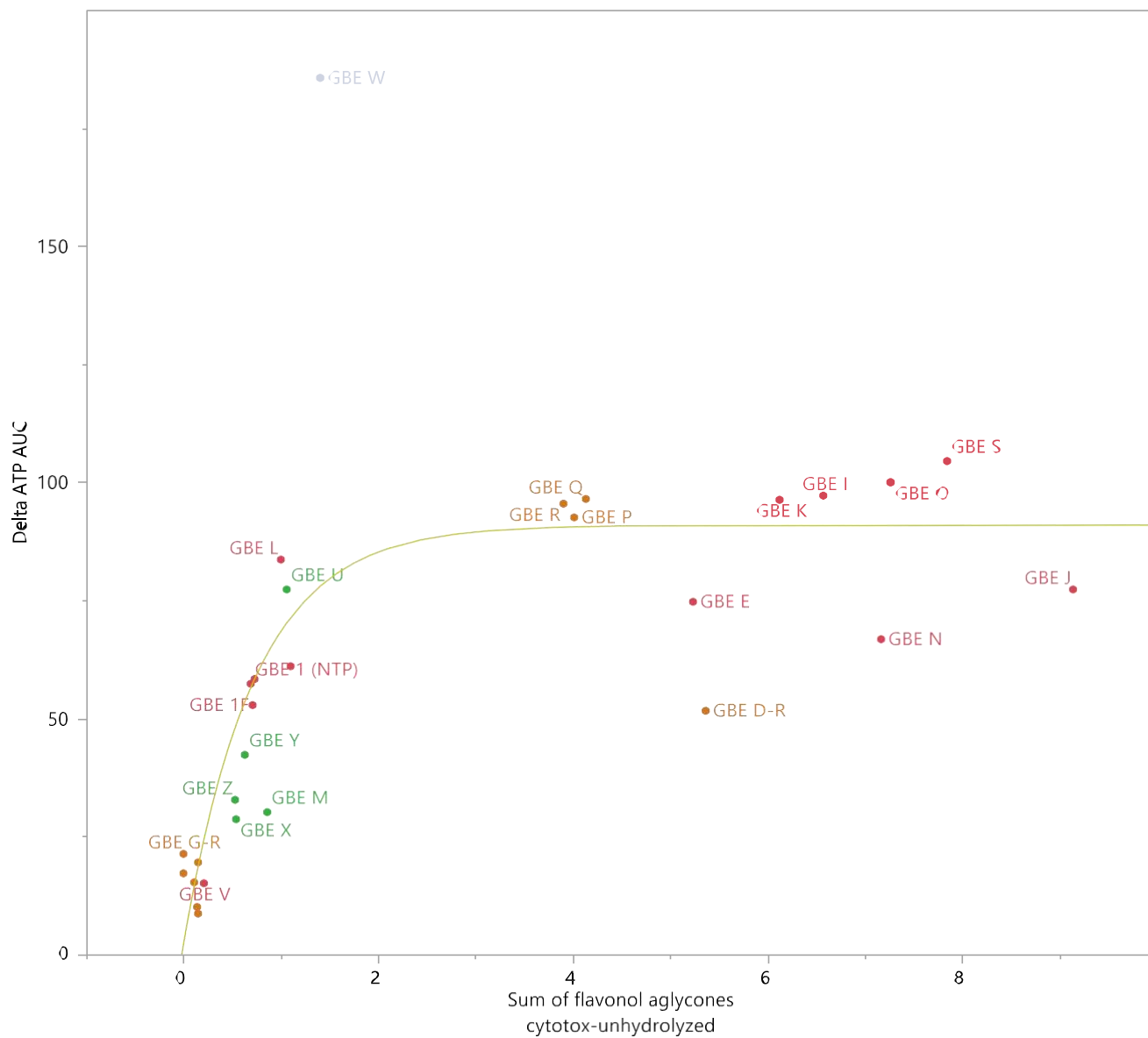


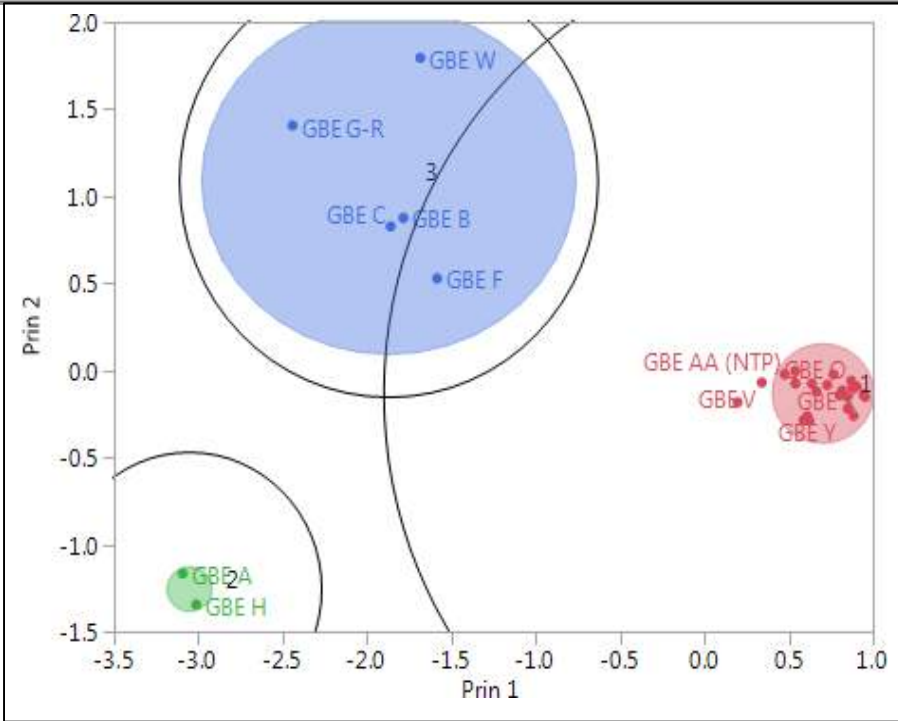
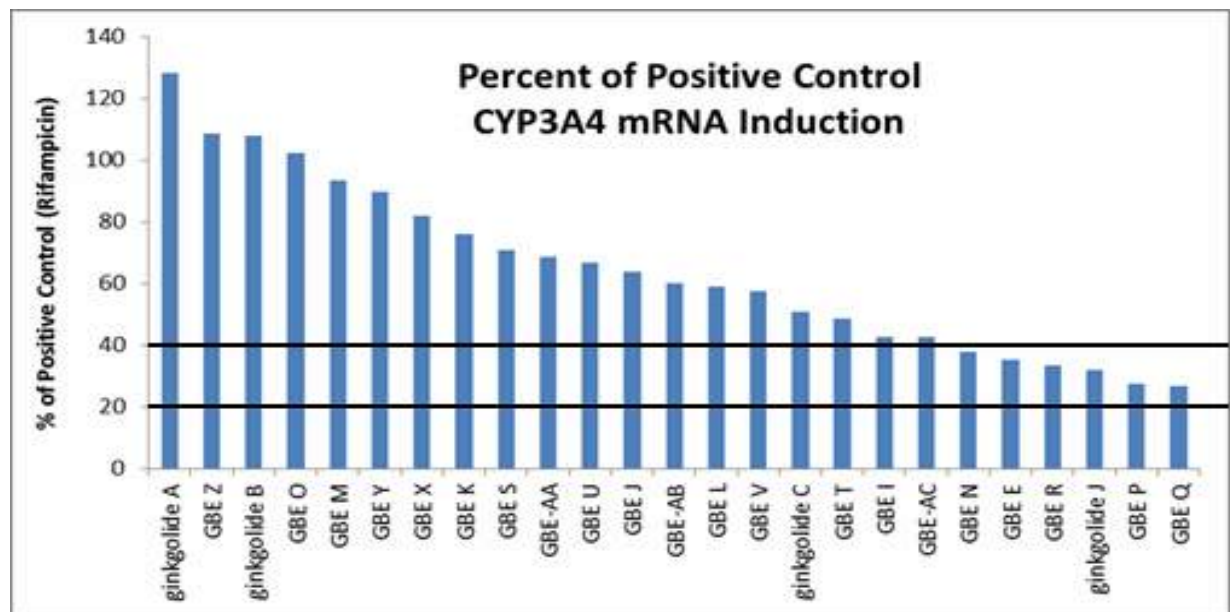
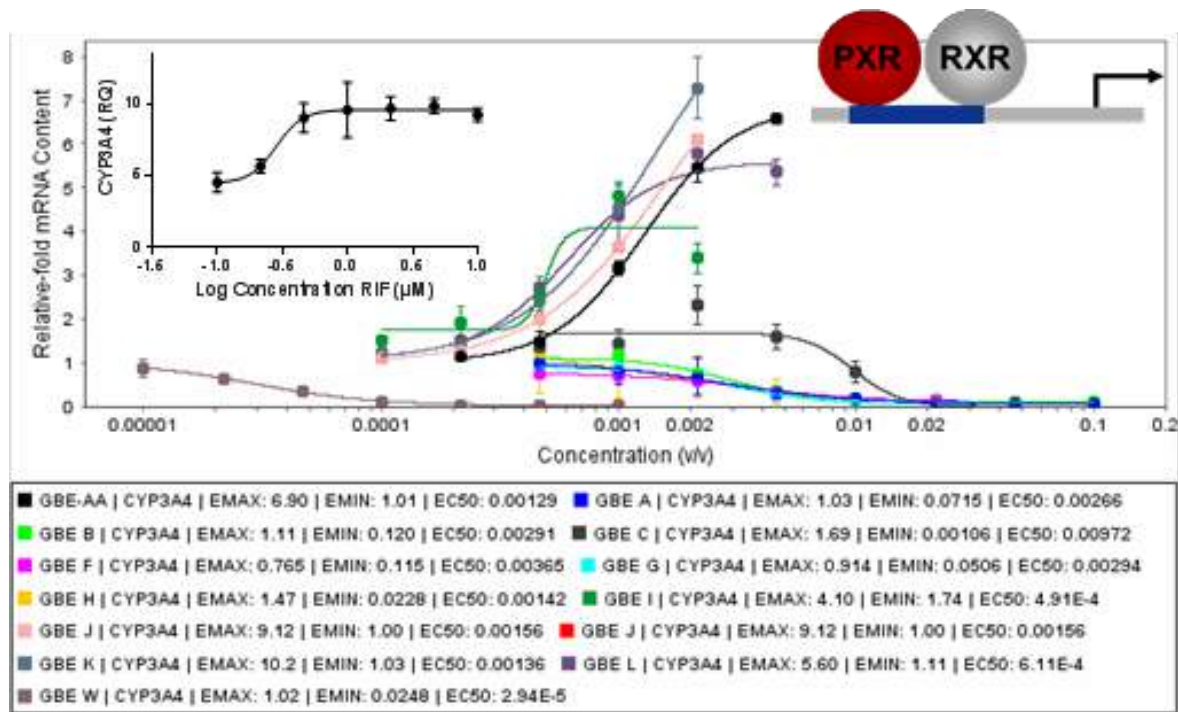
■ GBE 1 ATP EMAX: 1690000 EMIN: 1880 EC50: 0.0264	■ GBE 1F ATP EMAX: 1580000 EMIN: 5520 EC50: 0.0275	■ GBE X ATP EMAX: 1480000 EMIN: 8280 EC50: 0.0581	■ GBE Z ATP EMAX: 1730000 EMIN: 122000 EC50: 0.0510
■ GBE S ATP EMAX: 1410000 EMIN: 830 EC50: 0.00969	■ GBE T ATP EMAX: 1440000 EMIN: 269 EC50: 0.0248	■ GBE U ATP EMAX: 1540000 EMIN: 8.00E3 EC50: 0.0184	



