

Modeling Nuclear Receptor Mediated Pathways in Liver Cancer

Society of Toxicology
March 9, 2010

Virtual Liver Project (v-Liver™)

I Shah, DC Wolf, K Houck, R Judson, J Jack, J Wambaugh,
MT Martin, C Corton

UNITED STATES ENVIRONMENTAL PROTECTION AGENCY



This work was reviewed by EPA and approved for publication but does not necessarily reflect official agency policy.

Predictive Models of Liver Cancer

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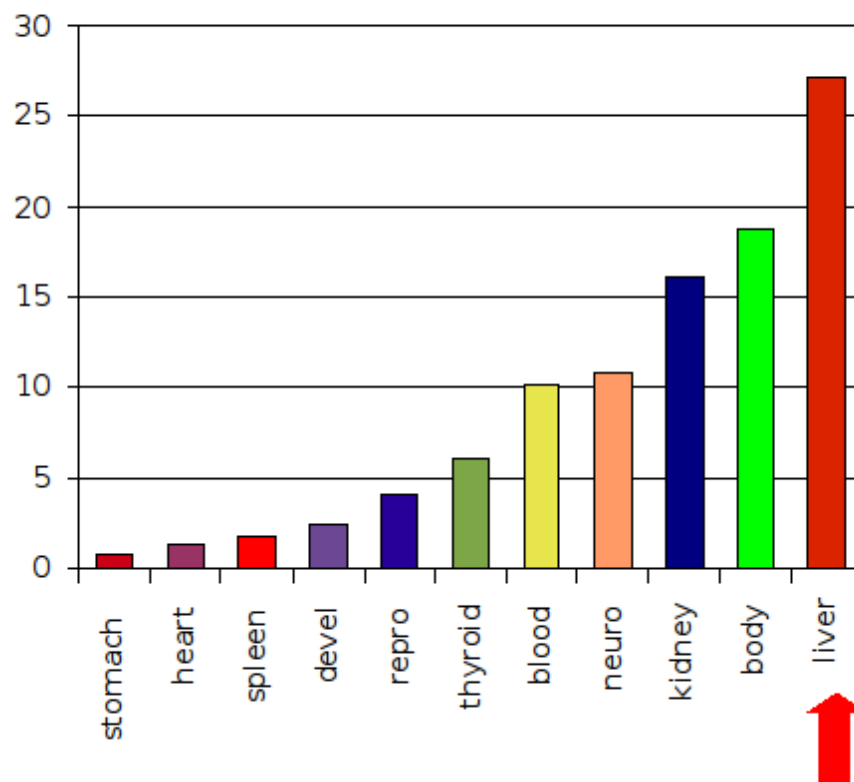
Motivation

- Thousands of environmental chemicals
- Insufficient information on human risk
- Rodent testing infeasible / uncertain results
- Need other decision support tools
- Develop proof-of-concept using
 - Existing *in vivo* data
 - New *in vitro* data
 - Predictive *in silico* systems

Hepatotoxicity

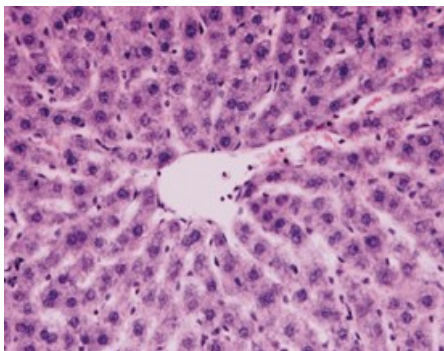
- Liver primary organ for detoxification
- Frequent site of adverse effects in rodents
- Human relevance of some effect uncertain
- Large amount of available data: *-omics*, histopathology, etc.

EPA Integrated Risk Assessment System (IRIS)
(Oral RfDs, Non-cancer endpoints)

