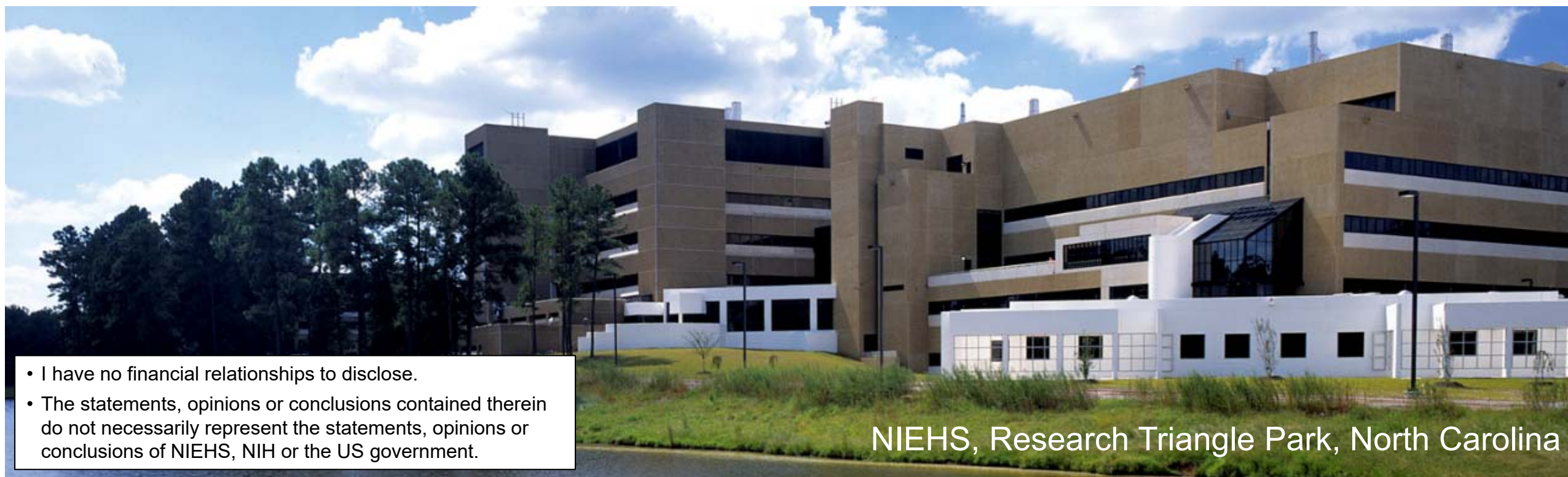


The Evolution of Tox21: Enhancing Physiological Relevance & Interpretability with Emerging Toxicological Approach Methods (TAMs)

Stephen S. Ferguson, Ph.D.



- I have no financial relationships to disclose.
- The statements, opinions or conclusions contained therein do not necessarily represent the statements, opinions or conclusions of NIEHS, NIH or the US government.

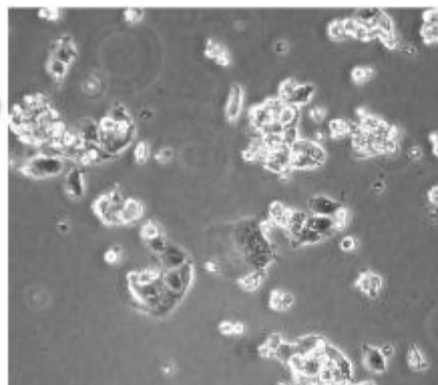
NIEHS, Research Triangle Park, North Carolina

- Numerous research programs addressing public health challenges
- EPA has >85,000 chemicals listed in Toxic Substances Control Act (TSCA)
- Tox21 Program (NIEHS, US EPA, NCATS, & FDA)
 - Address data-poor chemicals in context with pharmaceuticals and well-studied environmental chemicals
 - Prioritize chemicals for further study
 - ~9,000 chemicals evaluated
 - ~70 high-throughput assays
 - >125 biological targets/processes
 - >120 million data points
 - Publicly Available Resources:
 - PubChem
 - Tox21 Data Browser
 - EPA CompTox Chemicals Dashboard
 - Quantitative potency estimation through in vitro to in vivo extrapolation (IVIVE)
- Emerging technologies are providing tools estimate human bioactivity & toxicity potential using human cells and mechanistic signaling pathway interactions

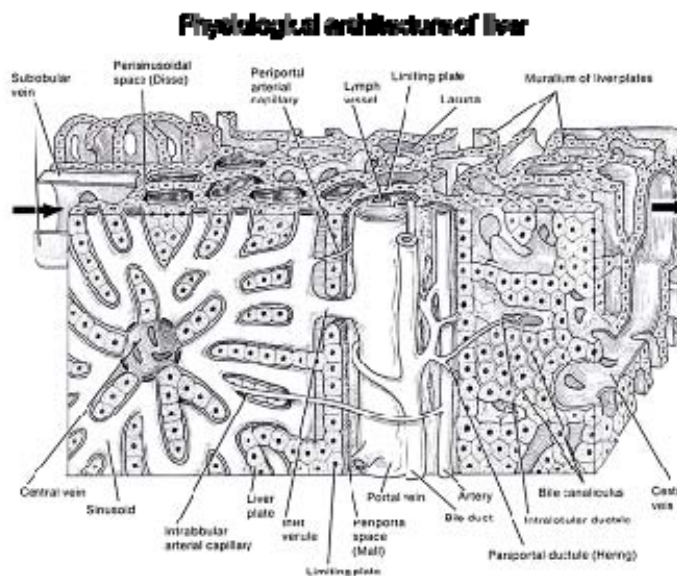


<https://tox21.gov/overview/about-tox21/>

In the beginning,
there were HepG2.



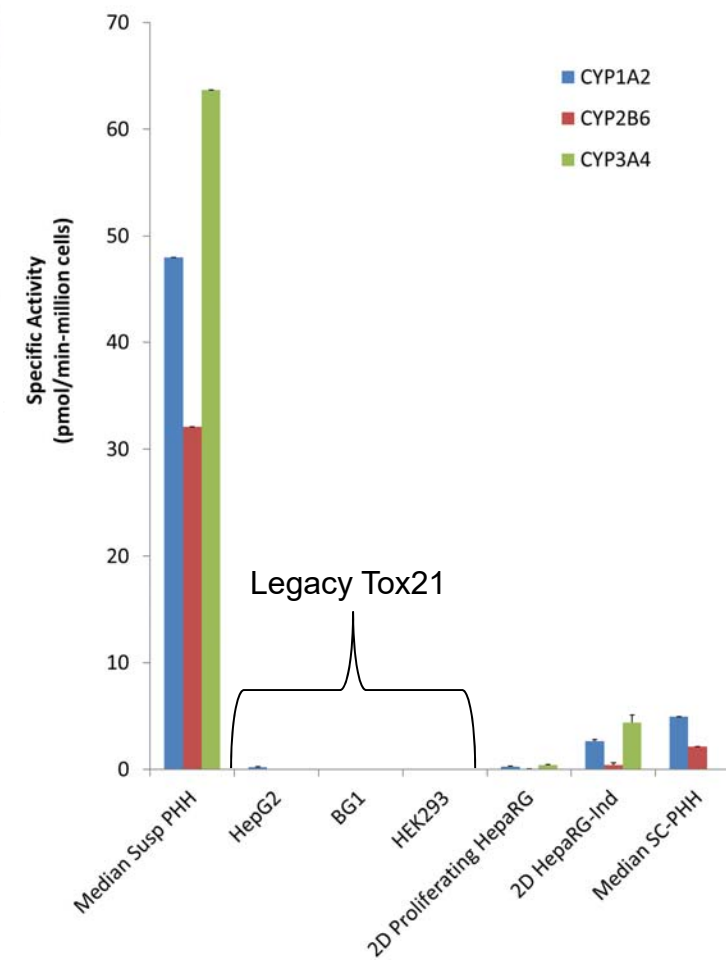
ATCC Number: **HB-8065**
Designation: **Hep G2**



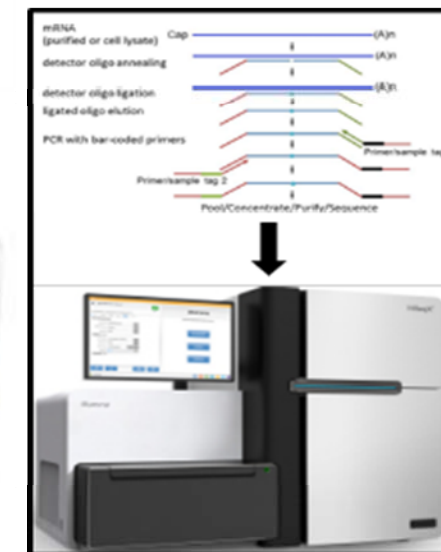
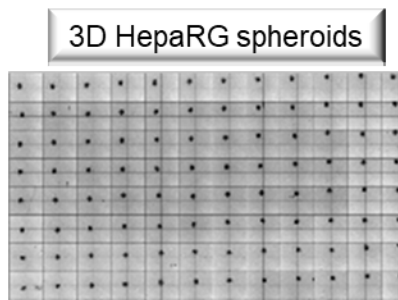
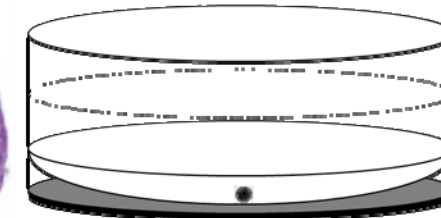
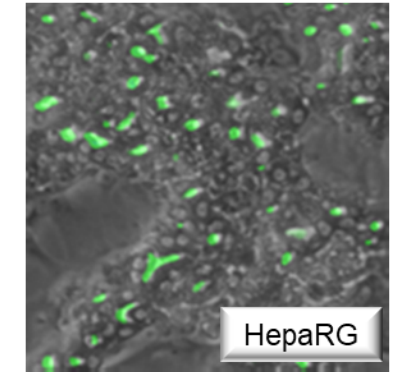
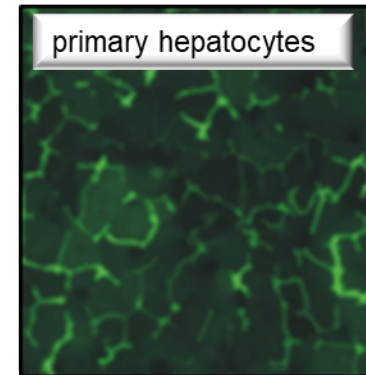
Physiological Architecture of Liver (Karnes, 1977; Karnes & Gundersen, 1979)

Limitations of Tox21 qHTS:

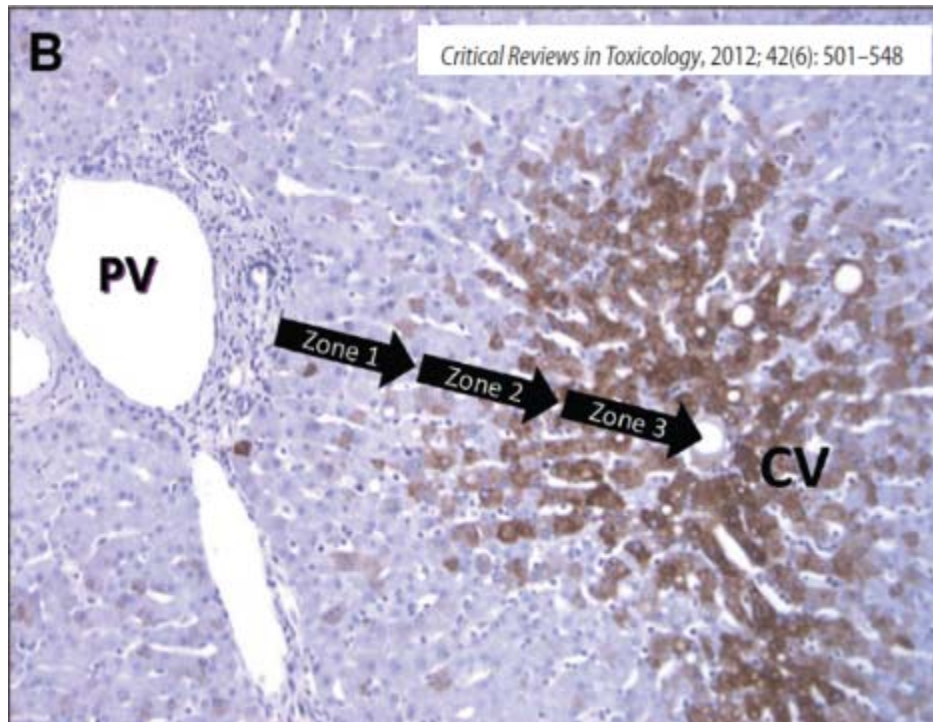
- Limited capability for xenobiotic metabolism
- Limited pathway coverage
- Focus on ‘individual’ cellular pathways lacking integrated biological/tissue-like functionality
- Use of immortalized and transformed cell lines
- Addition-only assays with ≤ 40 h exposure
- linking chemicals to AOPs, pathologies, and disease



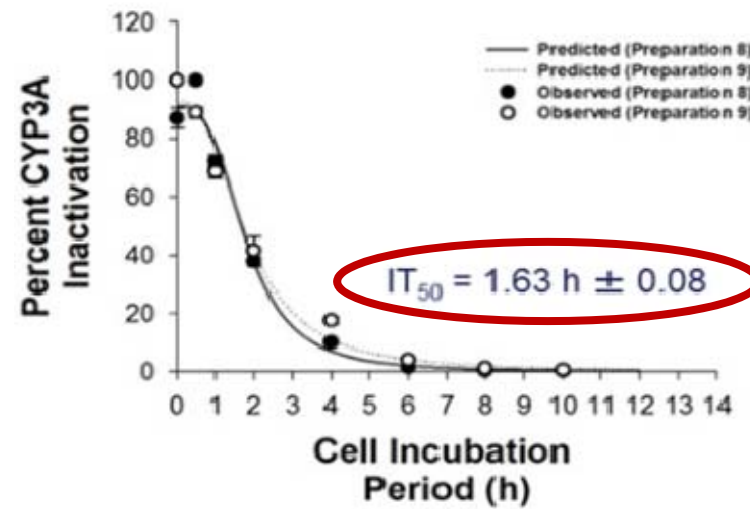
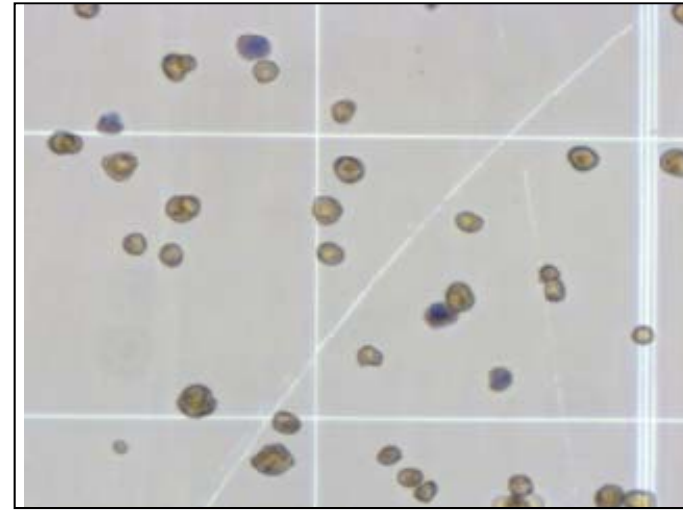
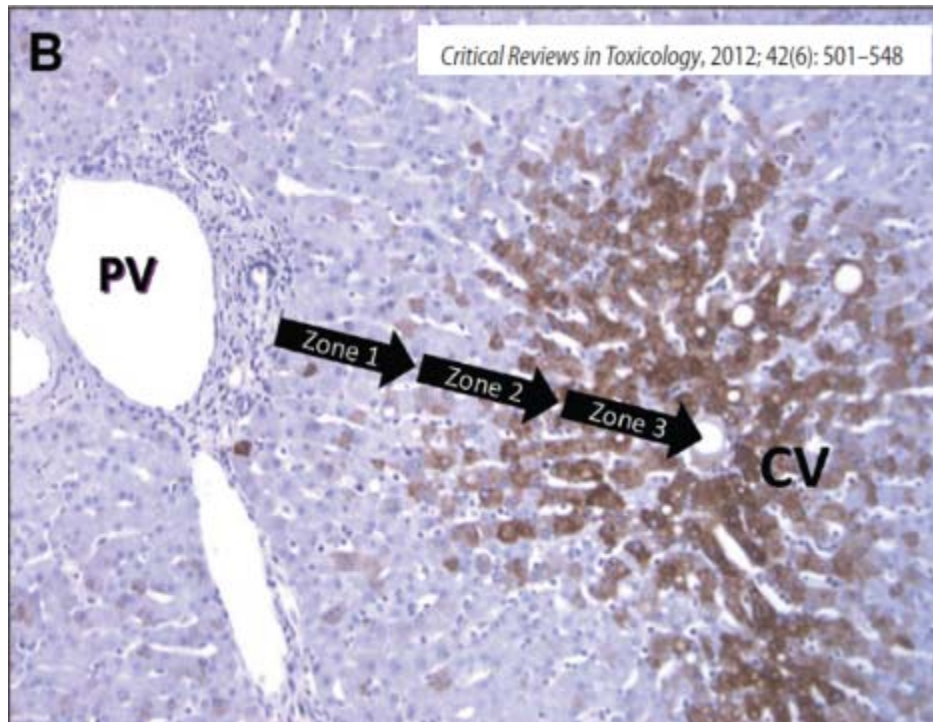
- Physiologically-relevant in vitro screening models
 - improved cellular differentiation/functionality
 - xenobiotic metabolism & bioactivation/detoxification
 - longevity to model progressions towards apical outcomes
 - pathological outcomes
 - xenobiotic clearance and drug interactions
 - pharmacology analogue case comparisons
 - species-specific response comparisons
- Multi-dimensional assay platforms (time/concentration)
 - cellular imaging
 - high throughput transcriptomics
 - metabolomics
- Quantitative translation to humans
 - IVIVE (e.g., BMC, AC_{50} , CL_{int} , f_{ub})
- Extend approach:
 - Extrahepatic tissues: kidney, lung, cardiovascular, intestine
 - Susceptibility models: developmental, disease, population



