

# v-Liver™ The Virtual Liver Project: Simulating Hepatic Lesions

*v-Tissues 2009, April 21-22, 2009*

UNITED STATES ENVIRONMENTAL PROTECTION AGENCY



**COMPUTATIONAL  
TOXICOLOGY**

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

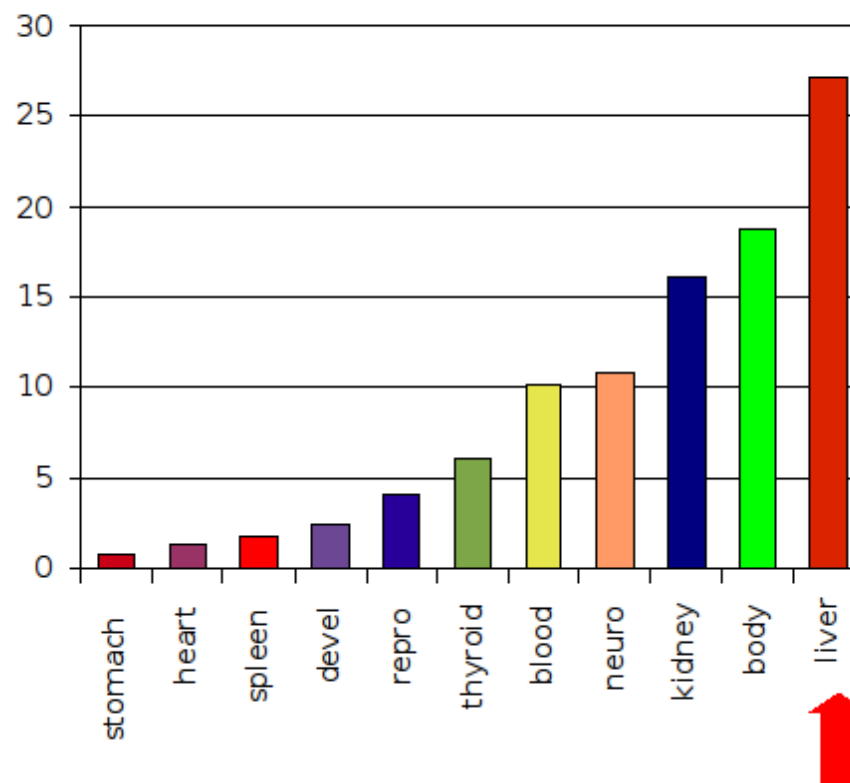
# Background

- ~10,000 HPV chemicals - little/no biological data !
  - Animal testing infeasible / uncertain value
  - 2007 NRC - Toxicity Testing 21<sup>st</sup> Century
  - 2008 EPA - Strategic Plan for Evaluating Chemical Toxicity
- Develop *in vitro-in silico* proof of concept
- Evaluate using environmental chemicals