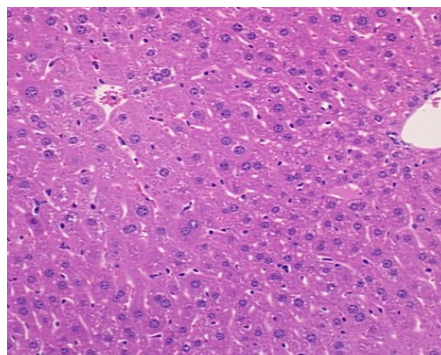


Goal: Predict Tissue Lesions

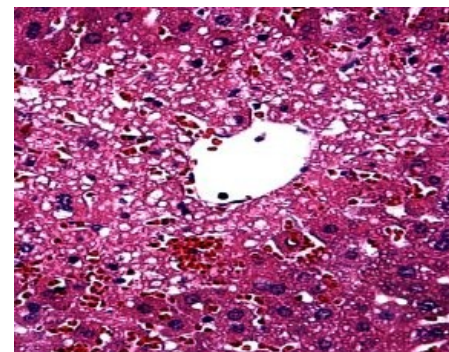
Hydropic Swelling



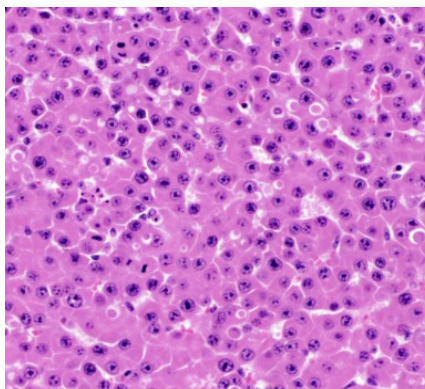
Hypertrophy



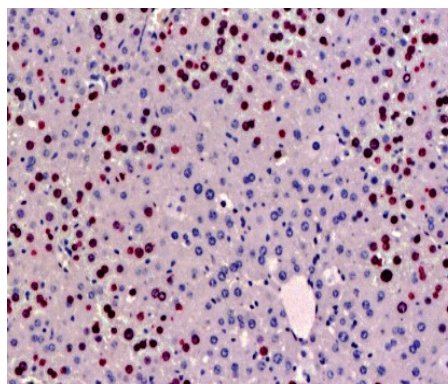
Necrosis



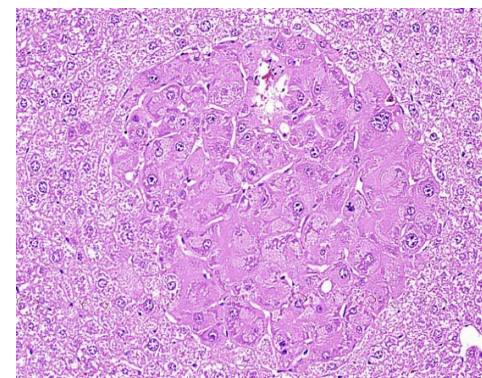
Apoptosis



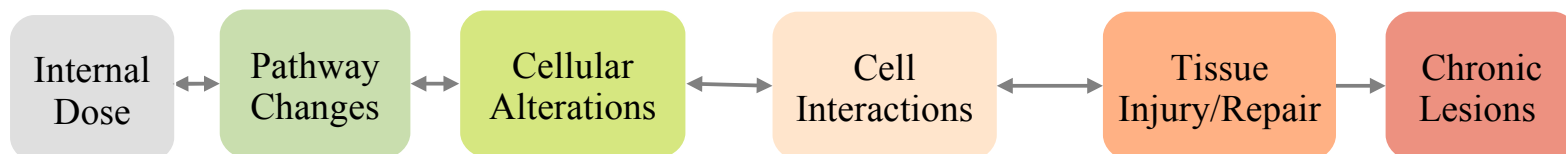
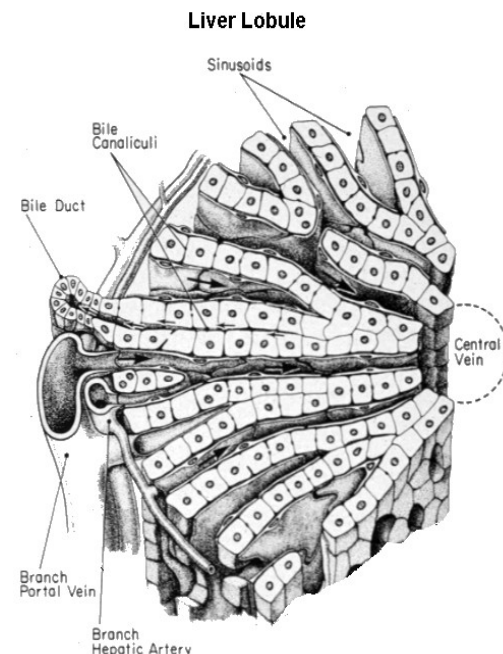
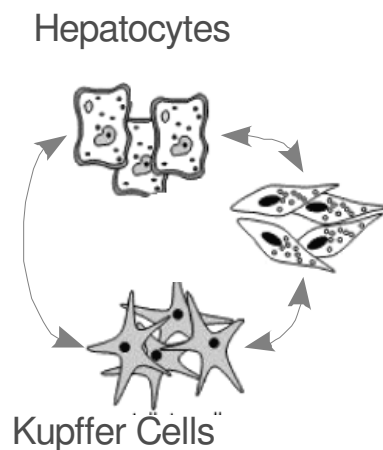
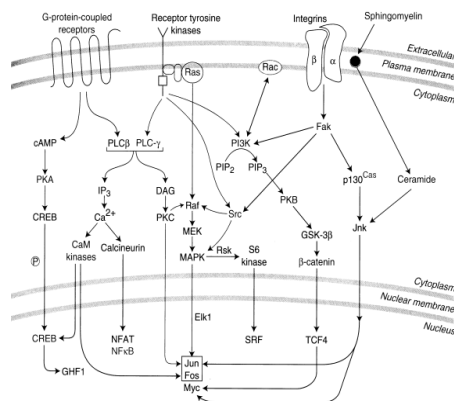
Regenerative Prol.



Altered Hep. Focus



Systems View of Lesion Formation

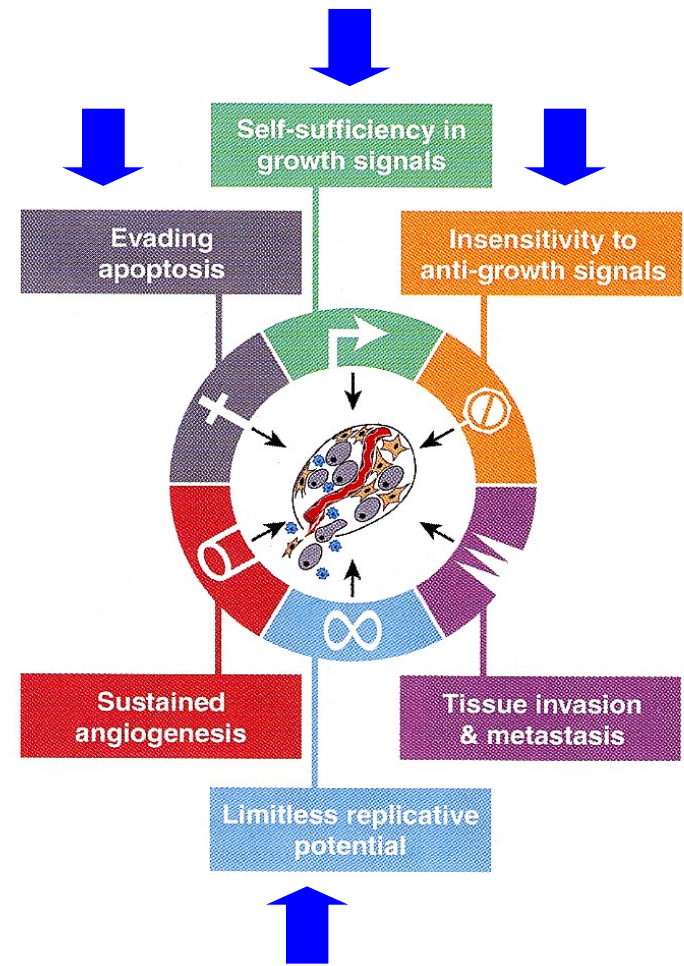


Cancer: Modes of Action

Multiple MOAs: Mutagen,
mitogen, cytotox
Nuclear Receptor (NR)
activation relevant in rodent
hepatocarcinogenesis

Can we extrapolate events to
humans ?