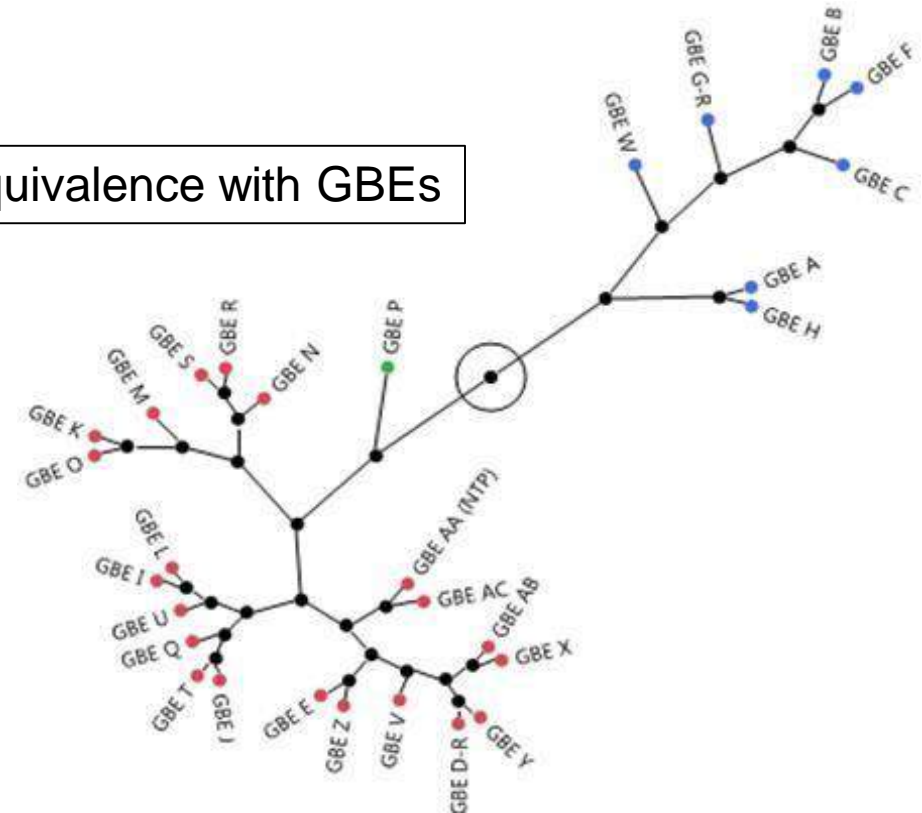


Linking GBE Constituents to Cytotoxicity via ATP Depletion in Primary Cultures of Human Hepatocytes





Phytoequivalence with GBEs



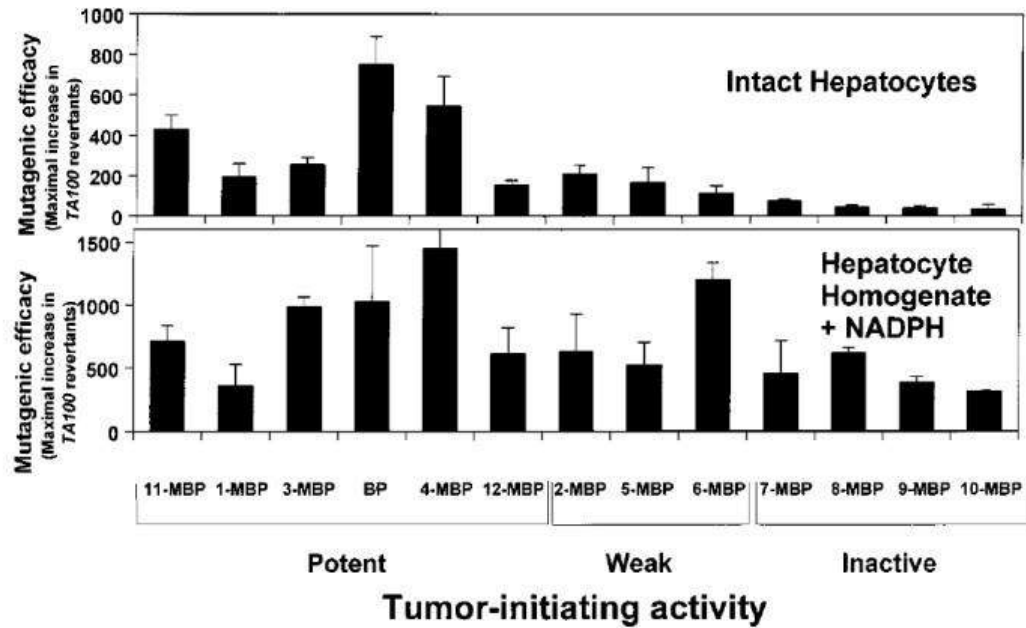


FIG. 1. Mutagenicity of benzo[*a*]pyrene (BP) and the 12 monomethylbenzo[*a*]pyrenes (MBP) using intact hepatocytes and NADPH-fortified hepatocyte homogenate as metabolizing systems. (From Ref. 1.)