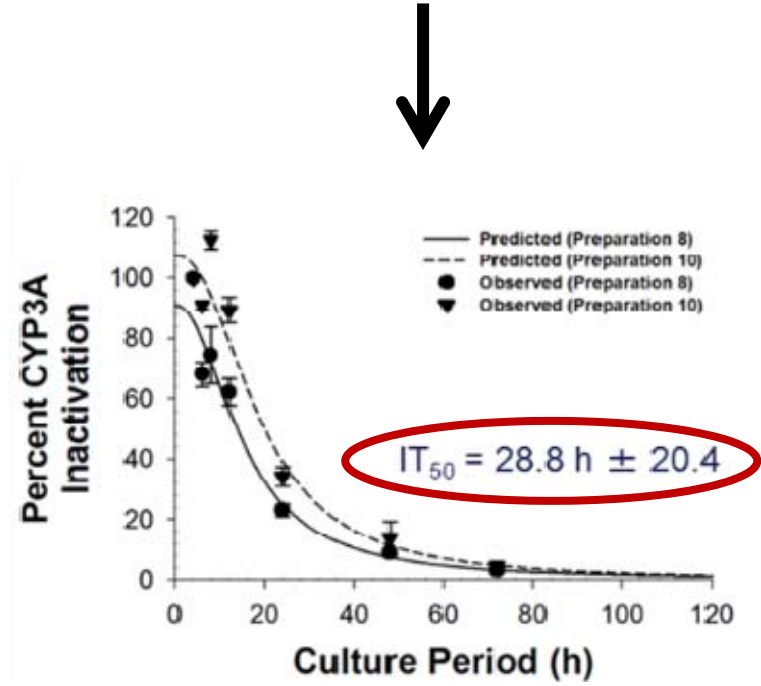
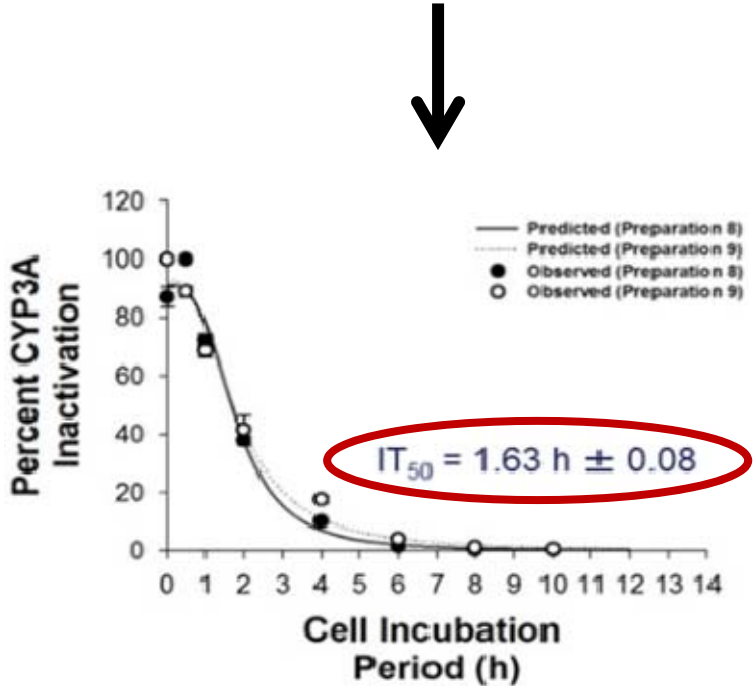
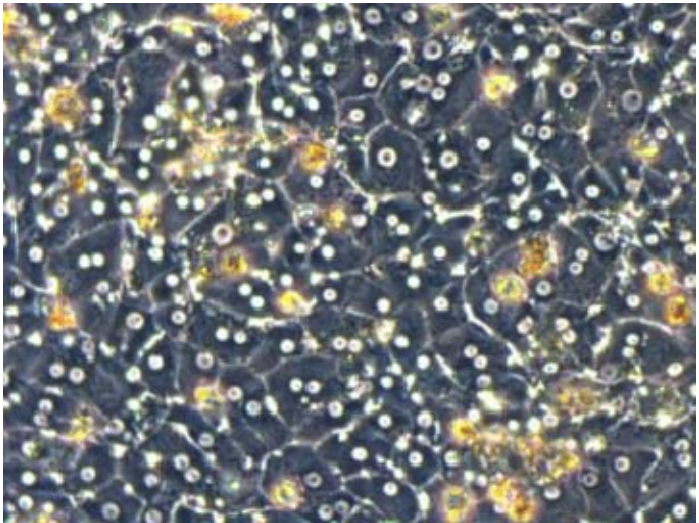
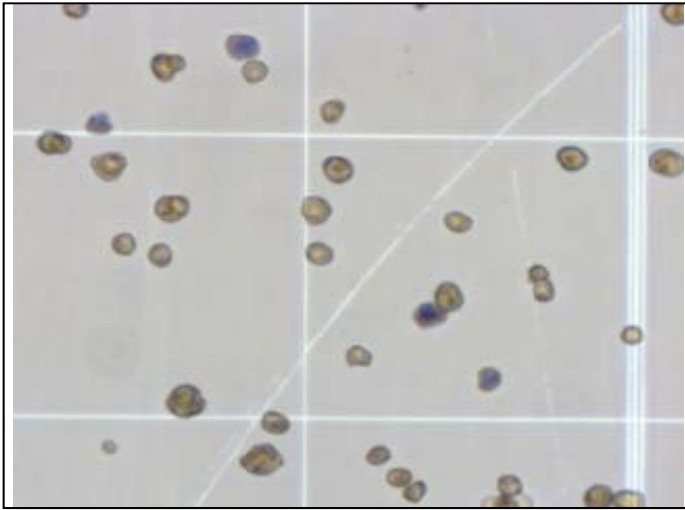
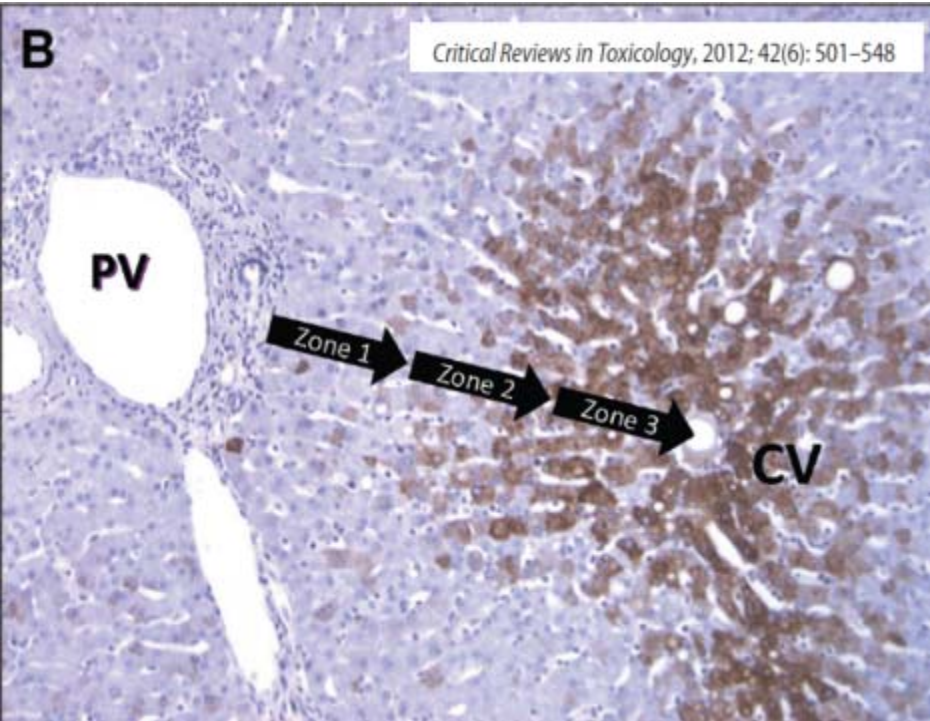
Isolated Primary Liver Cells Rapidly De-differentiate Once Removed from Liver Tissue



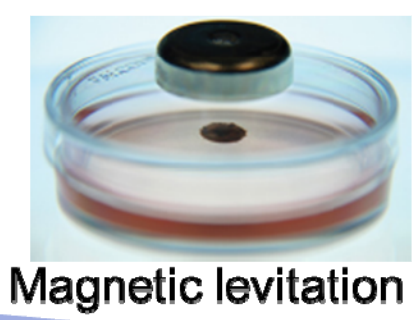
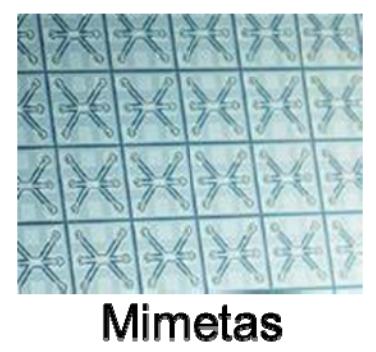
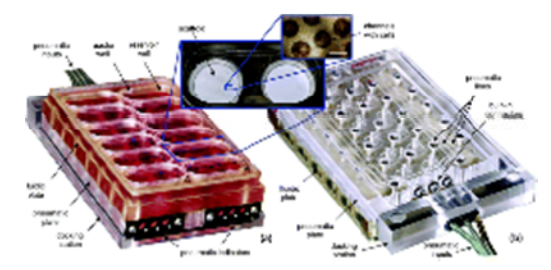
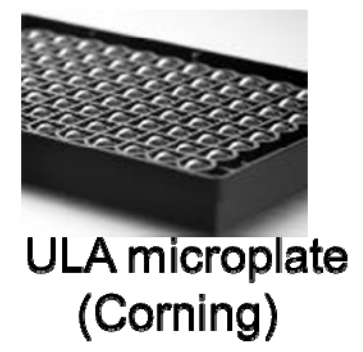
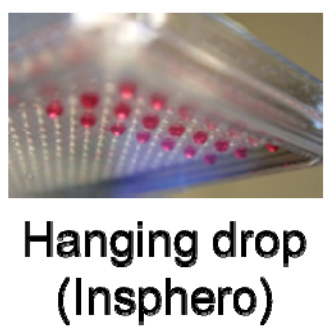
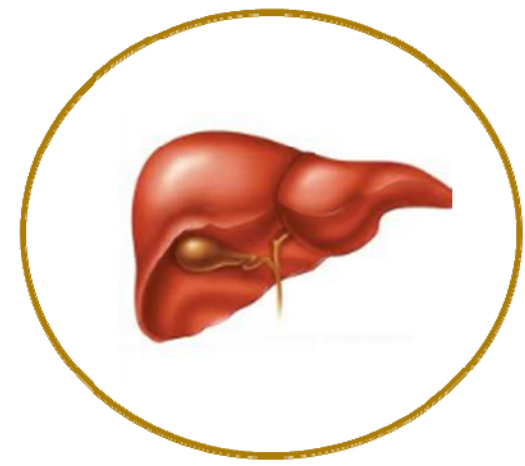
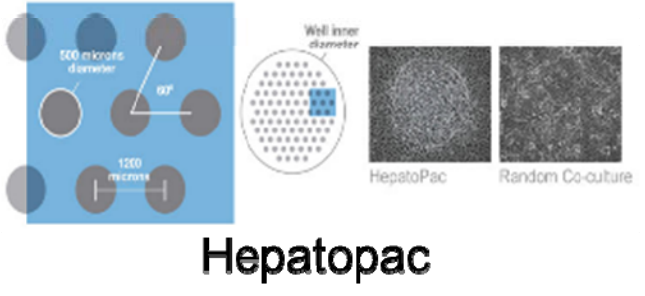
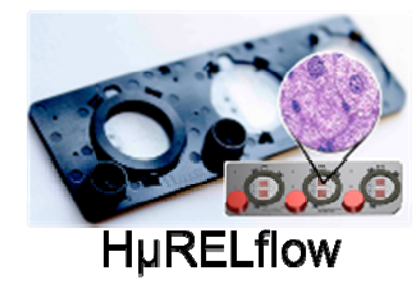
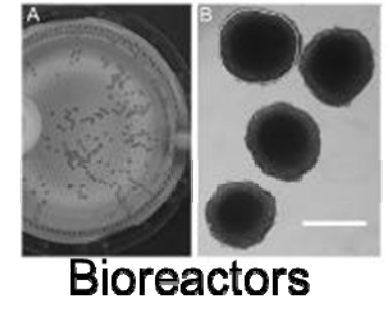
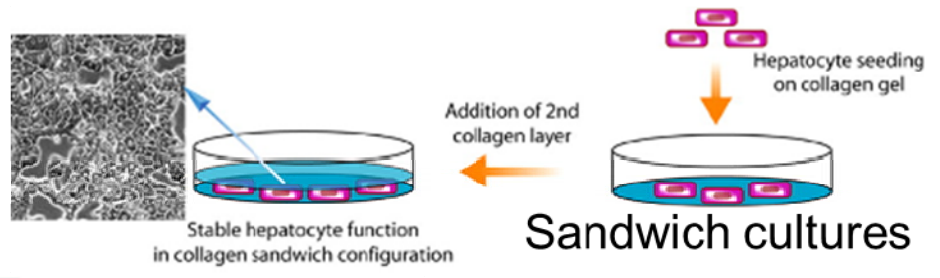
Isolated Primary Liver Cells Rapidly De-differentiate Once Removed from Liver Tissue



Isolated Primary Liver Cells Rapidly De-differentiate Once Removed from Liver Tissue



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



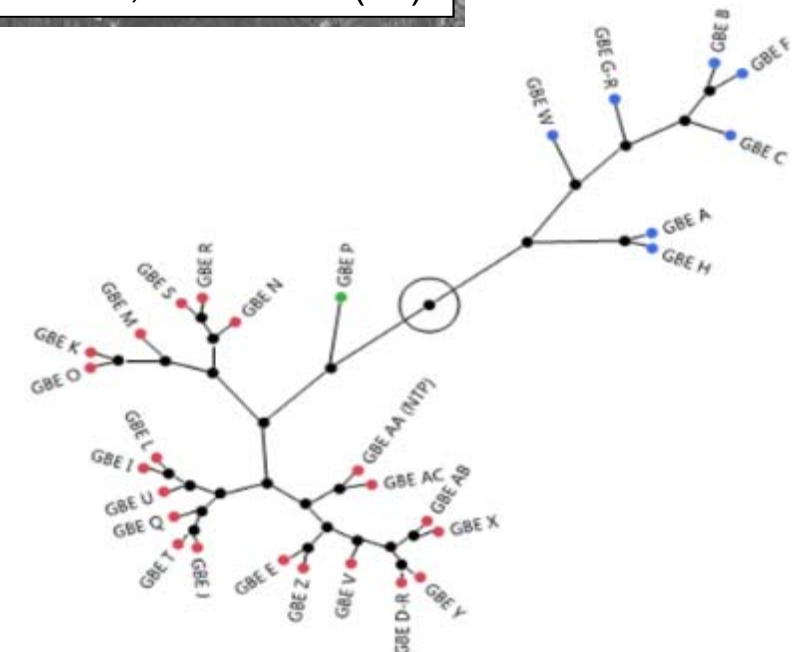
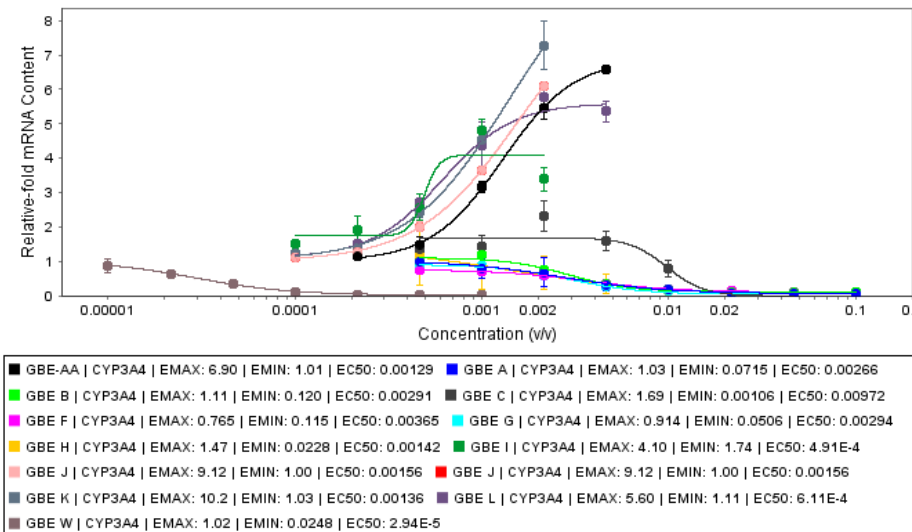
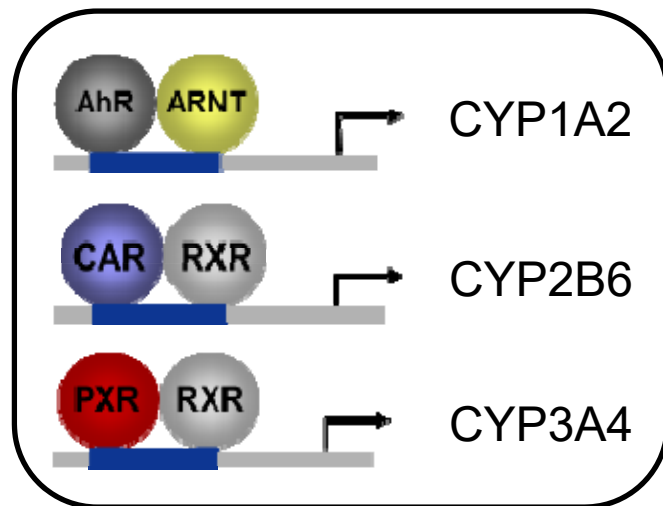
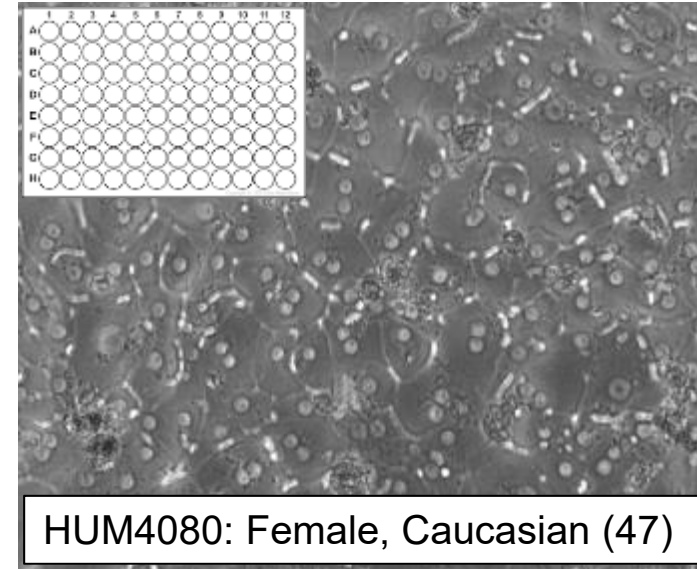
Throughput