- NR-activation
- Changes in apoptosis/proliferation
- Cancer lesion progression



“Hallmarks of Cancer”
Hanahan & Weinberg, 2000

Virtual Liver Overview

MOAs

Exposure

Internal
Dose

Pathway
Changes

Cell
Phenotypes

Cell
Interactions

Tissue
Injury/Repair

Chronic
Lesions

Data

Environ.
Chemicals

Reverse
TK

ToxCast™

Histomorphometry

ToxRefDB

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v-Liver™

KB

Knowledgebase of normal pathways and chemical effects

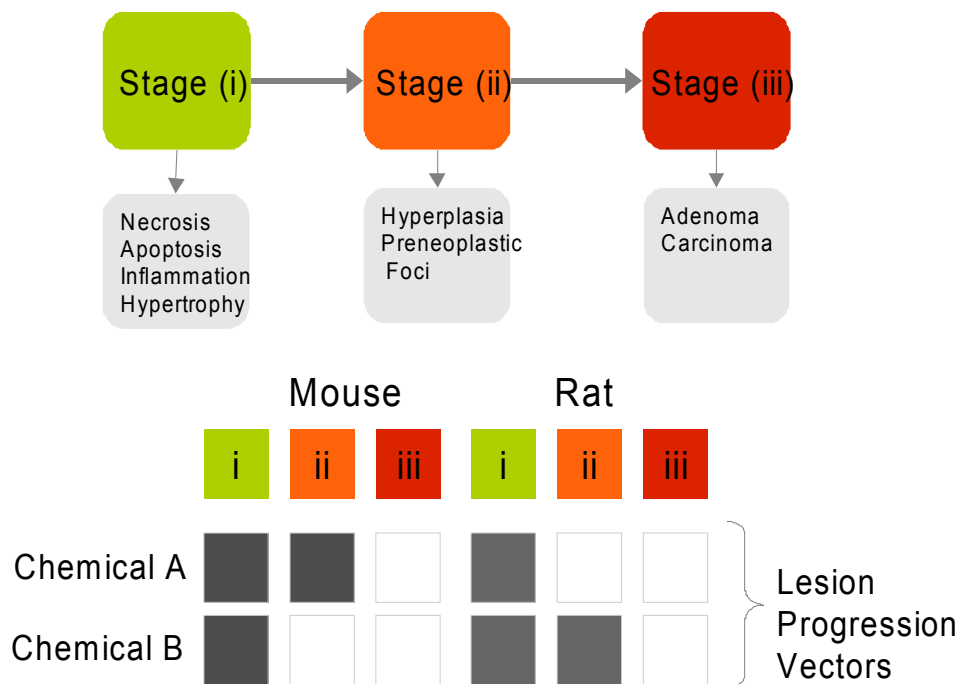
Simulator

Microdosimetry
Simulation

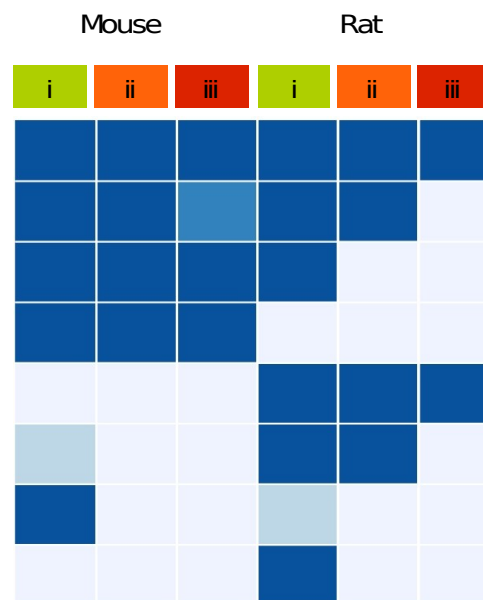
Cellular Response
Simulation

Tissue Response
Simulation

Selecting Chemicals Using Rodent Histopathology

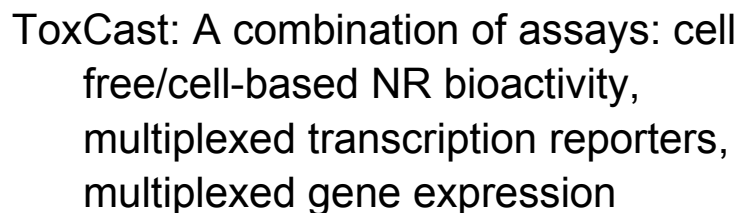


ToxRefDB

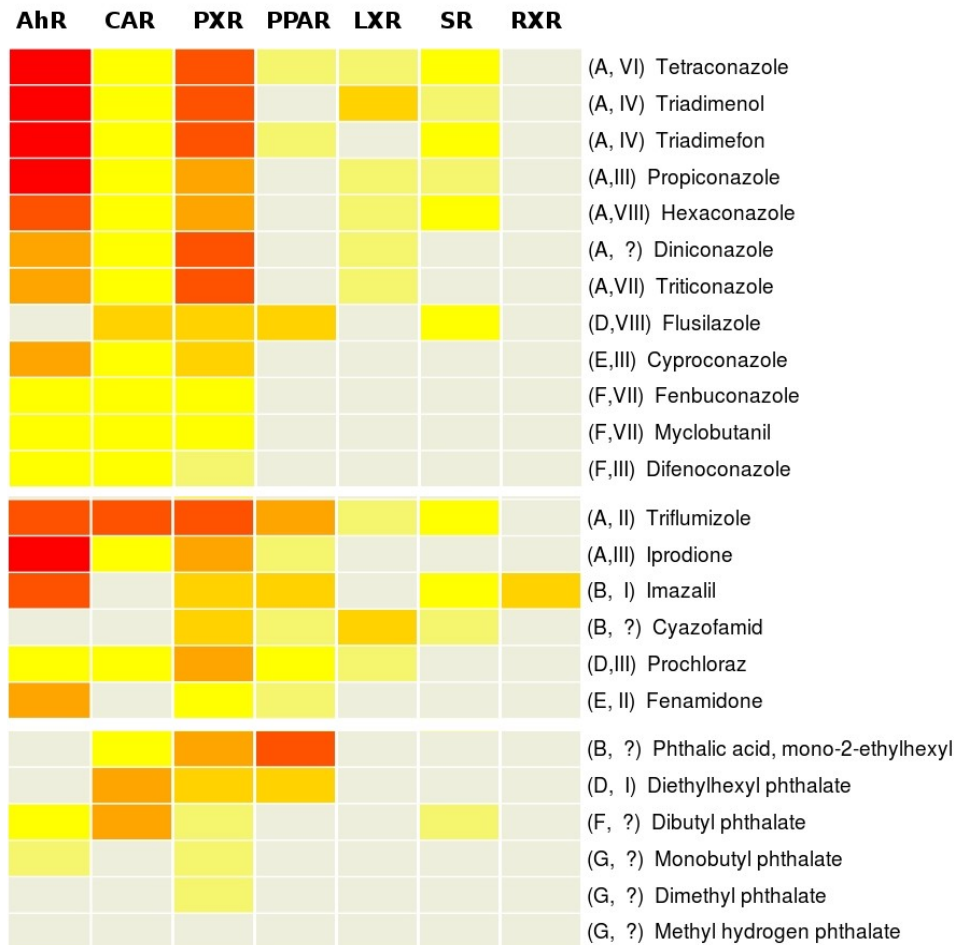


Shah, *et al.* (submitted)