Benchmarking Metabolic ‘Competence’

- **Specific Activity Assays** of major human drug metabolizing enzymes
- **Metabolic Clearance Assays** with high, moderate, & low turnover compounds
- **Metabolite Profiling Assays** with orbitrap LC-MS (HRAM)

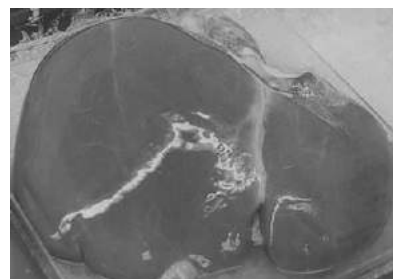
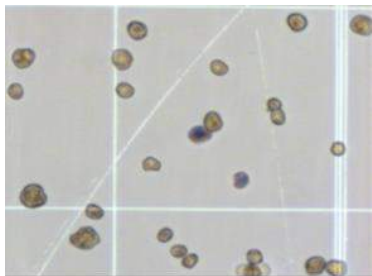
Modeling Metabolically-dependent Liver Toxicity

- Intact hepatocytes better distinguish potent tumor initiators from non-carcinogens (Ames)
- Short $t_{1/2}$ limits utility of PHH suspensions, lower activities & longevities limit SC-PHHs
- How “much” xenobiotic metabolism competence is sufficient?
- What about neonatal/developmental susceptibilities

TABLE 5
Separation of Database into Low, Intermediate, and High Clearance Chemicals

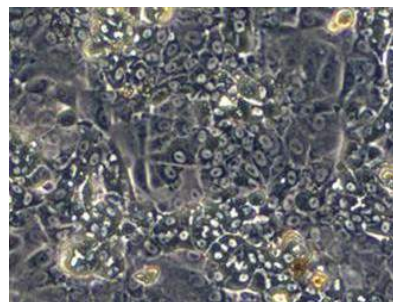
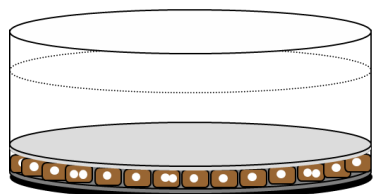
Age group	Ratio of child to adult $t_{1/2}$		
	Low clearance chemicals ^a	High clearance chemicals ^b	High + intermediate clearance ^c
Premature neonates	not available	3.34 ± 1.27	4.18 ± 1.26
Full-term neonates	3.40 ± 2.18 ^d	1.99 ± 0.67	2.38 ± 0.59
1 week–2 months	4.34 ± 0.62 ^e	1.85 ± 0.38	1.96 ± 0.41
2–6 months	1.25 ± 0.31	0.90 ± 0.26	0.94 ± 0.28
6 months–2 years	0.57 ± 0.16	0.26 ± 0.12	0.52 ± 0.14
2–12 years	0.60 ± 0.11	0.72 ± 0.24	0.72 ± 0.10