- Sandwich cultures of PHHs (SC-PHHs)



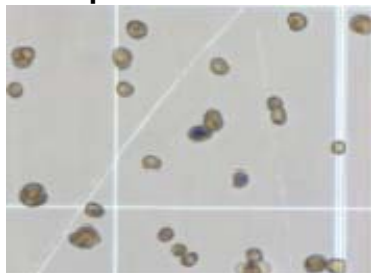
- Modeling liver weight increases, enzyme Induction

- Cell Health
- CYP450 induction/receptor activation
 - mRNA (TaqMan)
 - Liver enzymatic activity
- Linking constituents to biological responses

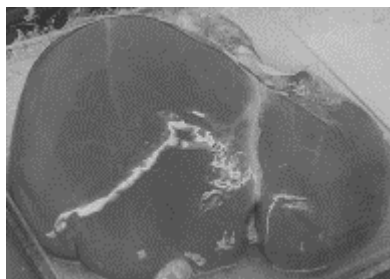


Metabolic Competence

Suspension PHHs



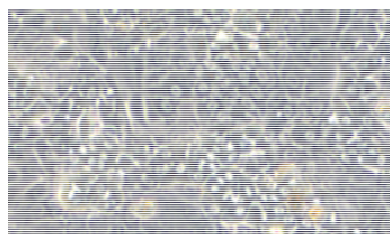
Human Liver



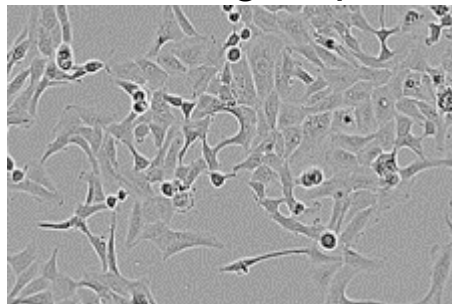
SC-PHHs



2D HepaRG

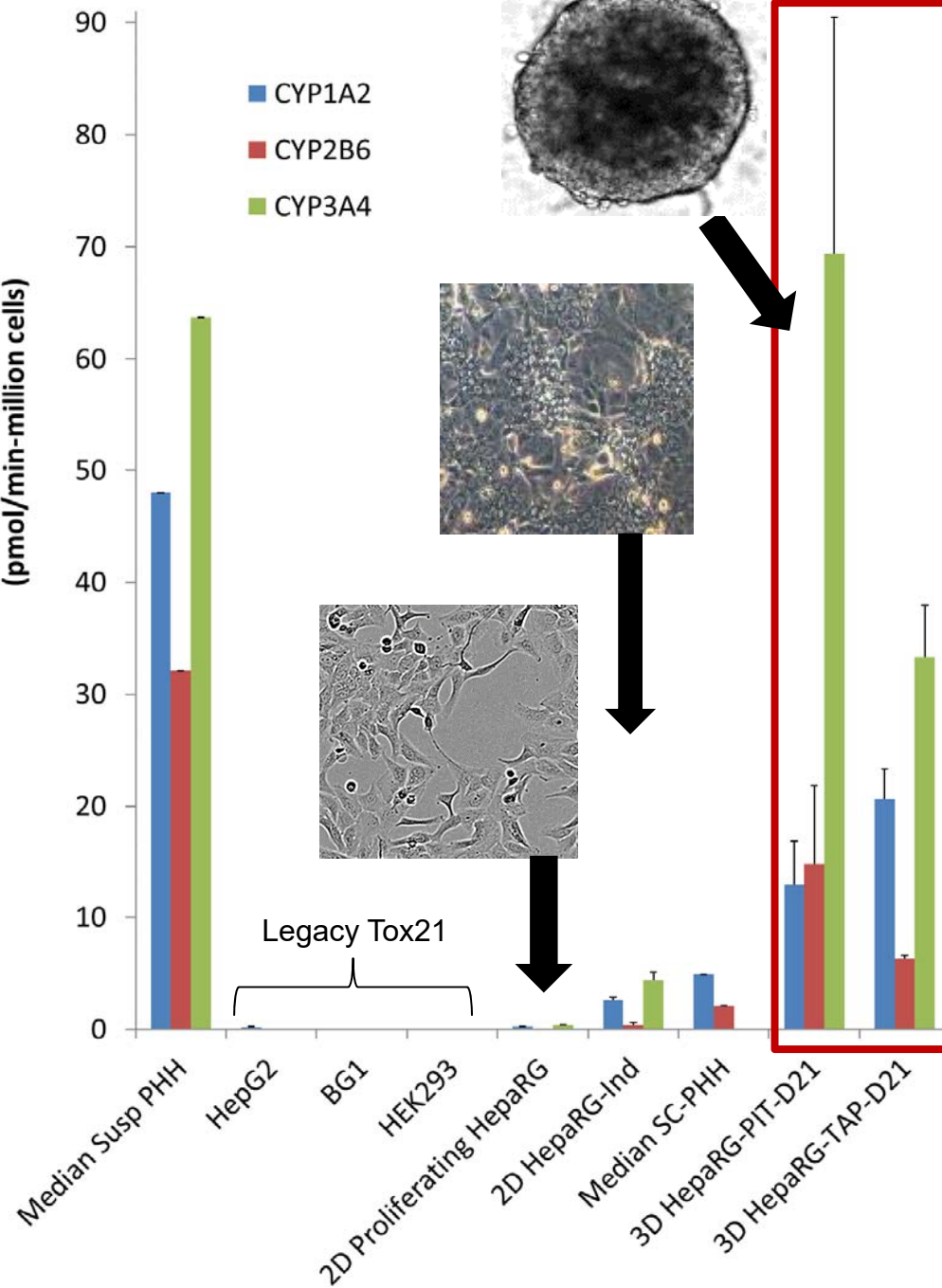


Proliferating HepaRG

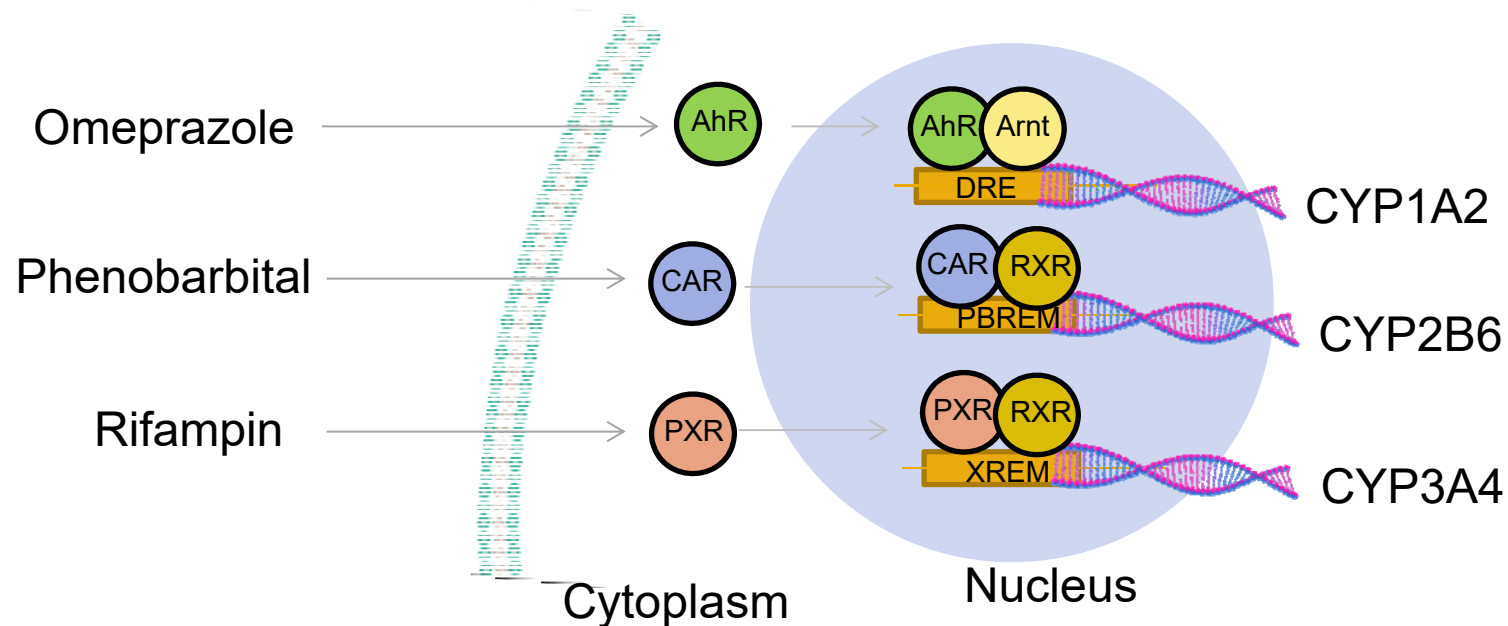
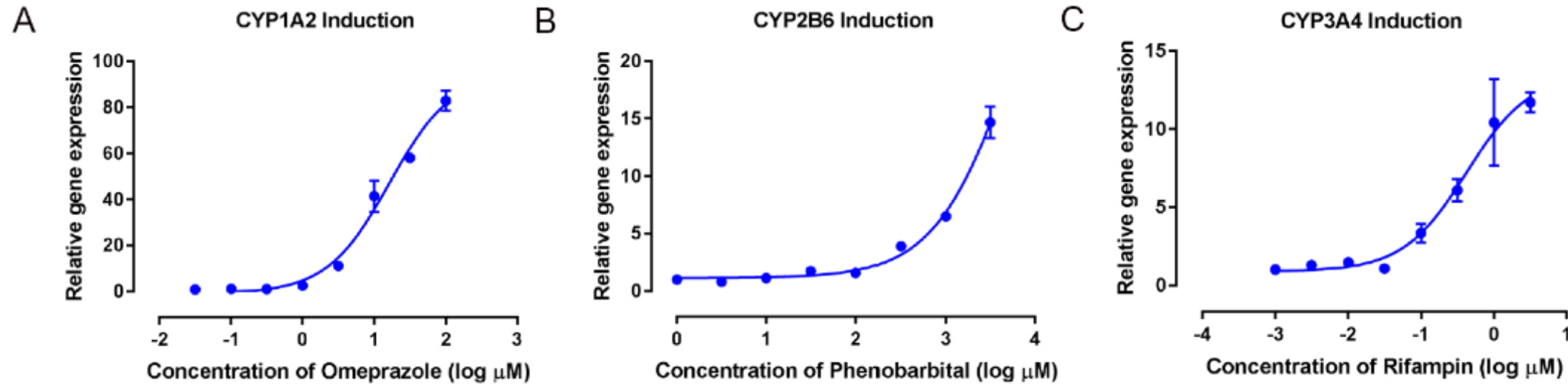


iPSC-derived hepatocytes
Transformed cell lines
(e.g., HepG2)

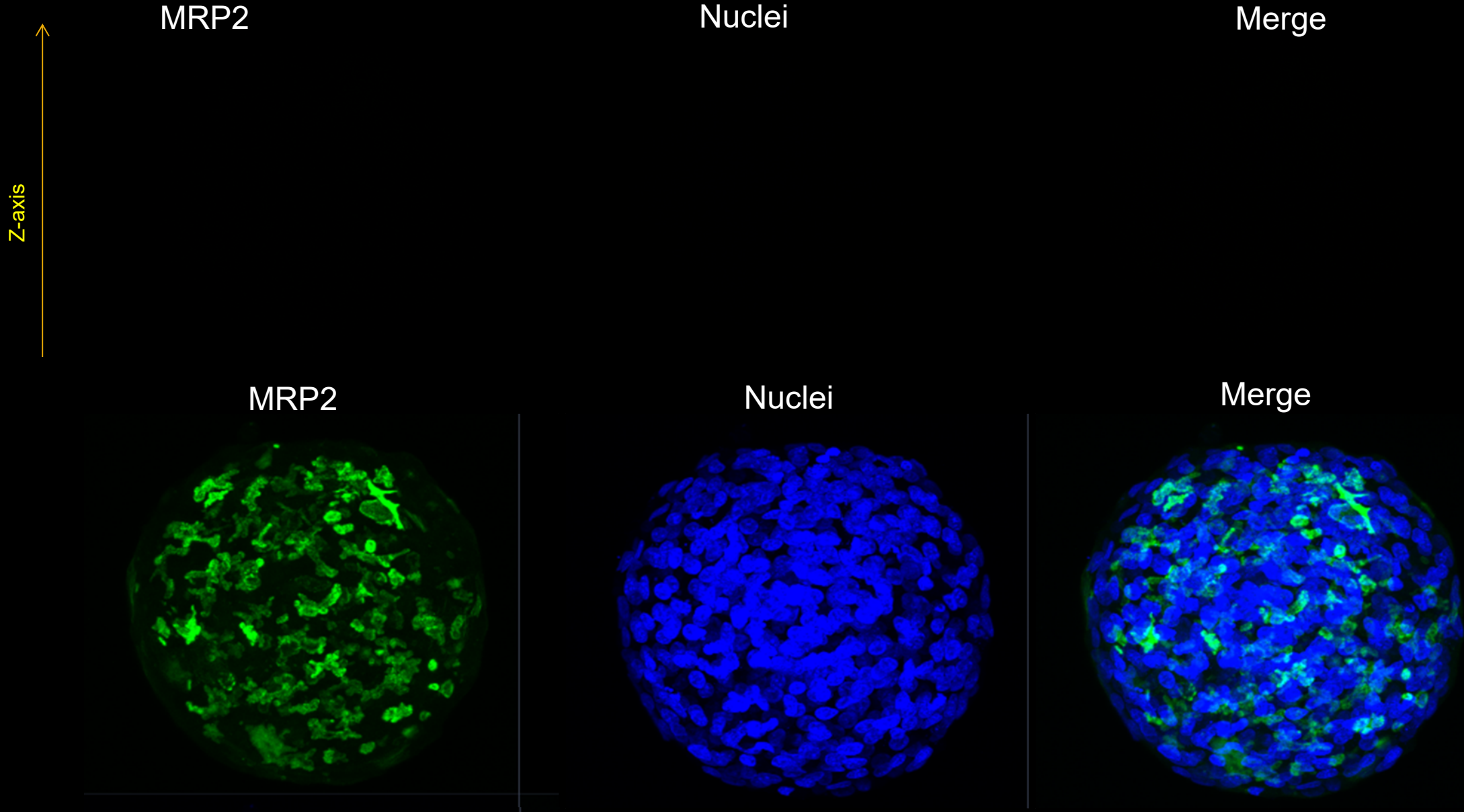
Specific Activity
(pmol/min-million cells)



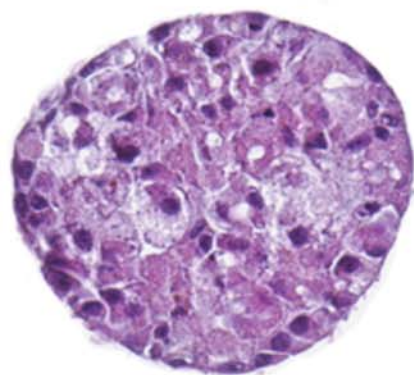
AhR-, CAR-, & PXR-Mediated Liver Enzyme Induction