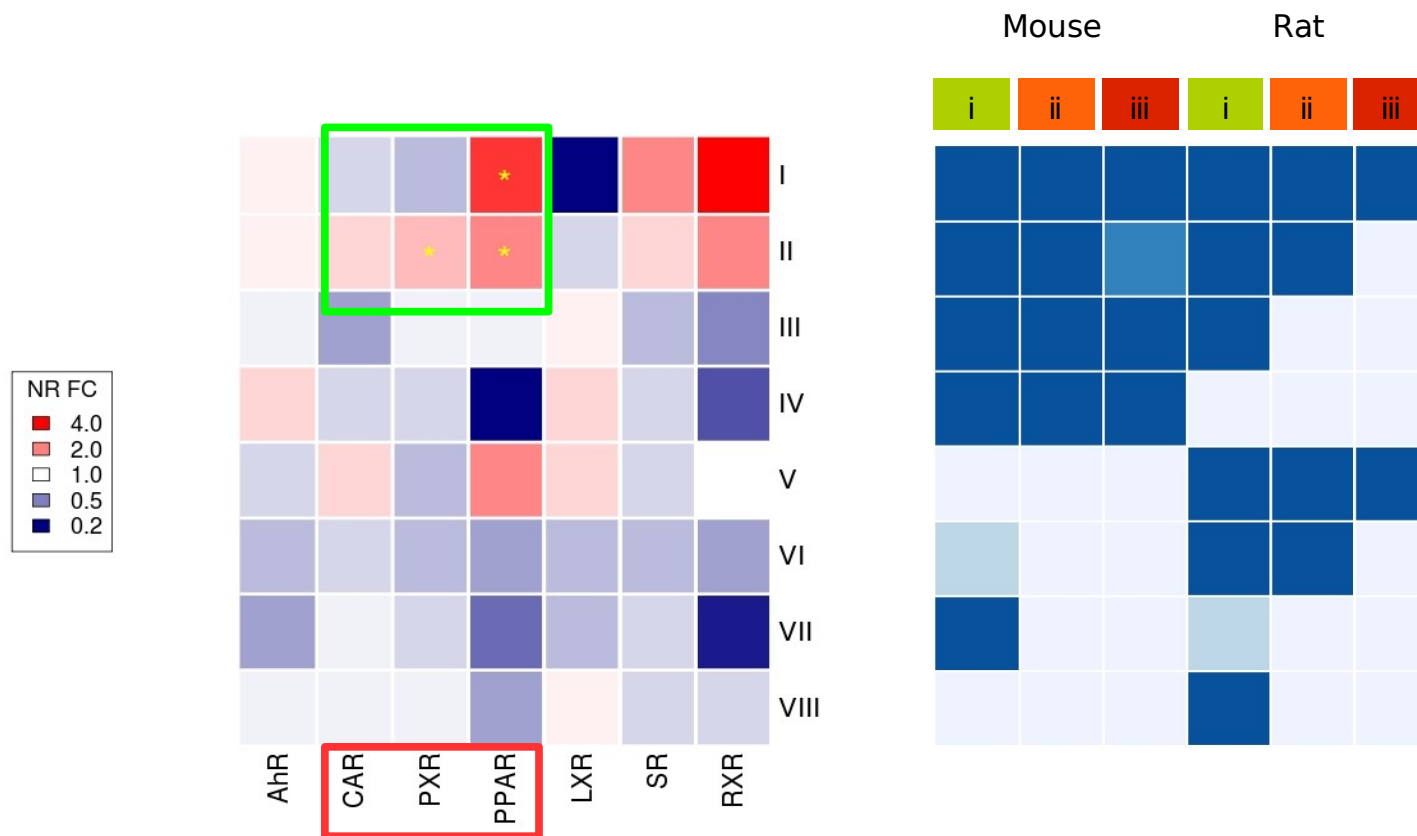
<http://www.epa.gov/ncct/toxrefdb>

Shah, *et al.* (submitted)

Aggregate NR Activities: Examples

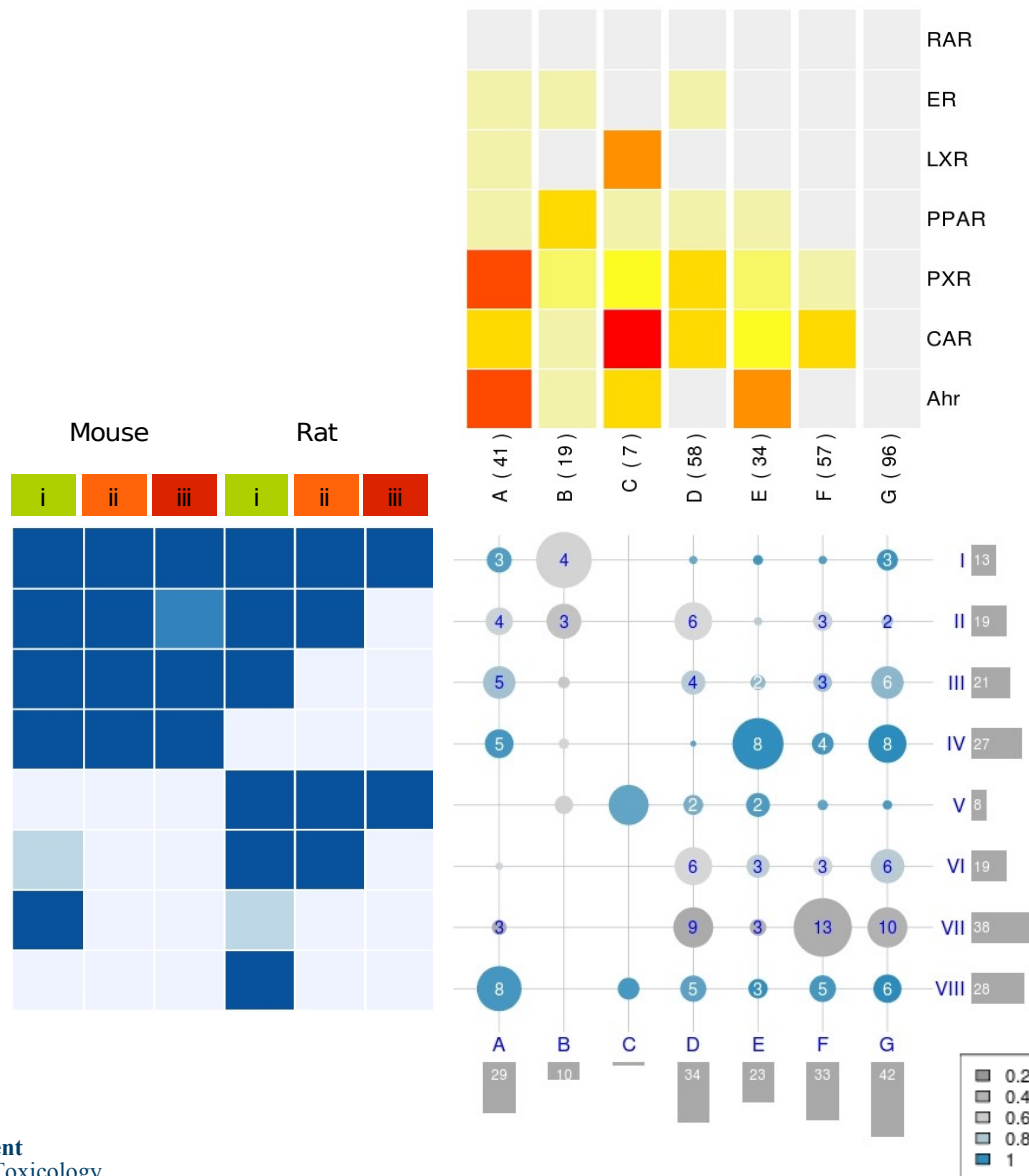


Human NRs and Rodent Cancer



Shah, *et al.* (submitted)