Suspension PHHs
isolated from human liver

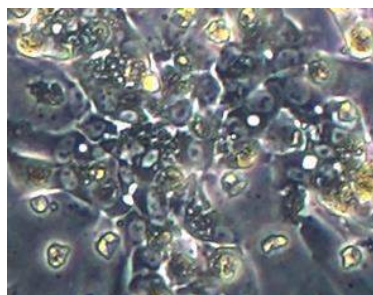


2D Differentiated
HepaRG

SC-PHHs



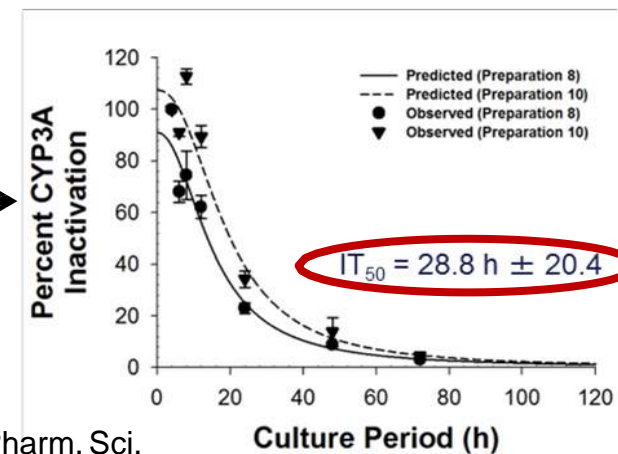
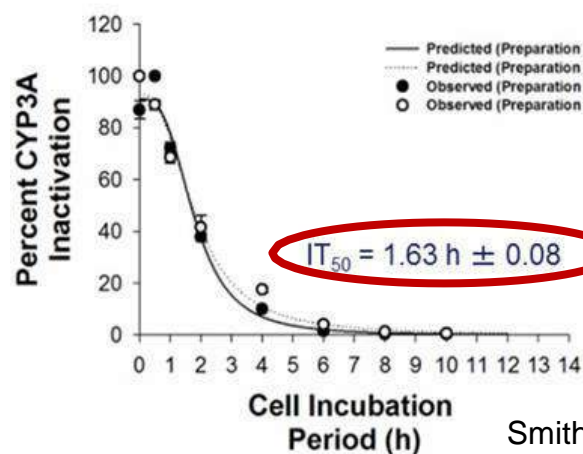
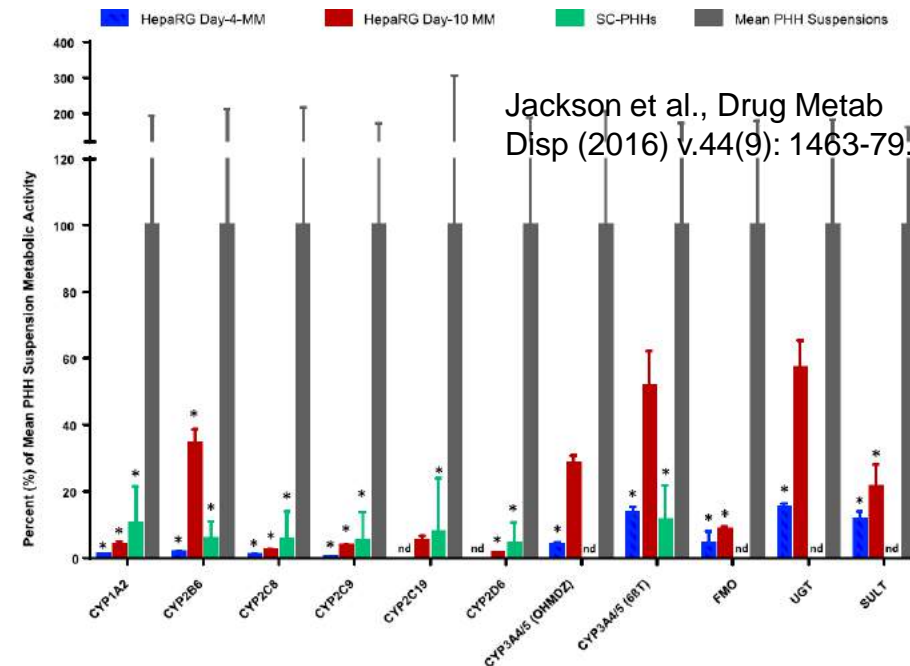
Day-4 HepaRG



stem cells,
transformed cell lines

Defining Xenobiotic Metabolism "Competence"

How much is needed, and over what time frame?



Smith et al. J. Pharm. Sci. 2012. v.101(10):3898.

To receive the rest of the presentation, please get in touch at info@hansonwade.com quoting PREDiCT: 3D Tissue Models Summit. We'll be delighted to send you the full slide deck asap.

We look forward to hearing from you!

Kind regards,

The PREDiCT Team