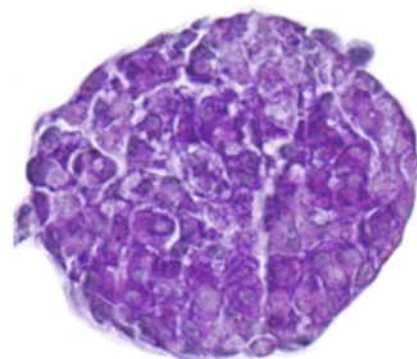
MRP2 Localization in HepaRG Spheroids



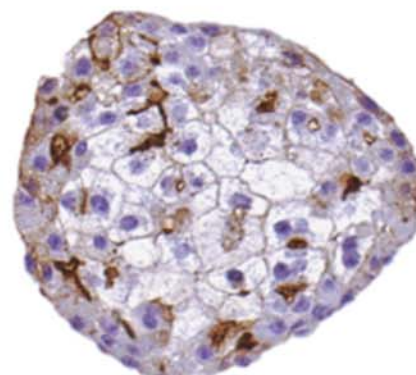
HepaRG cells form polarized spheroids



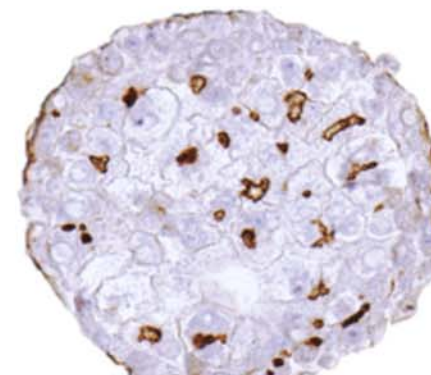
H&E



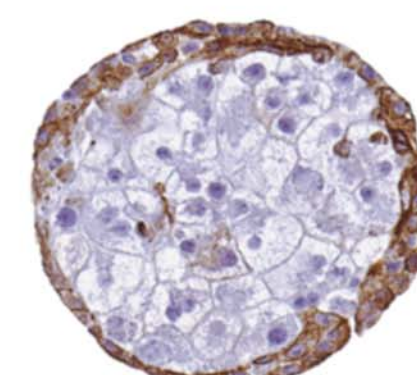
PAS



Poly CEA



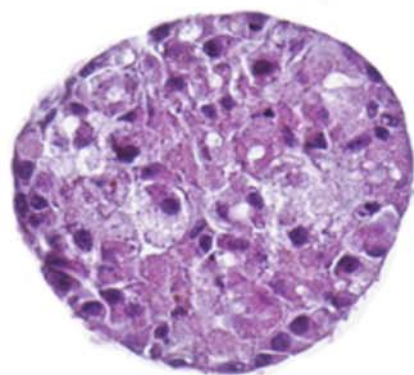
MRP2



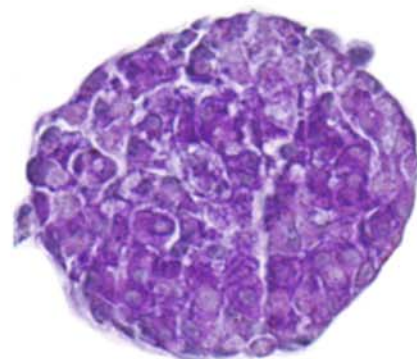
CK19

PAS: Glycogen storage
Poly CEA: Glycoprotein-1 on Bile Canaliculi (BC)
MRP2: Luminal transporter found at BC surfaces
CK19: Marker for Cholangiocytes

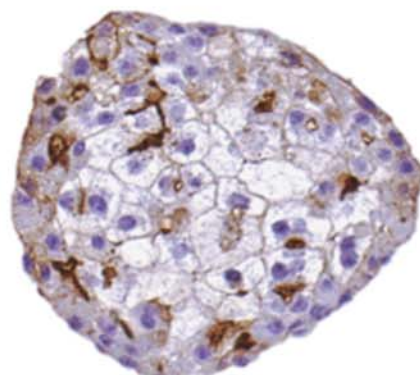
HepaRG cells form polarized spheroids



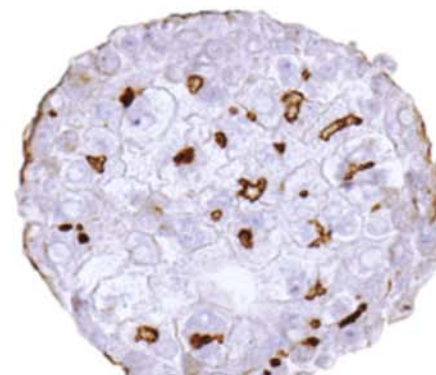
H&E



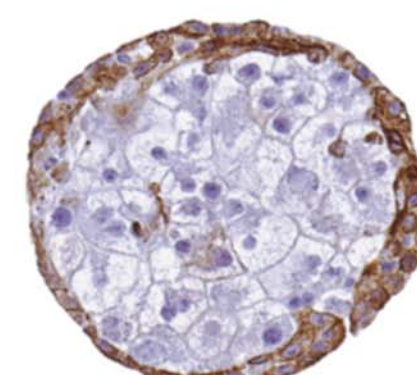
PAS



Poly CEA

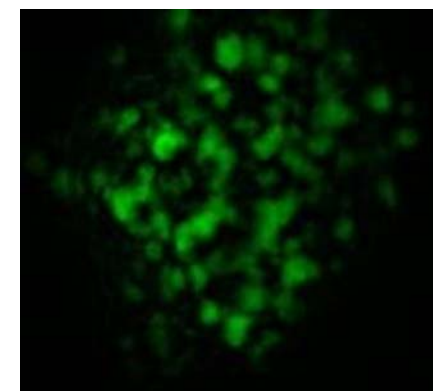


MRP2



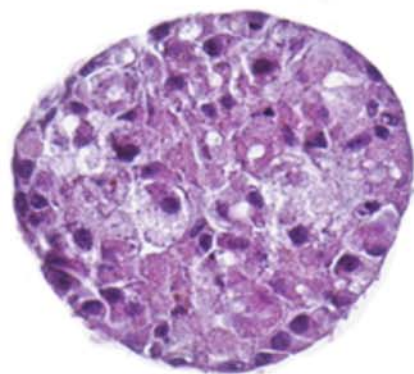
CK19

PAS: Glycogen storage
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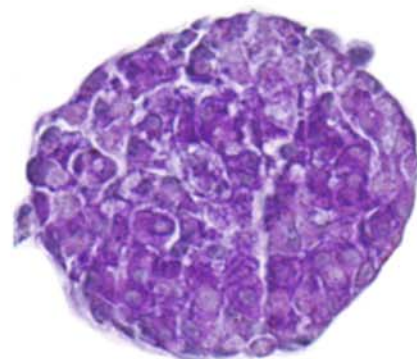


Live-cell CLF

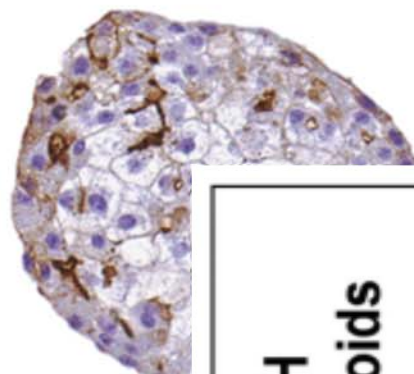
HepaRG cells form polarized spheroids



H&E

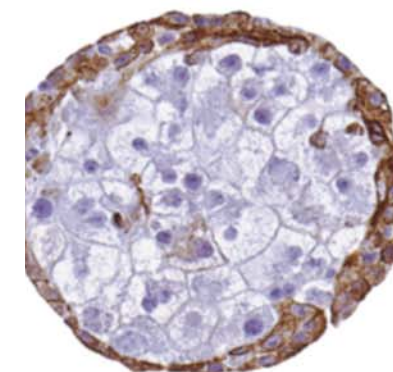


PAS



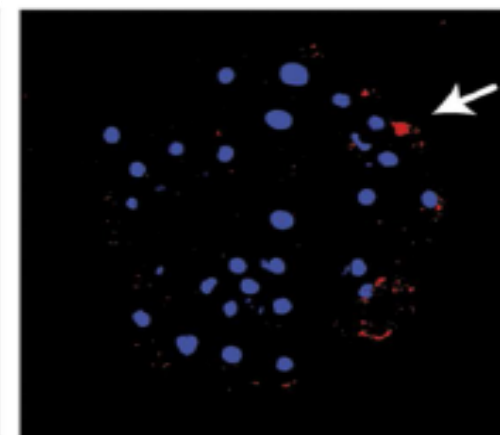
Pc

**CK19
(Biliary cells)**

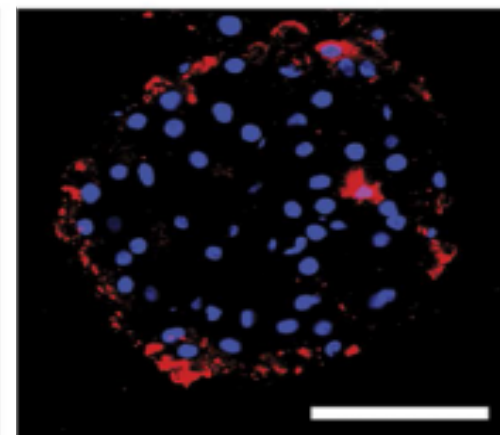


CK19

**PHH
spheroids**



**PHH + NPC
spheroids**

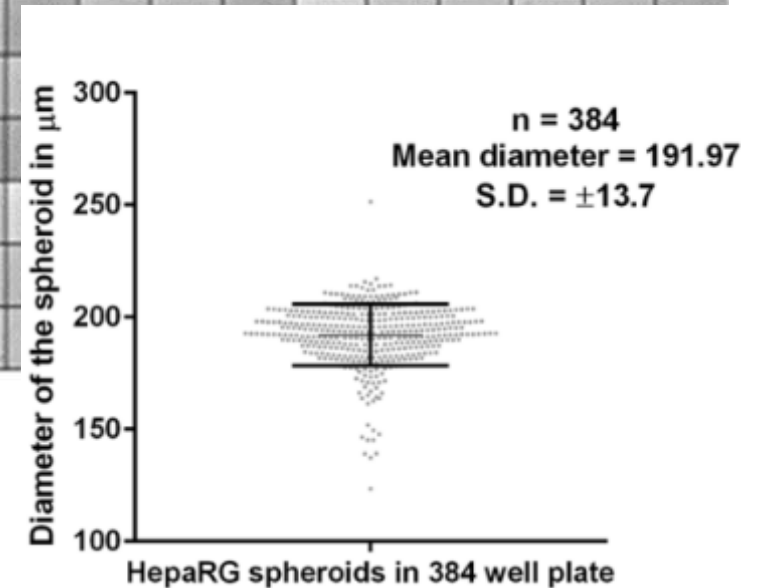
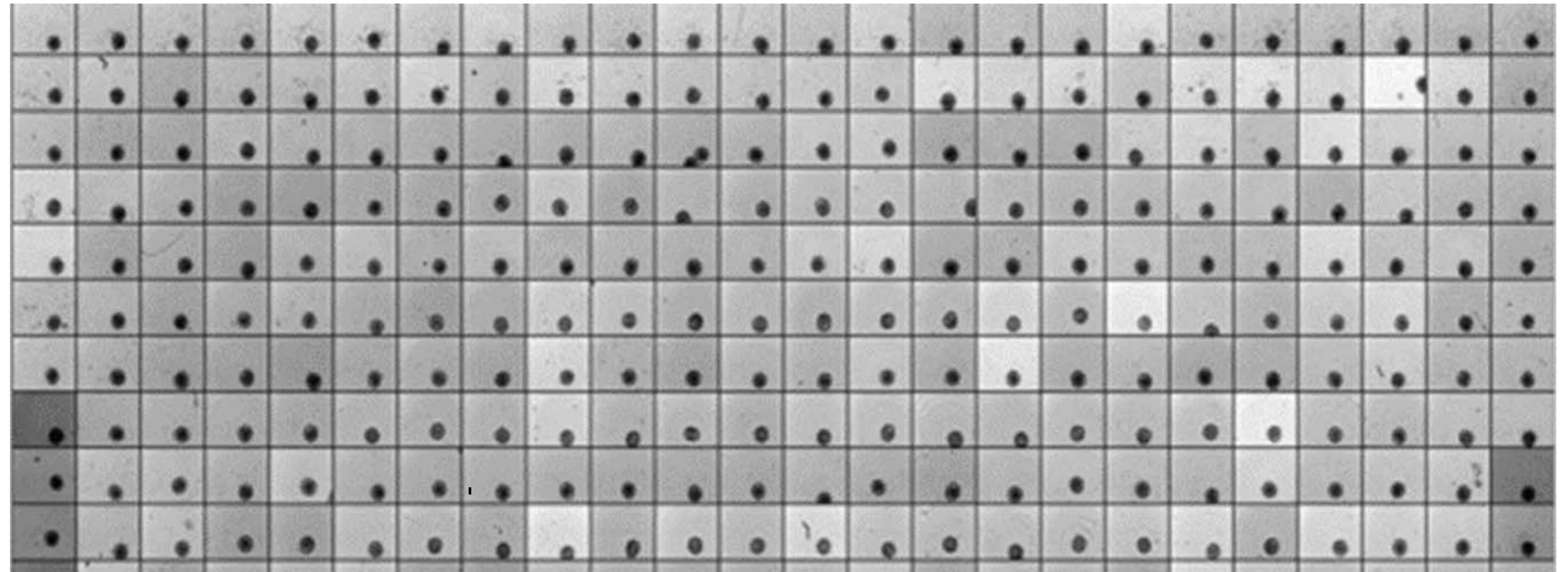


PAS: Glycogen storage
 Poly CEA: Glycoprotein-1 on Bile Canaliculi
 MRP2: Luminal transporter found at Bile Canaliculi
 CK19: Marker for Cholangiocytes

Bell et al., Sci Rep. 2016 May 4;6:25187.



3D HepaRG Spheroids (384-well)

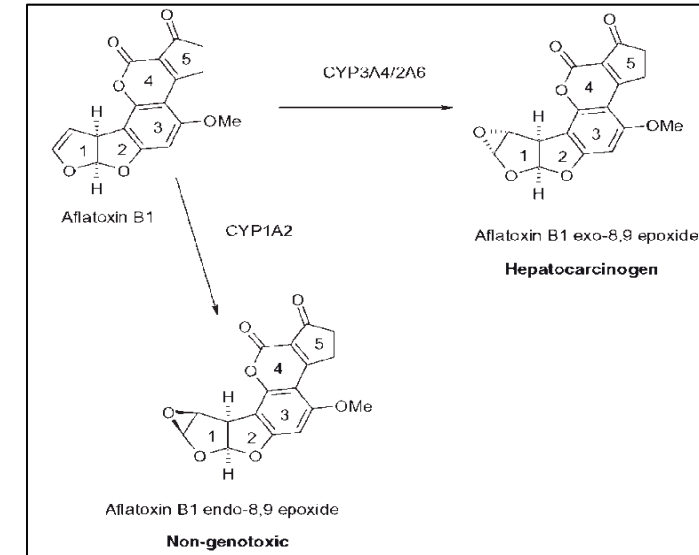
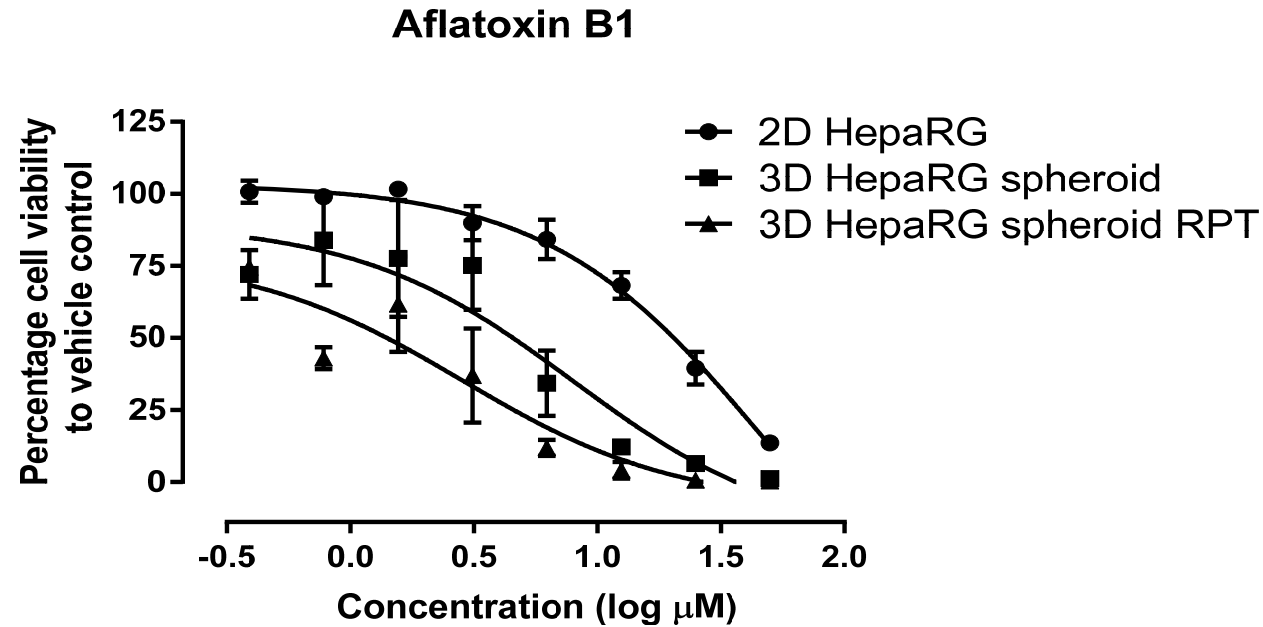