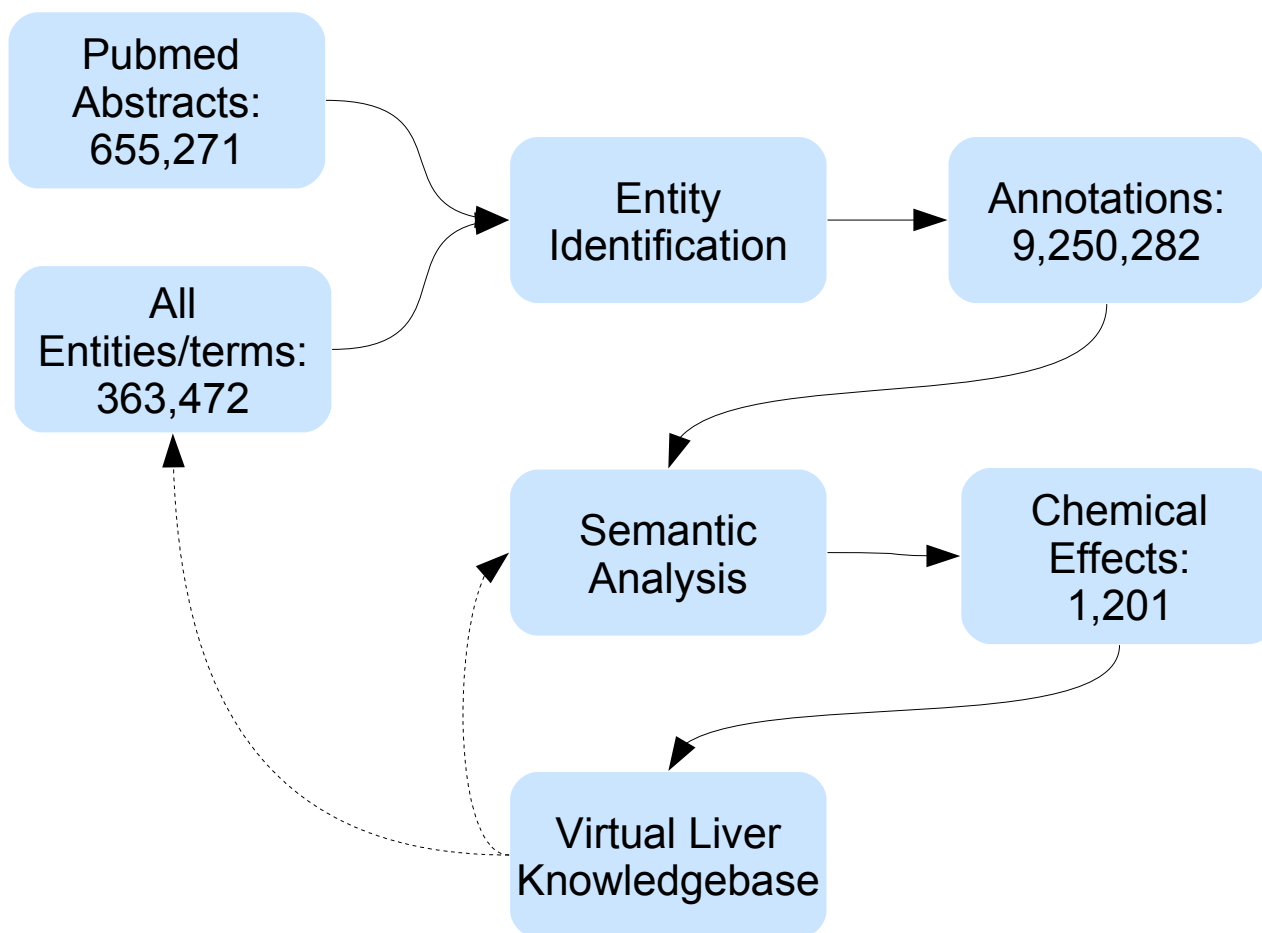
NR activity vs. Rodent Cancer



Relating NR Activity and Pathways

- Need to mine information from literature and public domain data
- Use natural language processing (NLP) tools
- Semi-automated curation of chemical-induced effects
- Store all information in v-Liver Knowledgebase

Literature Curation Overview



Shah & Haugh

Virtual Liver Knowledgebase

vLiverKB

KB Query

KB Search KB Browse

Class/Entity

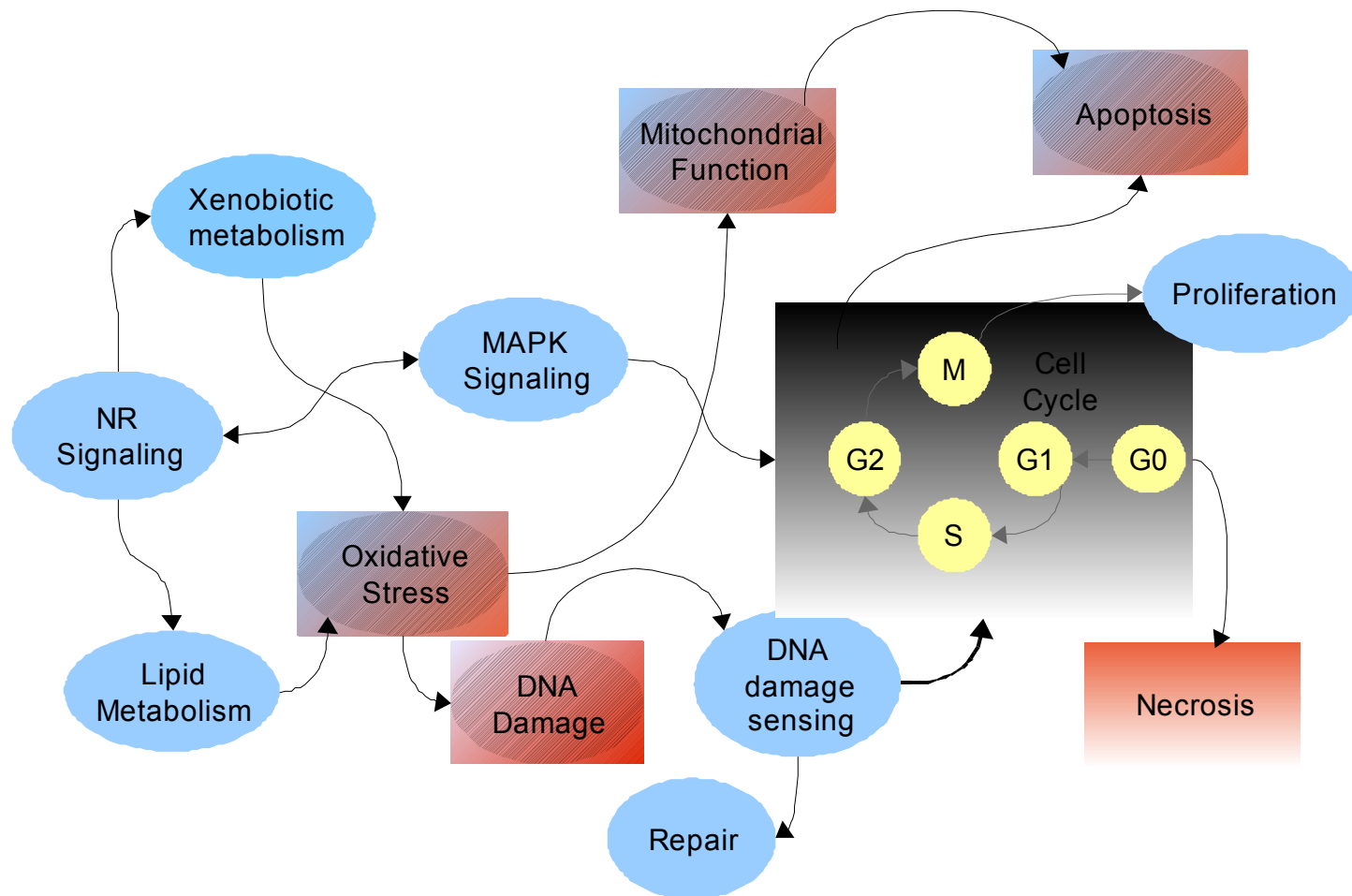
- bio:anatomic-location
- bio:cell
- bio:cell-event
- bio:cell-loc
- bio:chemical
- bio:complex
- bio:effect**
- bio:gene
- bio:mesh-cell
- bio:molecular-event
- bio:organism
- bio:phenotype
- bio:process
- bio:protein
- bio:protein-domain
- bio:ma
- bio:tissue-event