1 vial of 10 million cells
= 1 X 2D 384-well plate
= 12-25 X 384-well plates

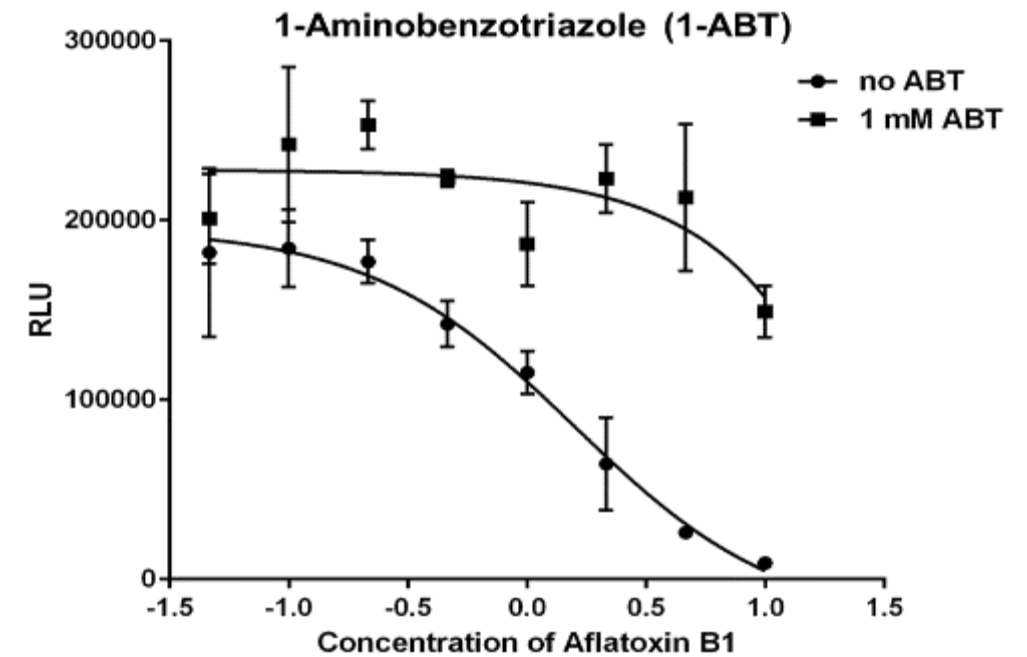


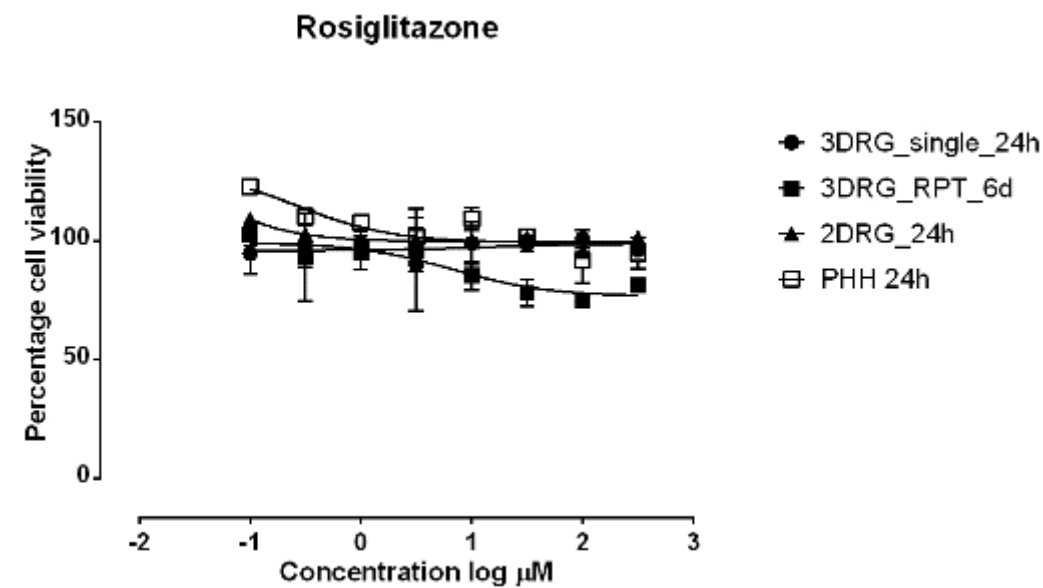
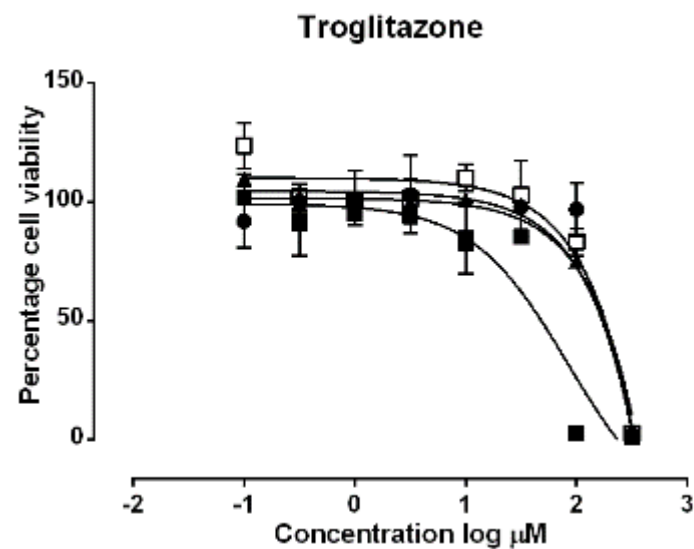
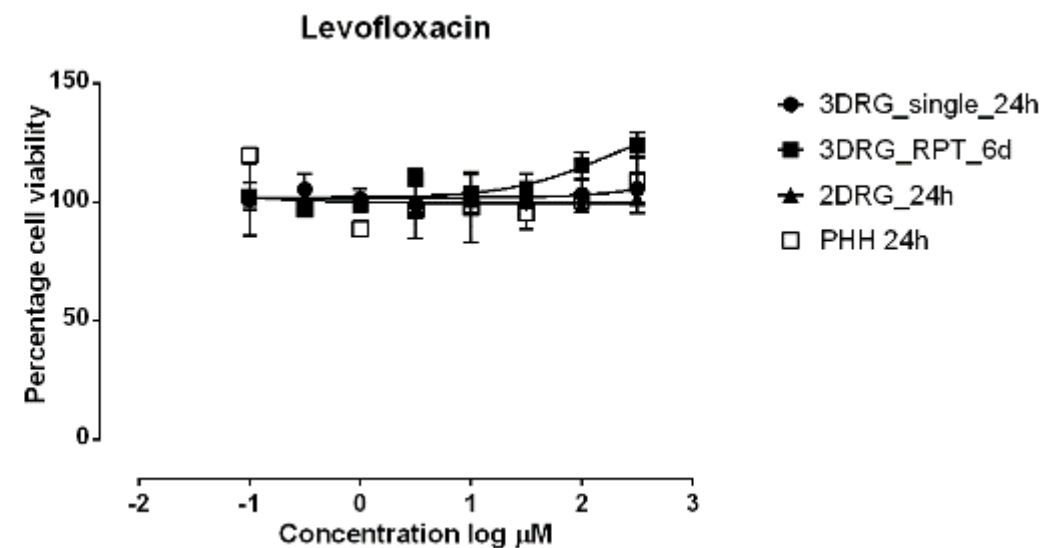
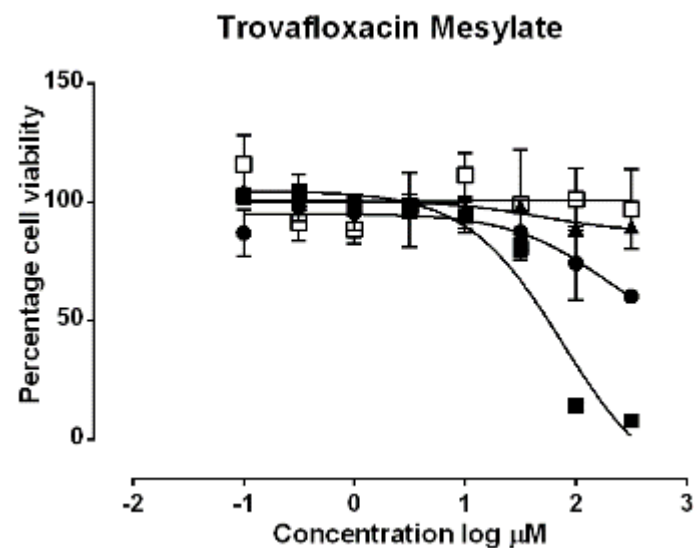
Dr. Sreenivasa Ramaiahgari

From the Cover: Ramaiahgari et al., Toxicol. Sci (2017) v.159 (1): 124-136



	2D HepaRG	3D HepaRG	3D HepaRG repeat exposure
TC ₅₀	46.54 μM	7.94 μM	2.83 μM





- 3 Culture Configurations of HepaRG Cells (384-well formats)

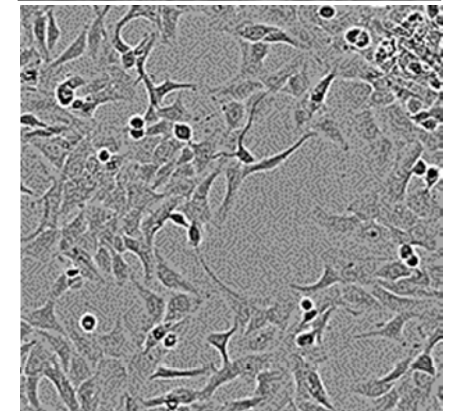
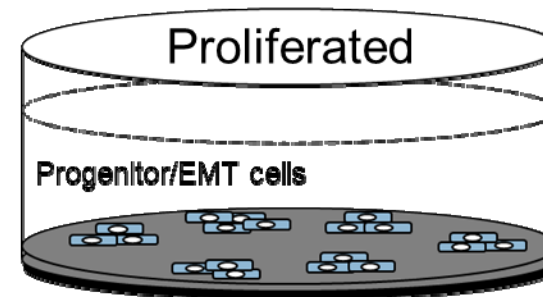
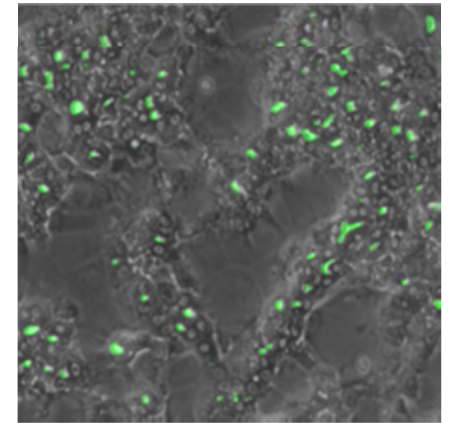
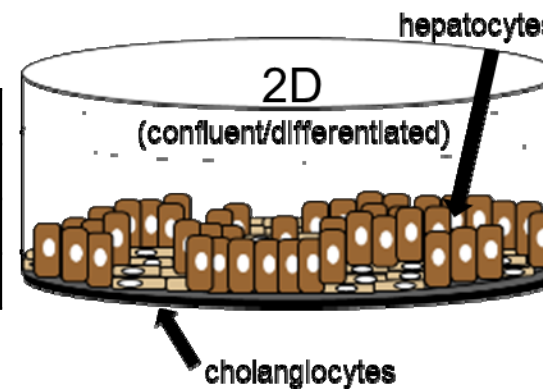
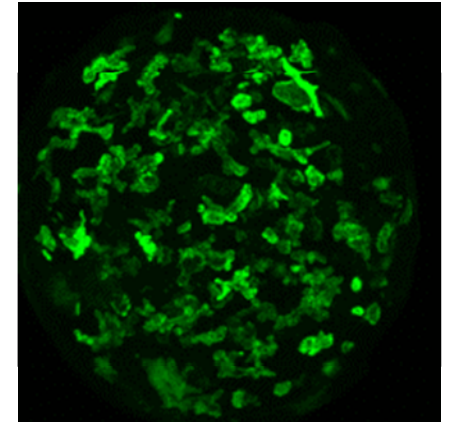
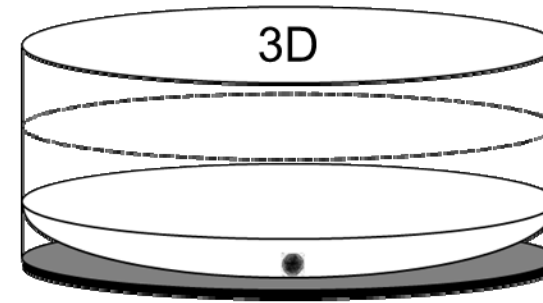
- 24 Compounds

- Liver injury/metabolically-activated toxicity
- Hepatic receptor activators
- Drug analogue comparisons
- 'Negatives' for liver injury

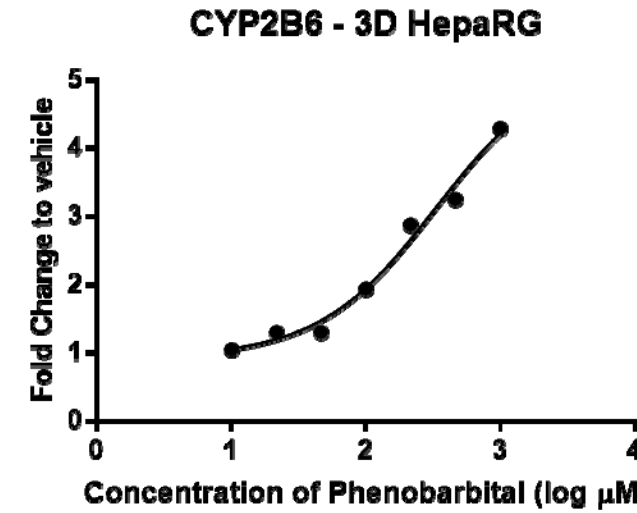
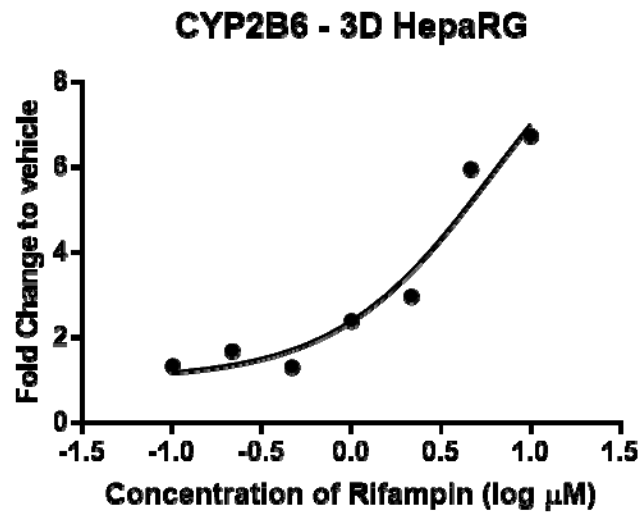
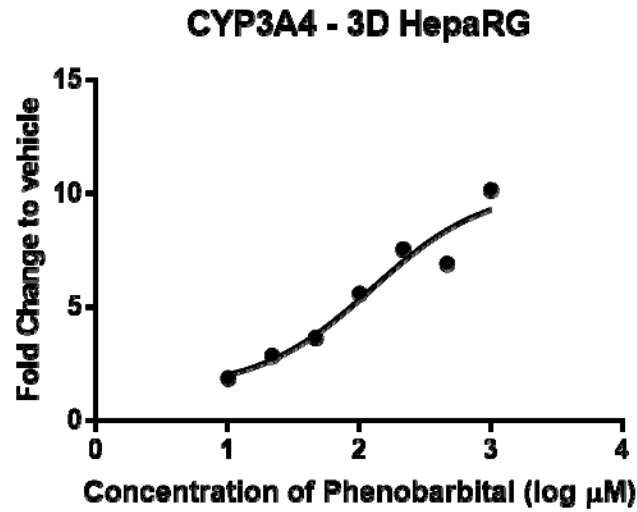
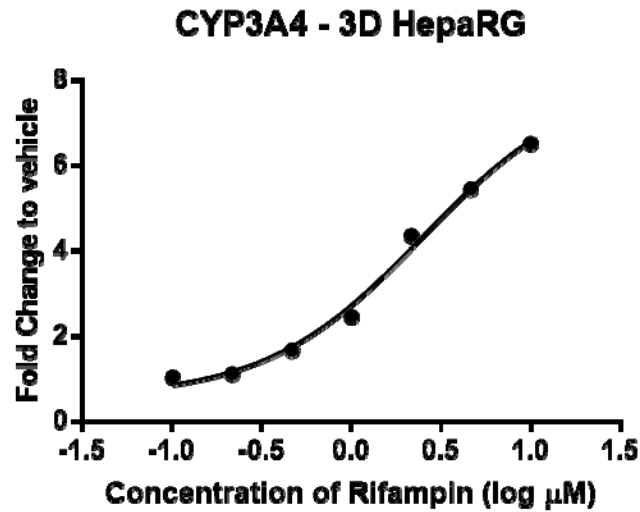
acetaminophen	caffeine	diphenhydramine	DMN	rifampicin	tamoxifen
aflatoxin B1	CDCA	fenofibric acid	omeprazole	ritonavir	troglitazone
aspirin	chlorpromazine	levofloxacin	phenobarbital	rosiglitazone	trovafloxacin
benzo(a)pyrene	cyclophosphamide	menadione	KCl	sucrose	valproic acid