Entities Models VizOpts

Property	Value
comment	Peroxisome proliferator-activated receptor-gamma
has-change	regulating
uri_s	vl:eff:1006
has-name	effect:1006
has-subject	vl:330039
uri_l	http://www.epa.gov/ncct/kb/vlkb#eff:1006
has-reference	pm:15150131
type	bio:effect
has-object	vl:177155

Layout spring + - Delete Print

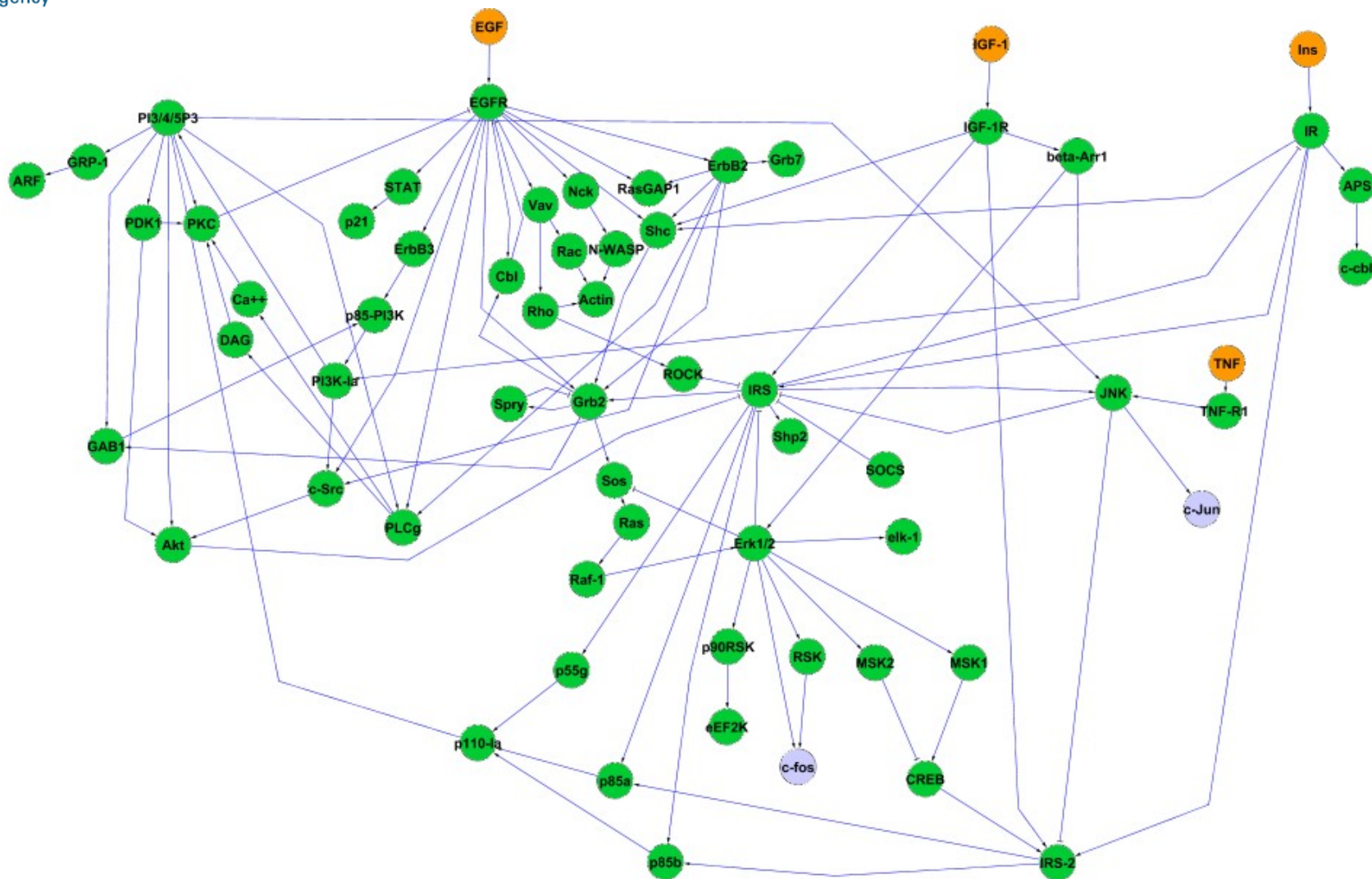
Knowledgebase: Cellular Processes



NR-Mediated Crosstalk Pathways

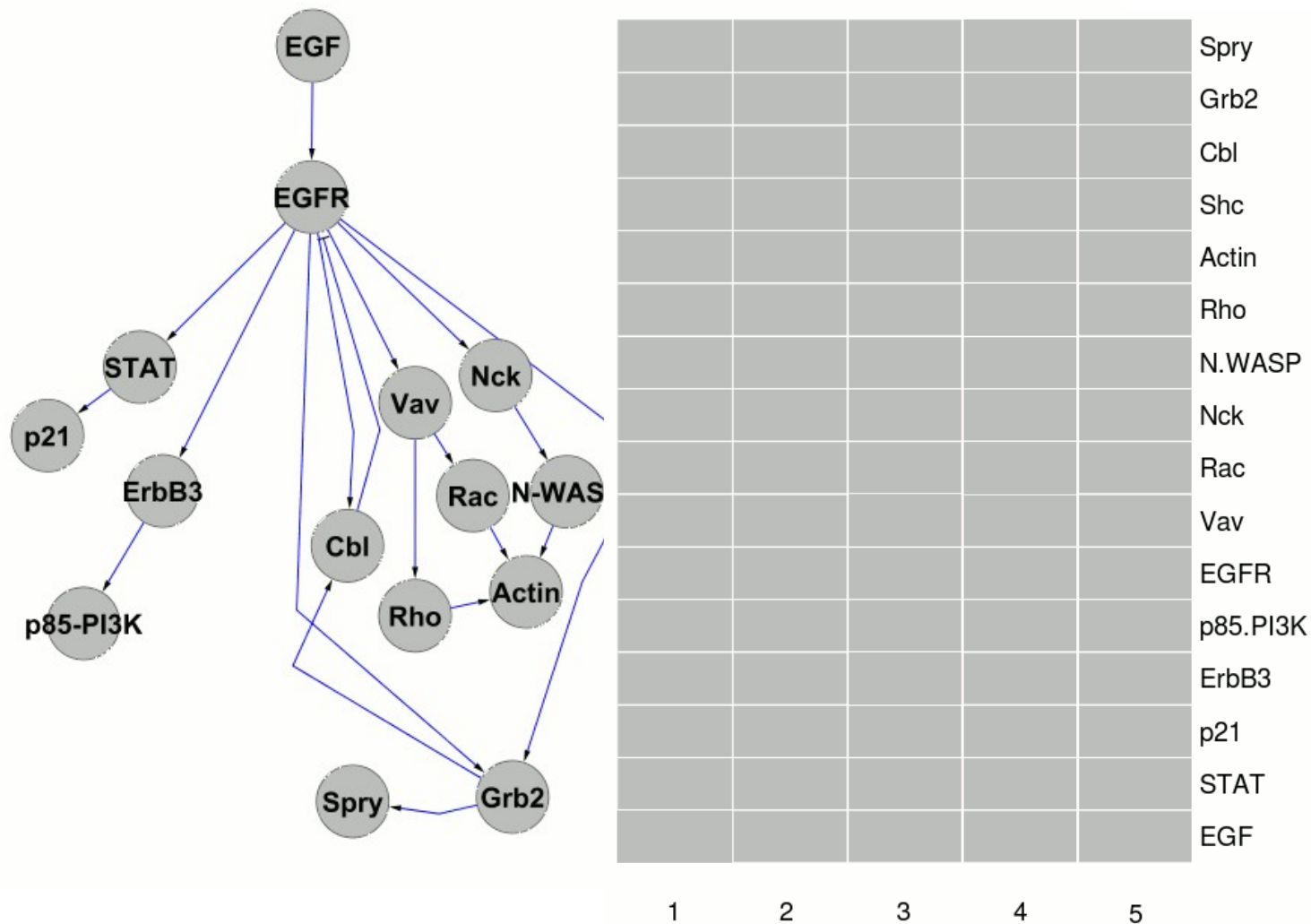
- Use KB to develop systems model of NR-mediated pathways
- E.g. synthesize evidence growth factor (GF) induced cell cycle changes (G1/S-transition)
- Evaluate potential crosstalk between NR and GF-signaling