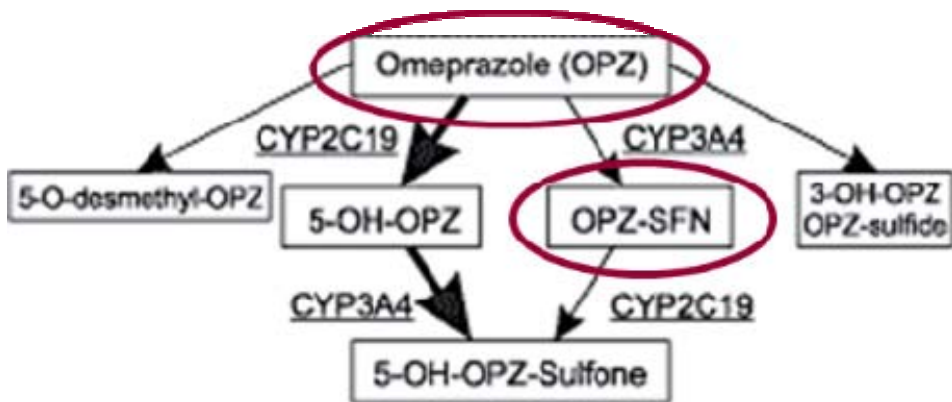
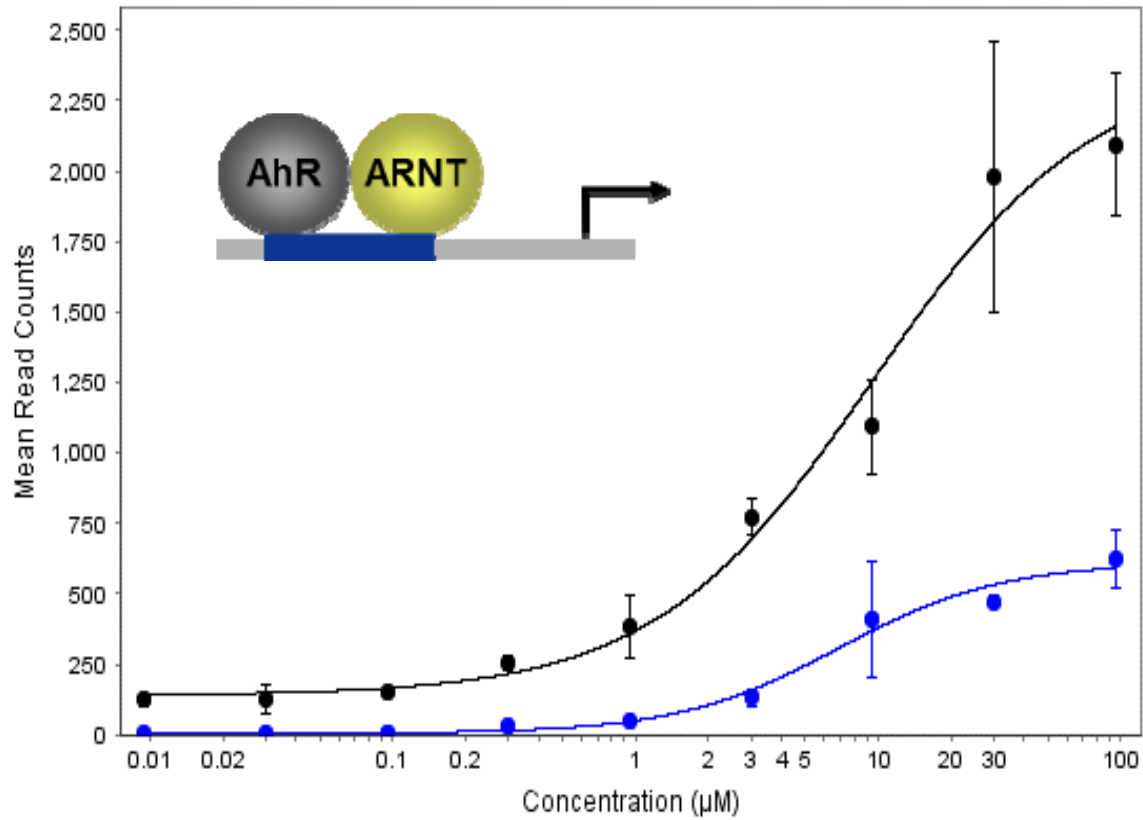
- Assays:

- cell morphology (Incucyte, daily for each culture well)
 - Image classifications, quantitative masking of confluence
- cytotoxicity (LDH leakage)
- high throughput transcriptomics (HTT with S1500+, TempO-Seq)



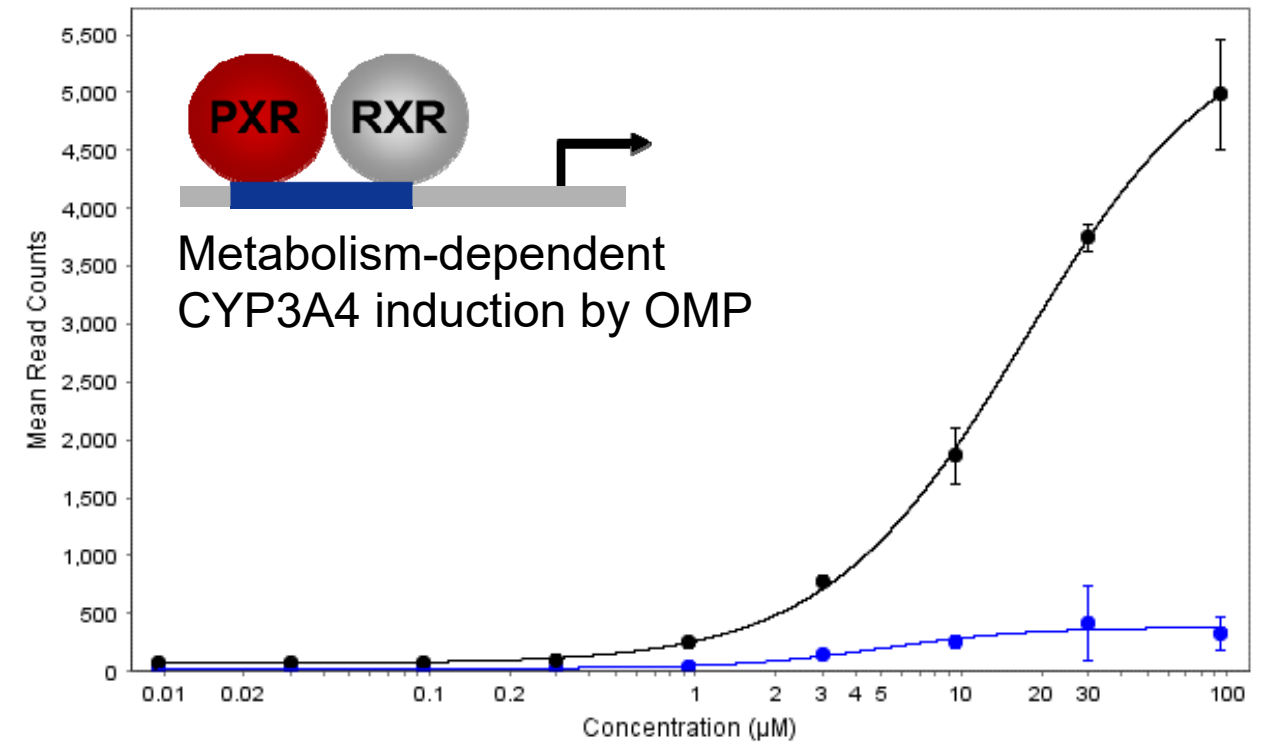
3D HepaRG Spheroids Model Gene- and Pathway-level Transcriptomic Functionality



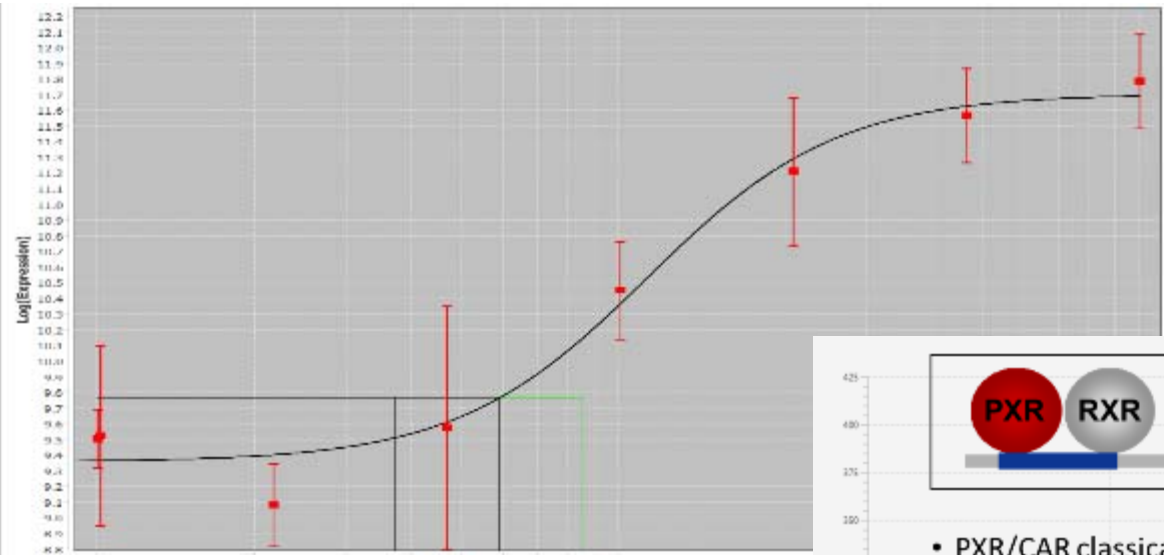


Omeprazole

- Elevated basal CYP1A1 expression in PROLIF HepaRG; linked to liver development
- AhR functionality in 2D & PROLIF
- Reduced xenobiotic metabolism competence impacts CYP3A4 inducibility (PXR)

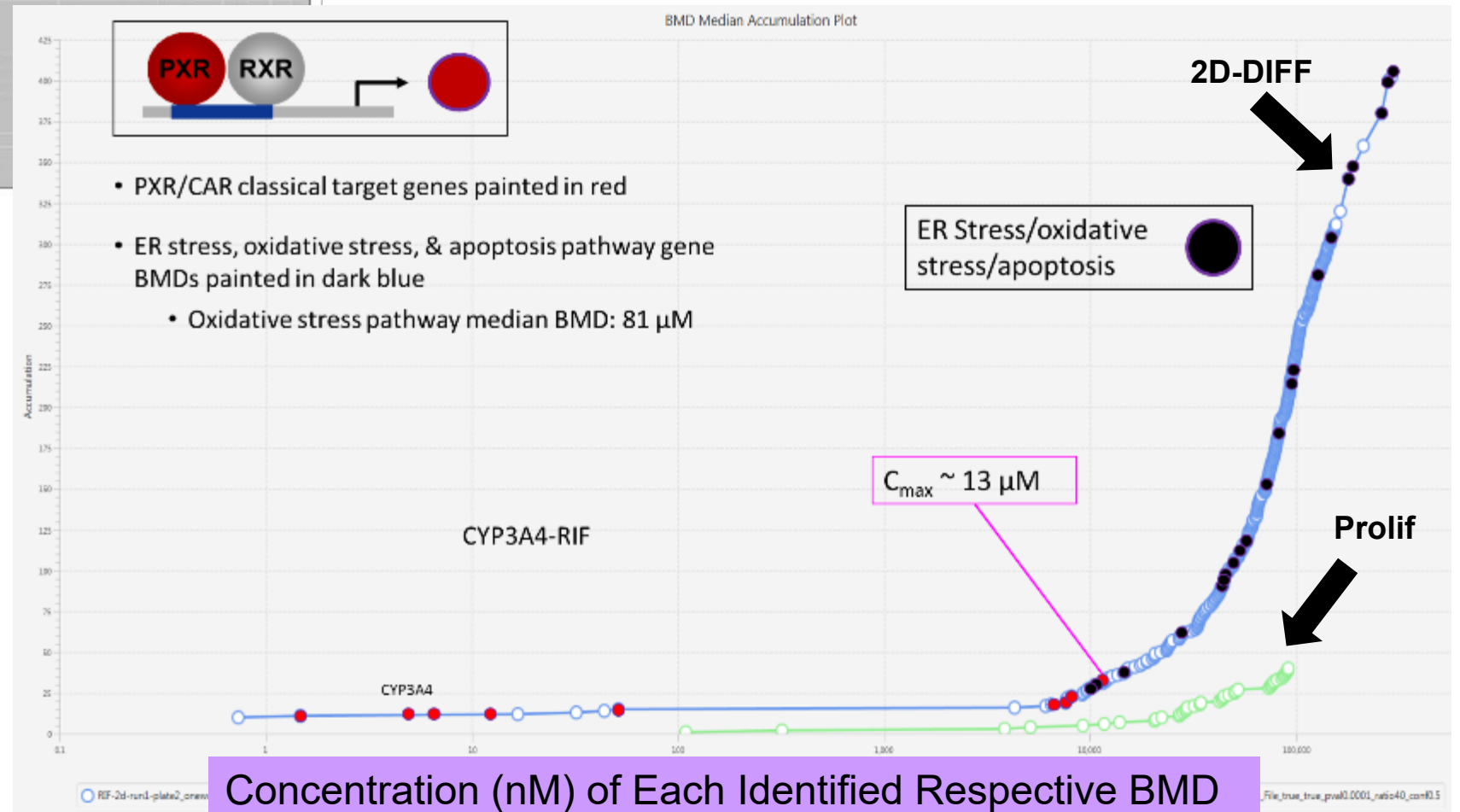


CYP3A4 BMC Curve



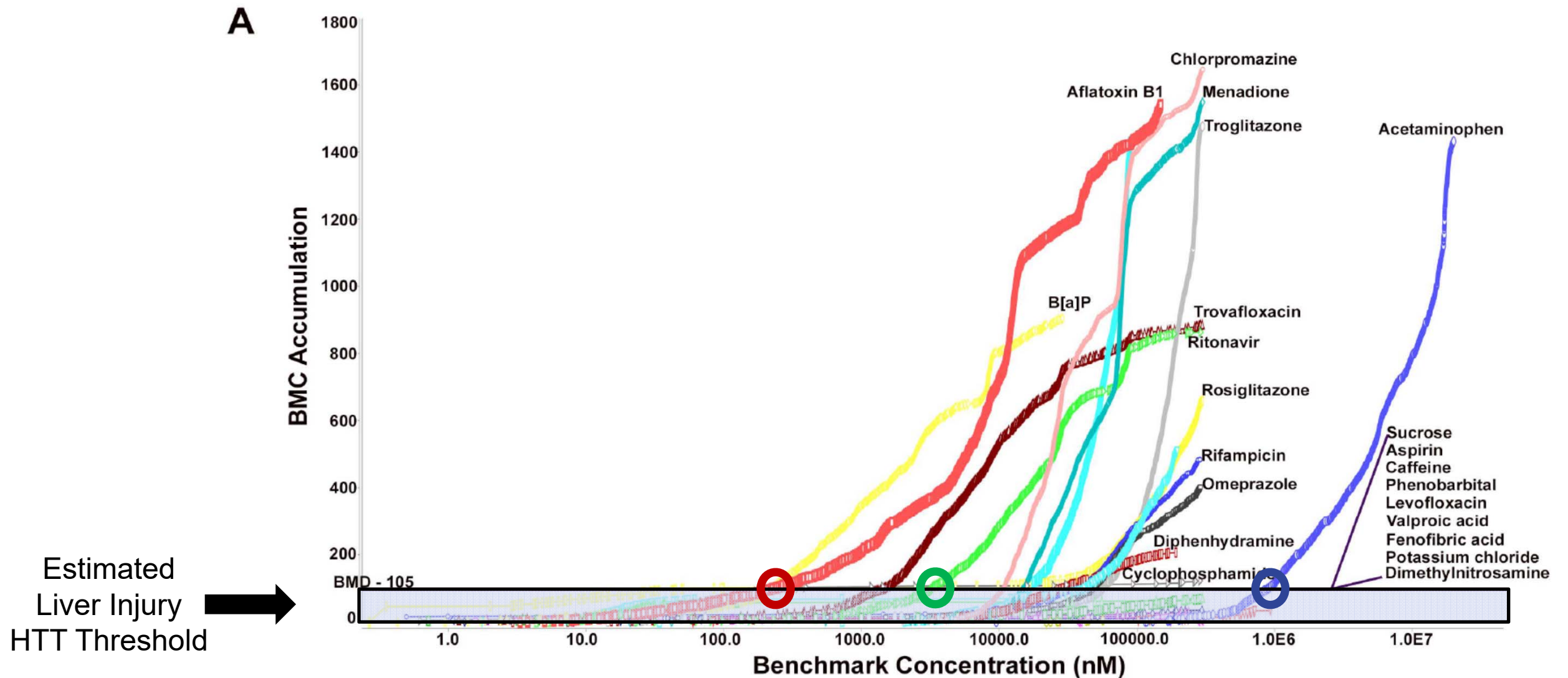
Benchmark Concentration Analysis of Transcriptomic Response

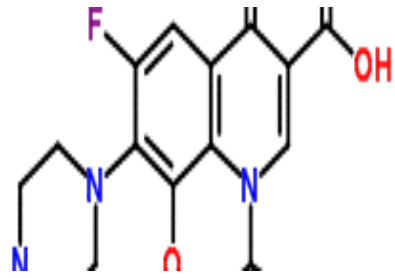
BMC Accumulation Plot (each circle an individual gene BMC)



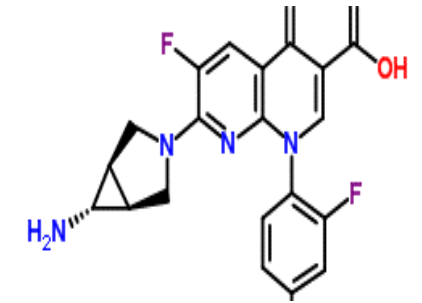
Scott Auerbach & Sciome

Estimation of Liver Injury Potential with Benchmark Concentration Analysis of High Throughput Transcriptomics (S1500+) with 2D Differentiated HepaRG

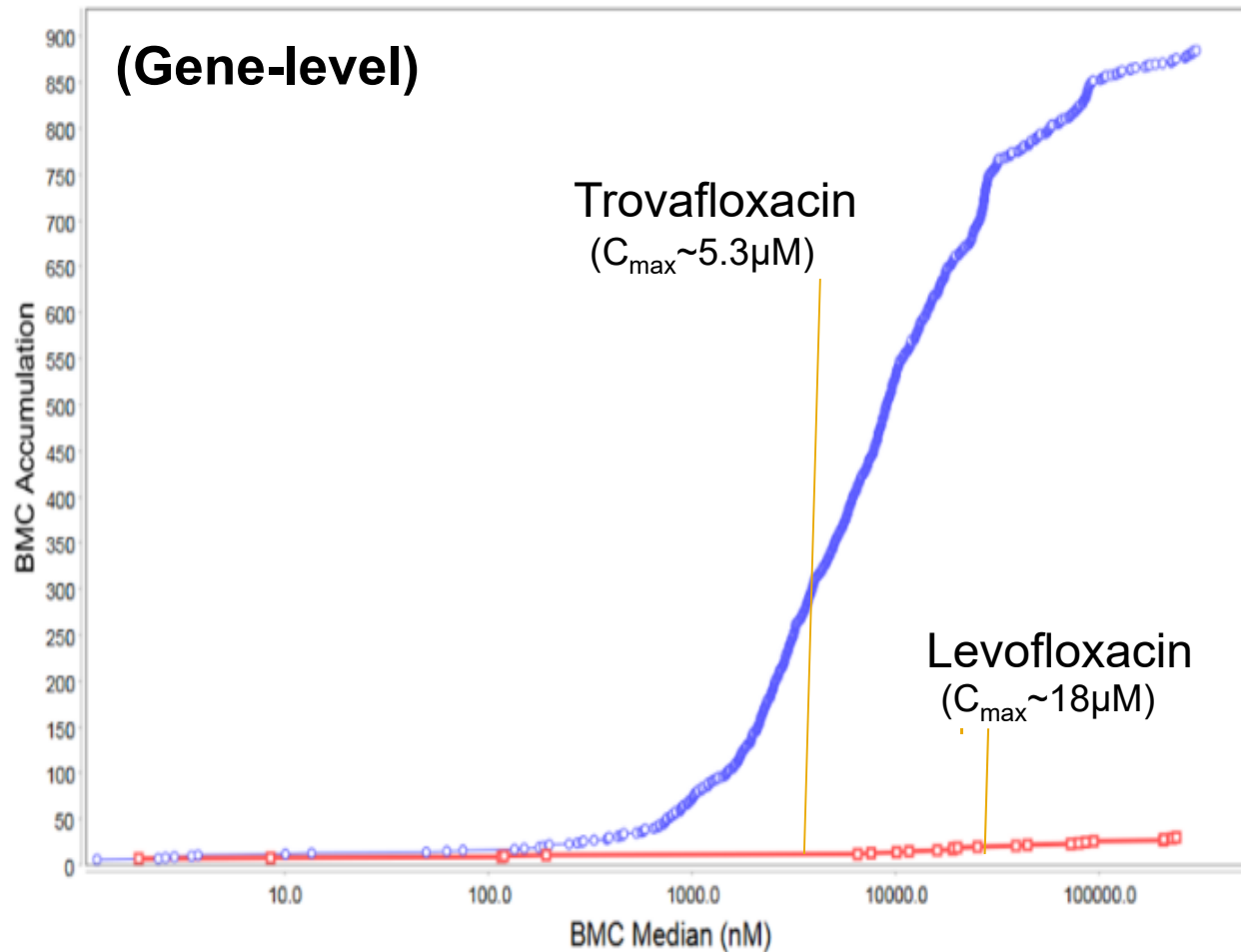




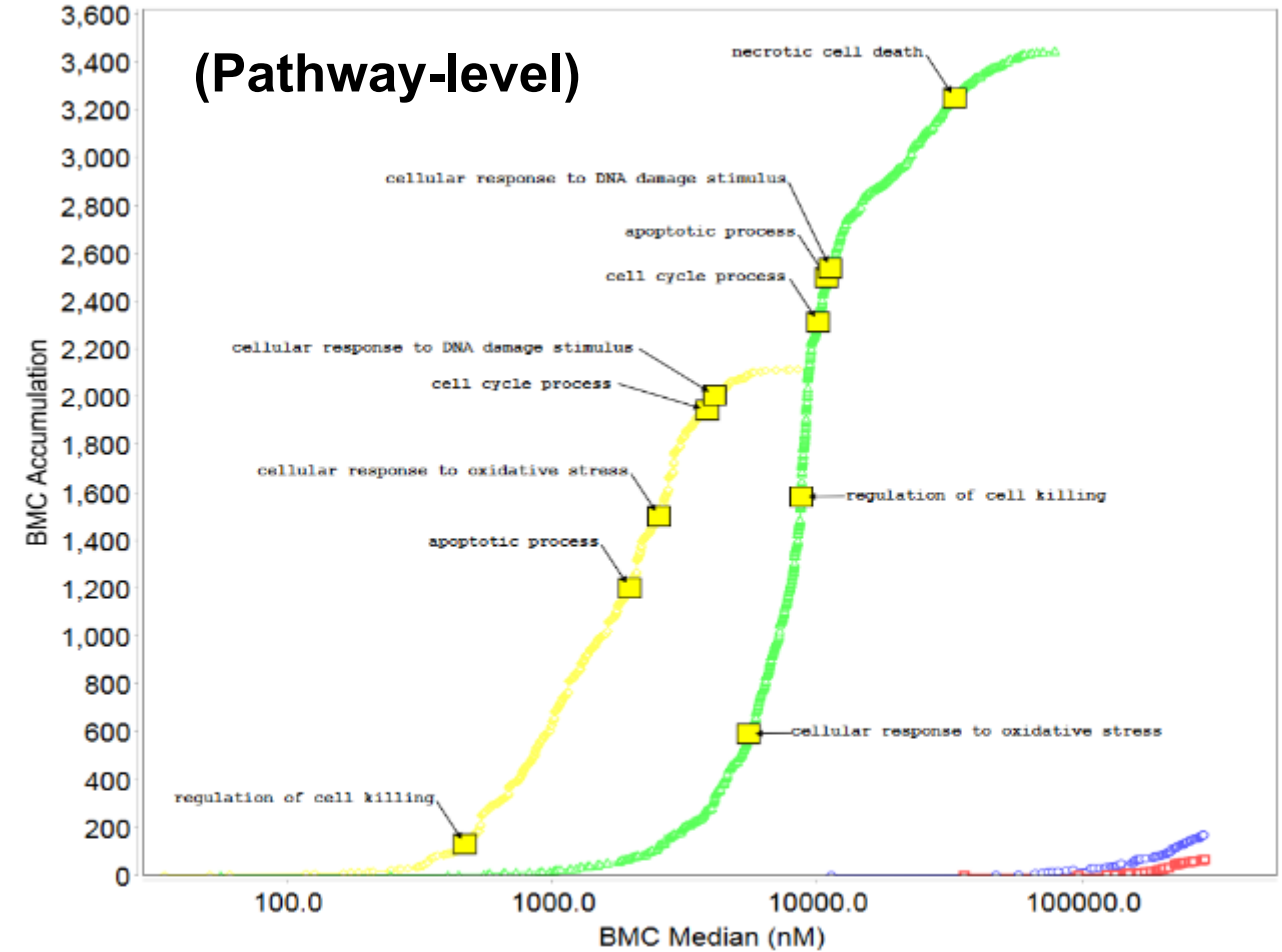
trovafloxacin vs. levofloxacin



BMC Median Accumulation Plot (Gene-level)



BMC Median Accumulation Plot



levofloxacin-2d-run1-plate3_williams_0.05_NOMTC_foldfilter1.5_BMD_S1500_Human_DEFINED-Category File_Human_Individual_Gene_true_true_pval0.0001_ratio40_foldchange2_conf0.5

levofloxacin-prol-run3-plate3_williams_0.05_NOMTC_foldfilter1.5_BMD_S1500_Human_GO_BP_true_true_pval0.0001_ratio40_conf0.5

trovafloxacin-2d-run3-plate3_williams_0.05_NOMTC_foldfilter1.5_BMD_S1500_Human_DEFINED-Category File_Human_Individual_Gene_true_true_pval0.0001_ratio40_foldchange2_conf0.5

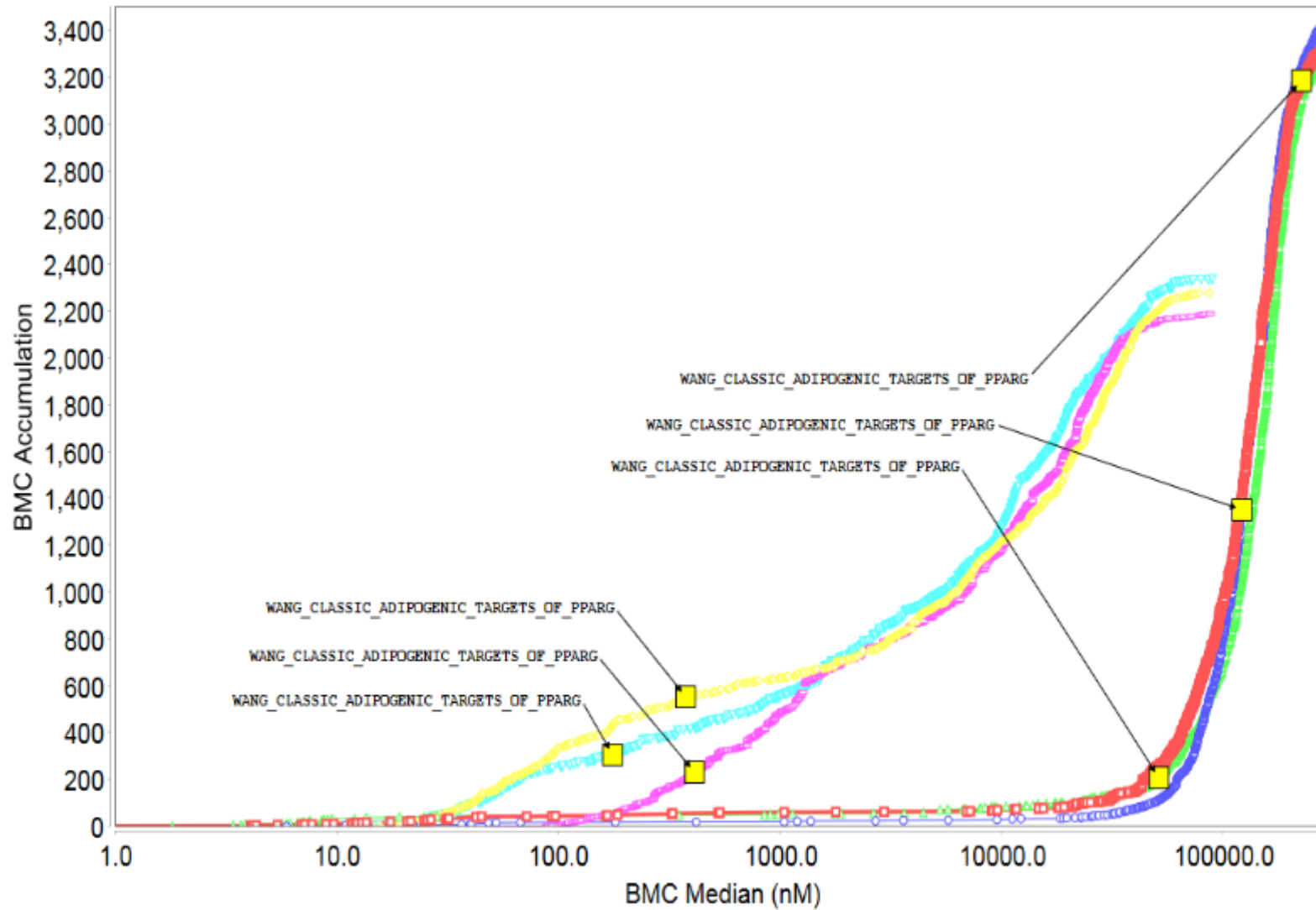
levofloxacin-2d-run3-plate3_williams_0.05_NOMTC_foldfilter1.5_BMD_S1500_Human_GO_BP_true_true_pval0.0001_ratio40_conf0.5

levofloxacin-prol-run3-plate3_williams_0.05_NOMTC_foldfilter1.5_BMD_S1500_Human_GO_BP_true_true_pval0.0001_ratio40_conf0.5

trovafloxacin-2d-run3-plate3_williams_0.05_NOMTC_foldfilter1.5_BMD_S1500_Human_GO_BP_true_true_pval0.0001_ratio40_conf0.5

trovafloxacin-prol-run3-plate3_williams_0.05_NOMTC_foldfilter1.5_BMD_S1500_Human_GO_BP_true_true_pval0.0001_ratio40_conf0.5

BMC Median Accumulation Plot



- rosiglitazone-2d-run1-plate1_williams_0.05_NOMTC_foldfilter1.5_BMD_S1500_Human_DEFINED-HG-U133_Plus_2_C2_msigdb_v5.0_true_true_pval0.0001_ratio40_conf0.5
- rosiglitazone-2d-run2-plate1_williams_0.05_NOMTC_foldfilter1.5_BMD_S1500_Human_DEFINED-HG-U133_Plus_2_C2_msigdb_v5.0_true_true_pval0.0001_ratio40_conf0.5
- rosiglitazone-2d-run3-plate1_williams_0.05_NOMTC_foldfilter1.5_BMD_S1500_Human_DEFINED-HG-U133_Plus_2_C2_msigdb_v5.0_true_true_pval0.0001_ratio40_conf0.5
- troglitazone-2d-run1-plate1_williams_0.05_NOMTC_foldfilter1.5_BMD_S1500_Human_DEFINED-HG-U133_Plus_2_C2_msigdb_v5.0_true_true_pval0.0001_ratio40_conf0.5
- troglitazone-2d-run2-plate1_williams_0.05_NOMTC_foldfilter1.5_BMD_S1500_Human_DEFINED-HG-U133_Plus_2_C2_msigdb_v5.0_true_true_pval0.0001_ratio40_conf0.5
- troglitazone-2d-run3-plate1_williams_0.05_NOMTC_foldfilter1.5_BMD_S1500_Human_DEFINED-HG-U133_Plus_2_C2_msigdb_v5.0_true_true_pval0.0001_ratio40_conf0.5

Ramaiahgari et al., 2019 (Jun) Toxicological Sciences, v.169 (2), 553-566.

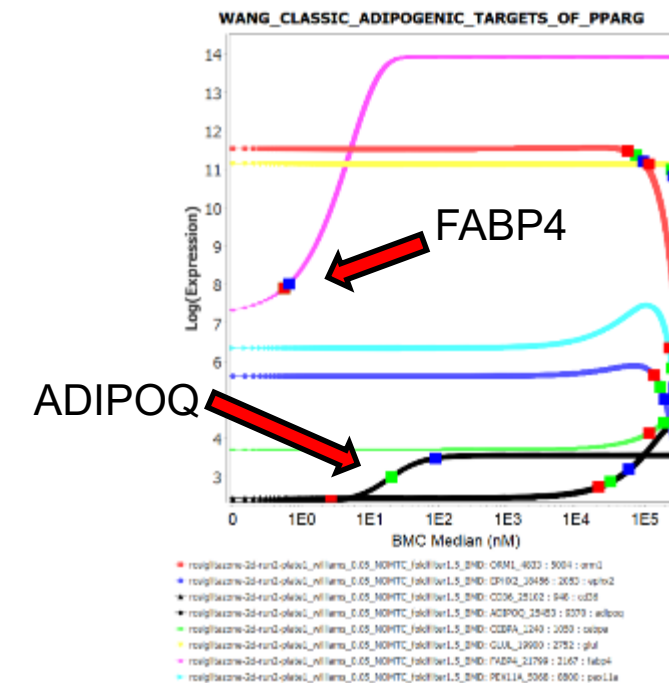
Therapeutic Target ID via BMC Modeling

PPAR γ report gene assay potencies:

Troglitazone: EC₅₀ = ~550 nM

Rosiglitazone: EC₅₀ = ~18 nM

Chen, R. et al. Pharm Biol 55, 503-509 (2017).