Growth Factor Crosstalk: Cell Cycle



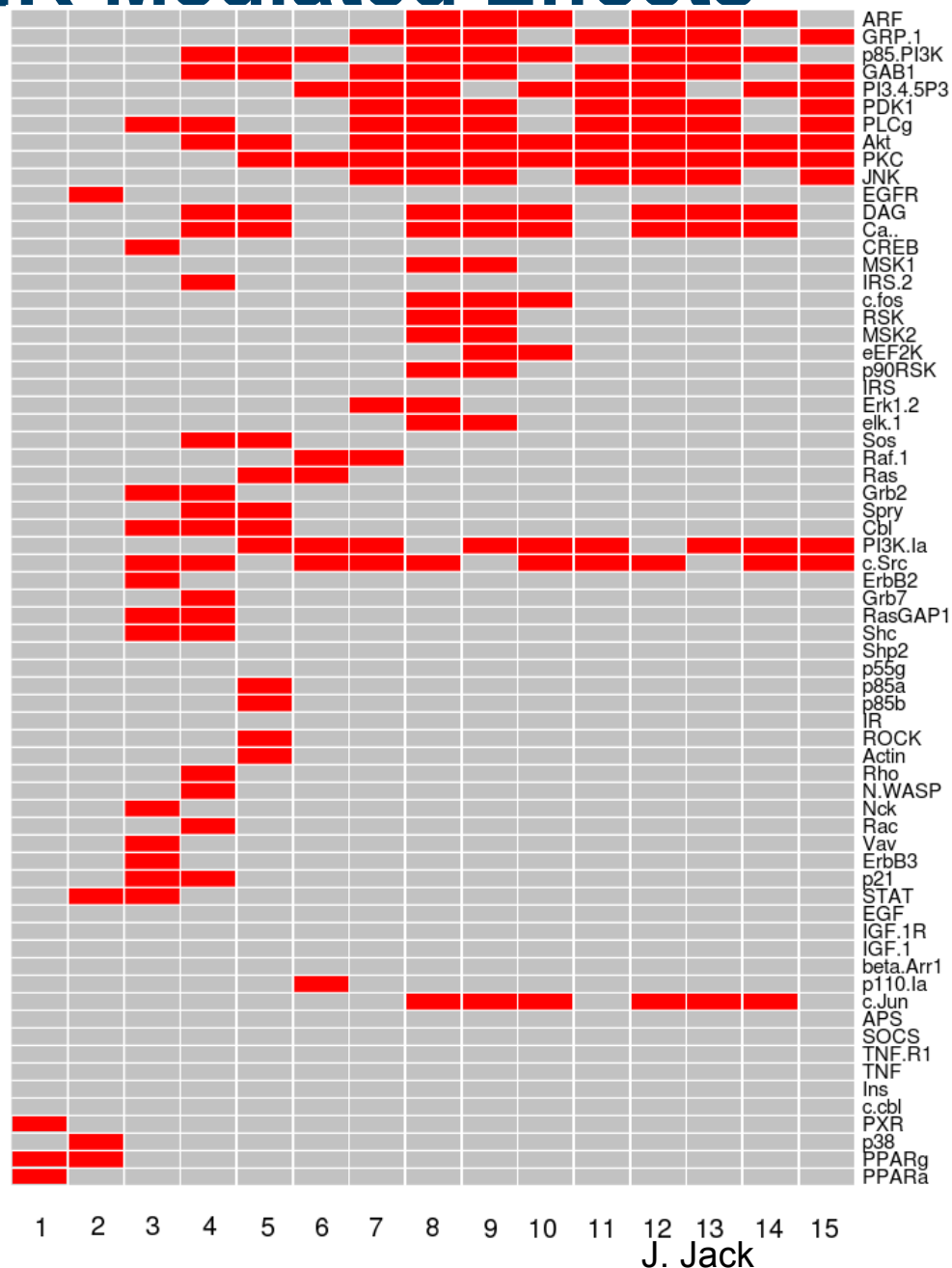


Simulating Signal Propagation



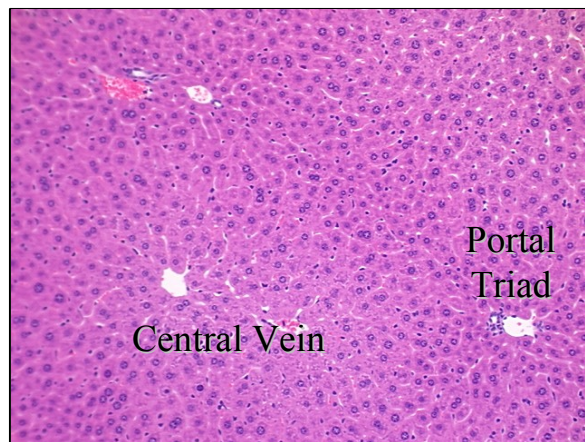
Simulate NR-Mediated Effects

- Transient Activation of PPAR α , PPAR γ , and PXR
- Effects:
 - cFos activation (t=8)
 - cJun activation (t=8)
 - Erk1/2 activity (t=7)
 - p21 activation (t=3)
 - Akt activation (t=7)

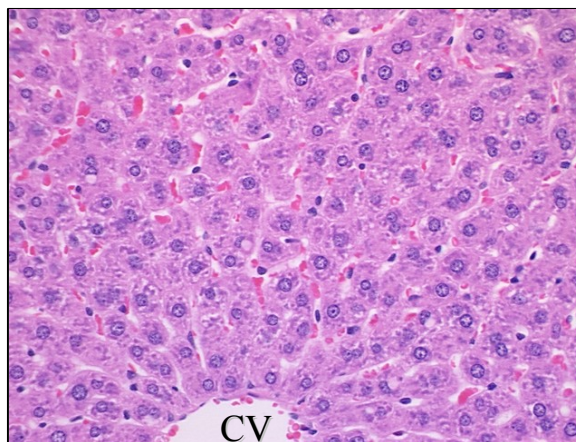


Quantitative Tissue Modeling

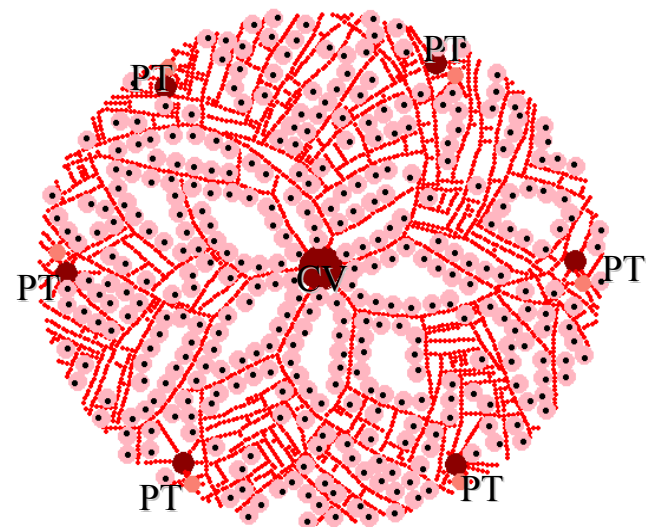
Dose-Response for a Virtual Lobule



The classic lobule consists of a single central vein fed venous and arterial blood via multiple portal triads



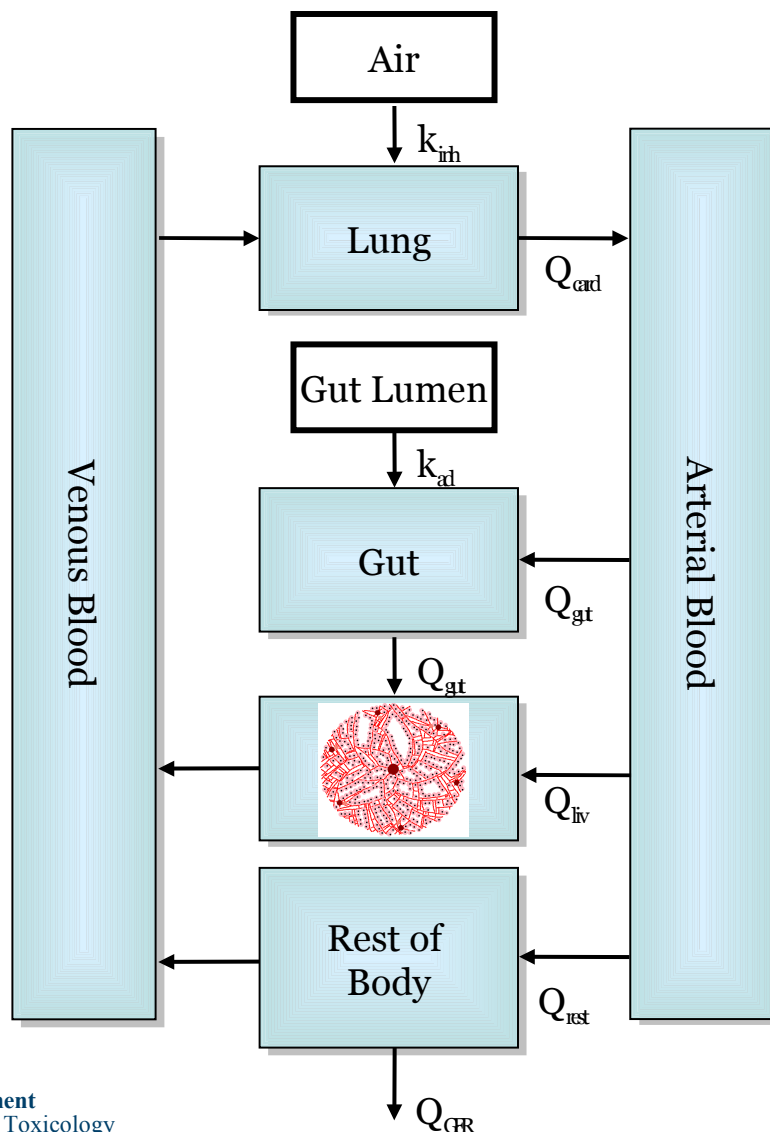
Blood flows through a network of sinusoids, supplying and exposing hepatocytes



Synthetic lobule sufficiently complex to determine that approach would work with actual lobule morphology (e.g. Drasdo et al.)

Wambaugh & Shah (submitted)

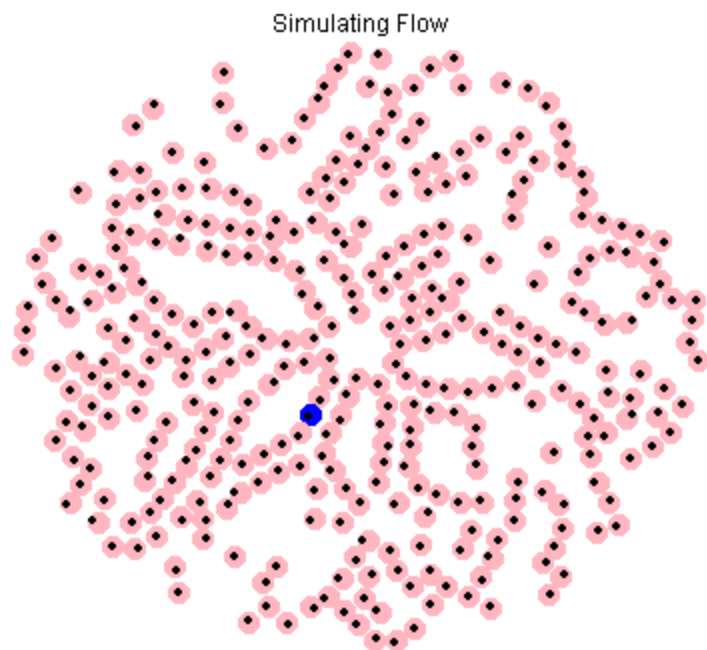
Integration with Pharmacokinetic Models



- Our lobule module is directly integrated with a pharmacokinetic model.
- Determines specific amount of chemical delivered to any given hepatocyte in μmol .
- Suitable for endogenous and xenobiotic compounds.

Simulating Microdosimetry and Effects

Slow Metabolism
($Cl_{it} = 1 \mu\text{L}/\text{min}/10^6 \text{ hep.}$)



- H^o (Normal/Quiescent)
- H^{adt} (Stressed/Adaptive)
- H^{inj} (Stressed/Injured)
- H^{nc} (Necrotic)
- H^{pd} (Proliferative)
- H^{ap} (Apoptotic)

Concentration

Above Threshold

Below Cytotoxicity Threshold

Wambaugh & Shah (submitted)

Summary

- Goal: NR-mediated cellular pathways, alterations and lesions
- Integrative computational and experimental paradigm is vital for success
- Next steps: *short-term in vivo* study design and assays evaluating *in vitro* data, MOA/dose-response

Multi-disciplinary Team: Cross-EPA/ORD & External Collaborations

v-Liver

John Jack
John Wambaugh
Chris Haugh
Woody Setzer
Lockheed Martin

v-Embryo

Thomas Knudsen
Amar Singh
Nisha Sipes
Nicole Kleunstreuer
Michael Roundtree

ToxCast

David Dix
Keith Houck
Richard Judson
Matt Martin
Holly Mortensen
David Reif
Ann Richard

ExpoCast

Elaine Hubal-Cohen

Robert Kavlock