NOT all genes within a historical 'pathway' (i.e., gene set) are diagnostic, correct relative potency for FABP4 and ADIPOQ

Cyclophosphamide

(2-fold filter)

□ 3D HepaRG Spheroids

○ 2D-DIFF HepaRG

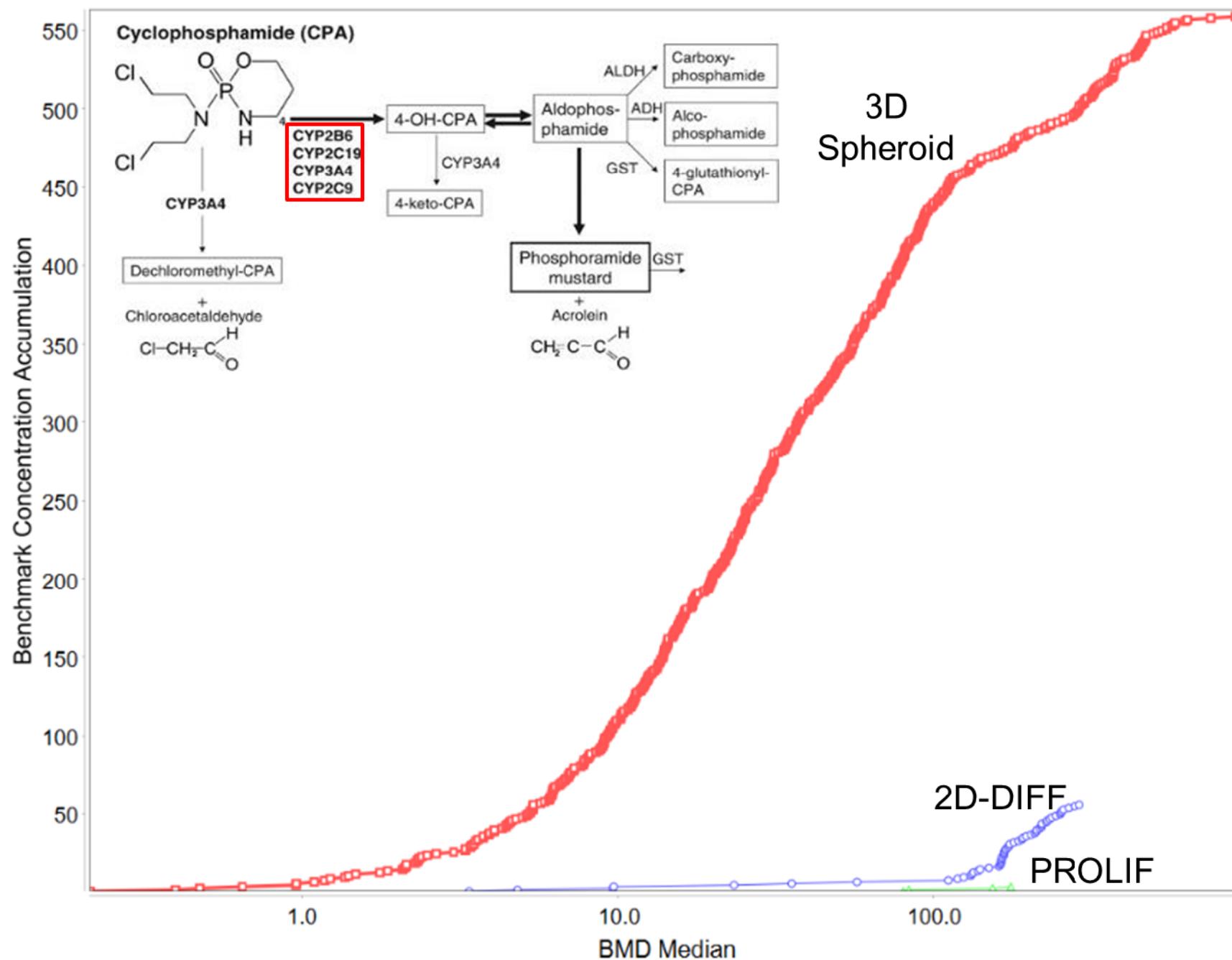
△ PROLIF HepaRG

• Plausibly-relevant response pathways including:

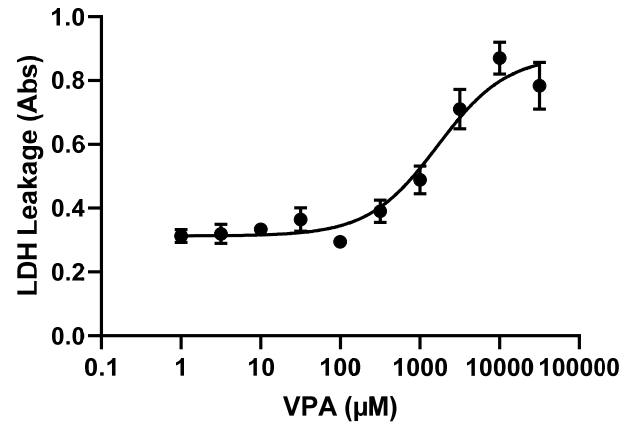
- Lipid hydroxylation
- P450 metabolism
- Cell cycle
- ROS
- DNA damage
- Hypoxia

Unpublished Data

BMC Median Accumulation Plot

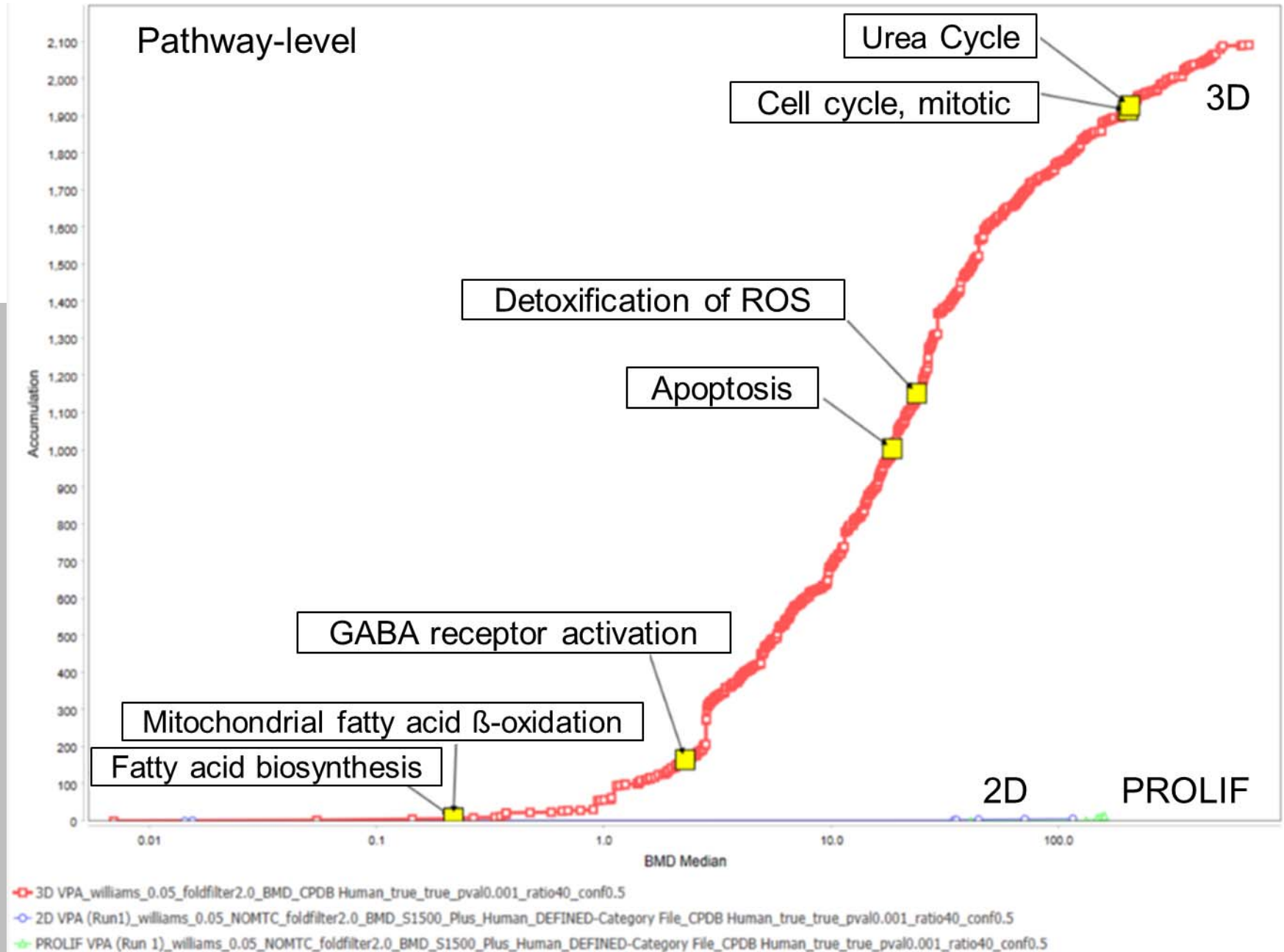


Valproic Acid-3D HepaRG Spheroids

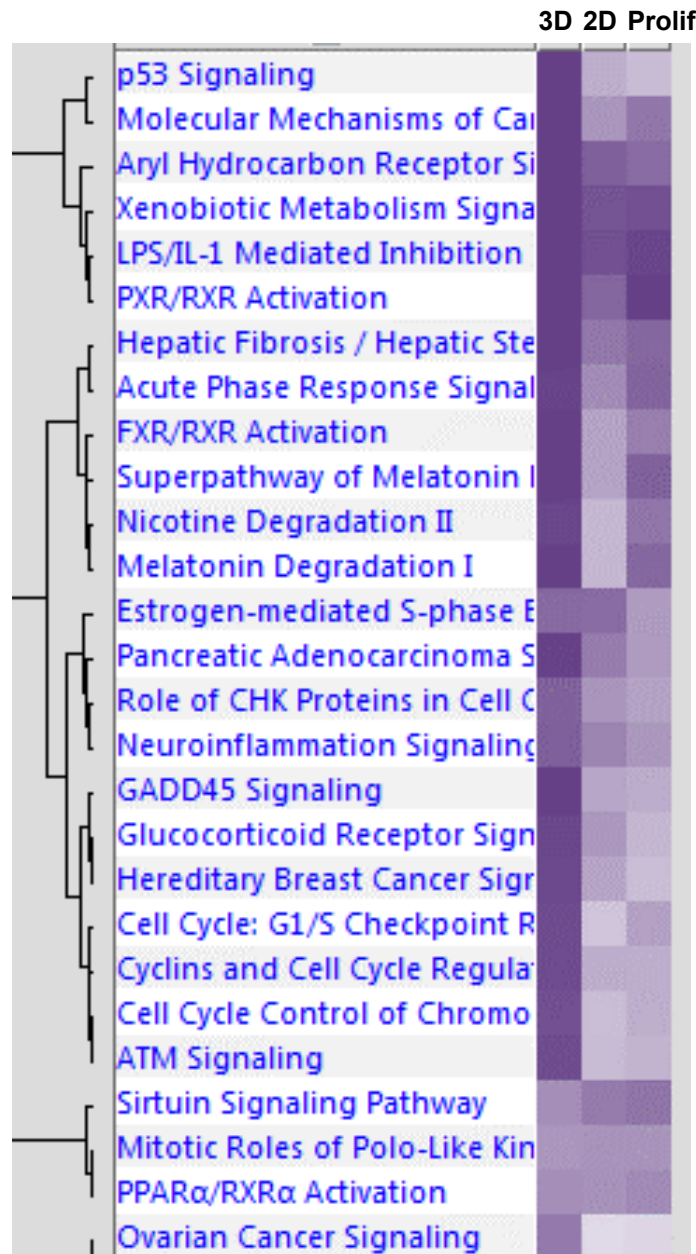


- C_{max} ~240 µM (human plasma)
- Extensively metabolized (P450s)
- Cytotoxicity @ 1000 µM (3D only)
- Alters lipid & fatty acids levels
- Therapeutic target GABAergic receptor
- Hepatic mitochondrial toxicity & hyperammonemia
- Idiosyncratic liver injury compound

Valproic Acid HTT in 3D HepaRG Spheroids



3D Spheroids & Biological Pathway Enrichment



Benzo(a)pyrene (Group 1 carcinogen (IARC))
exposure on HepaRG cell culture models

Canonical Pathways	Significantly changed genes		
	3D_3 μ M	2D_DIFF_3 μ M	2D_PROLIF_3 μ M
P53 Signaling	39	14	14
Molecular Mechanisms of Cancer	71	32	40
AhR Signaling	38	23	23
Xenobiotic Metabolism Signaling	52	35	39
PXR/RXR Activation	27	16	20
Hepatic Fibrosis	38	24	28
Acute Phase Response Signaling	36	22	28
Pancreatic Adenocarcinoma	30	19	18
GADD45 Signaling	12	7	7
ATM Signaling	26	12	14

Dietary Intake of Heterocyclic Amines and Benzo(a)Pyrene: Associations with Pancreatic Cancer

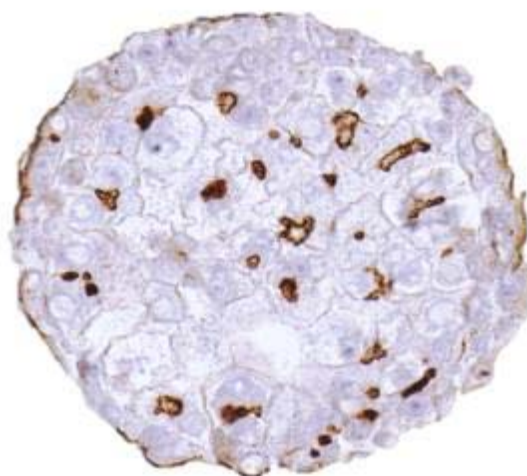
Kristin E. Anderson,¹ Fred F. Kadlubar,² Martin Kulldorff,³ Lisa Harnack,¹ Myron Gross,¹ Nicholas P. Lang,⁴ Cheryl Barber,¹ Nat Rothman,⁵ and Rashmi Sinha⁵

¹School of Public Health, University of Minnesota, Minneapolis, Minnesota; ²National Center for Toxicological Research, Jefferson; ³Department of Ambulatory Care and Prevention, Harvard Medical School and Harvard Pilgrim Health Care, Boston, Massachusetts; ⁴Central Arkansas Veterans Healthcare System and Department of Surgery, College of Medicine, University of Arkansas for Medical Sciences, Little Rock, Arkansas; and ⁵National Cancer Institute, Rockville, Maryland



Opportunities

- Enhanced hepatocyte functionality & population of transcriptomic pathways
- Long-term differentiation for repeated exposure, time-course, & reversibility
- Simple model system readily compatible with most cell culture labs
- Efficient use of hepatocytes
- In vitro pathology to image microtissues for toxicological outcomes (e.g., steatosis, cholestasis, fibrosis)
- Explore potential mechanisms linked with in vitro pathology characteristics



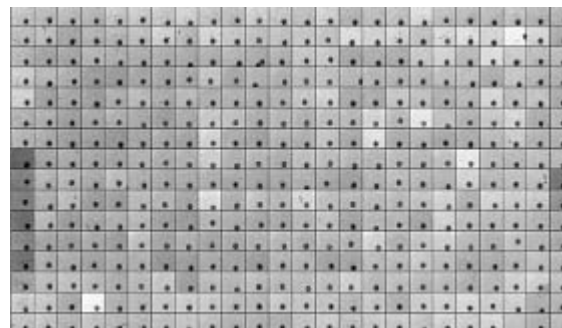
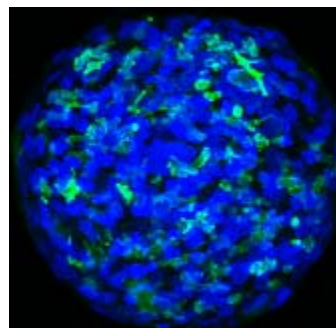
**Biliary Efflux Transporter MRP-2
Immunostaining of HepaRG Spheroids
(21d)**

Challenges

- Changing culture media without liquid handling, aspirating floating spheroids
- High throughput imaging of 3D spheroids
- Recent plate manufacturing issues
- Insufficient knowledge of spheroid maturation & context of use (e.g., phenotypes & outcomes)
- Allometric scaling & biomass challenges (e.g., metabolite profiles over time)
- Inadequate optimization of cell culture media largely adopted from 2D (e.g., DMSO, hydrocortisone)

• Tox21 Cross-partner Project #5

- Develop robust chemical reference dataset for interpretation of high-throughput transcriptomic data: 3D HepaRG spheroids (NTP), MCF-7 (EPA)
- Exposures to >300 pharmaceuticals & chemicals with established high affinity linkages to specific molecular targets (e.g., agonist, antagonist, inhibitor)
- Define signatures of transcriptomic response to reference chemicals to contextualize environmental chemicals

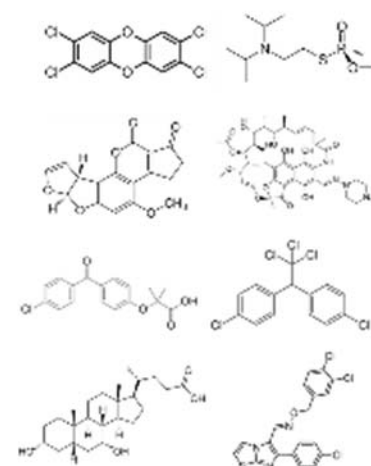


• NTP interspecies parallelogram evaluation of 20 historical chemicals

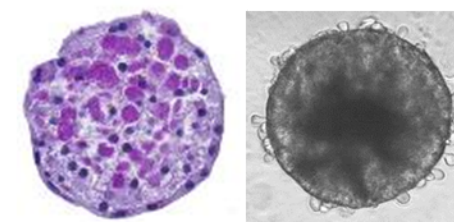
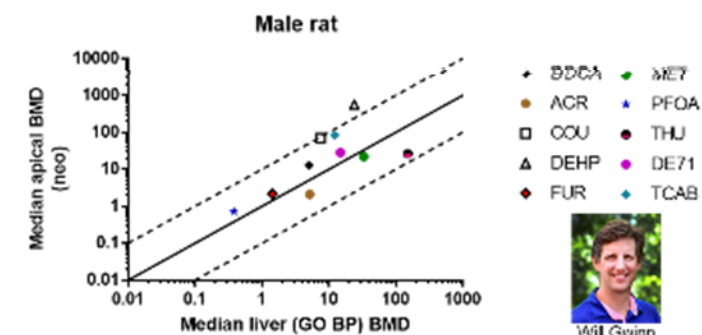
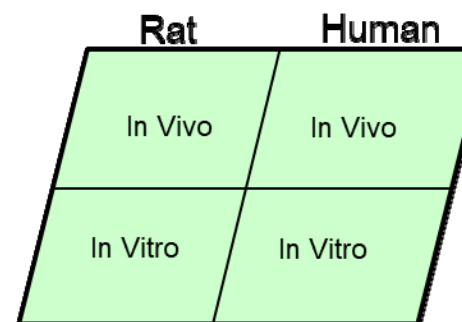
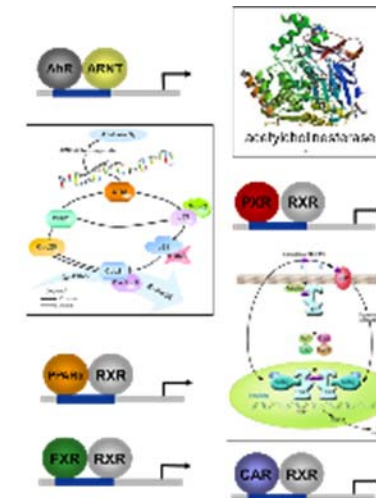
- Chronic in vivo bioassay results (e.g., 2-year carcinogenicity)
- 5-day in vivo rat liver/kidney transcriptomics
- 3D in vitro rat hepatocyte spheroids
- 3D in vitro human HepaRG spheroids

• Histo- and clinical-pathology modeling of liver and renal proximal tubule

Reference Chemicals



Biological-response Pathways



Primary Rat Hepatocytes



Biomolecular Screening Branch



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Rick Paules
Scott Auerbach
Trey Saddler



Alison Harrill
Jui-Hua Hsieh
Fred Parham



Kristine Witt
Stephanie Smith-Roe
Alex Merrick



Stephen Ferguson
Sreenivasa Ramaiahgari
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Jennifer Fostel
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Sciome

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