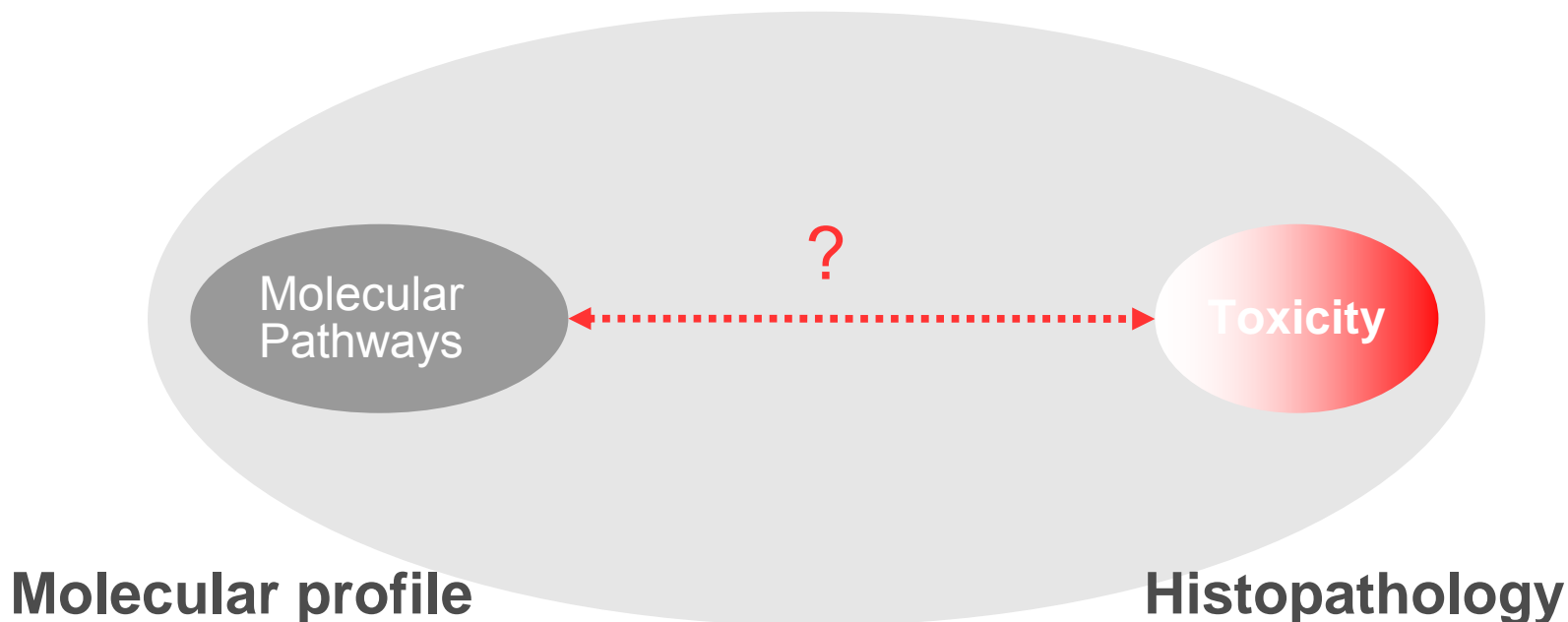
# Why Liver?

- **Primary** organ for environmental chemical **detoxification**
- **Most frequent** site of **adverse effects** (IRIS & ToxRefDB) in rodents – **relevant** to EPA
- **Human relevance** still uncertain
- **Large** amount of **available molecular** and **tissue** data
- **Unmet Need**: Extrapolation across doses, chemicals, species, populations

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



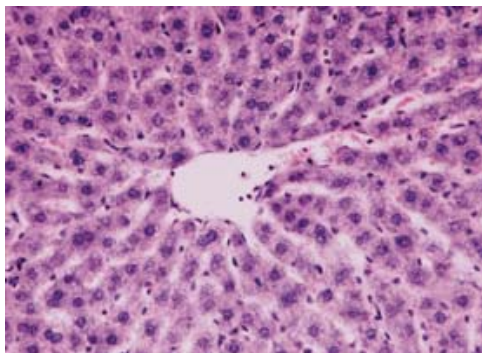
# How to Predict Liver Toxicity ?



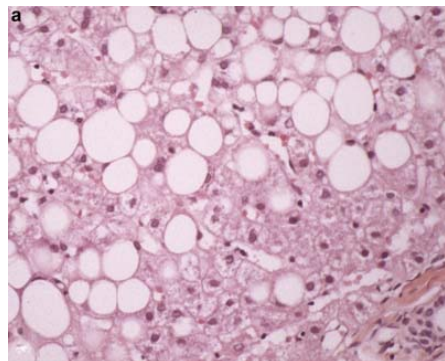
Phenotypic anchoring valuable for prioritization  
**Difficult to elucidate mode-of-action / dose-response**

# Toxicity: Cell Alteration

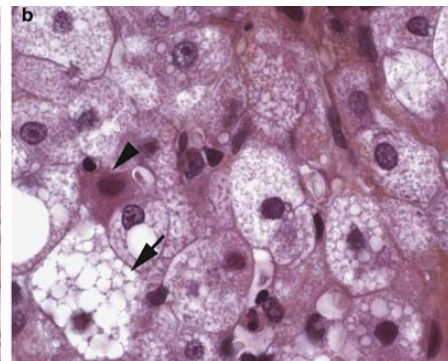
Swelling



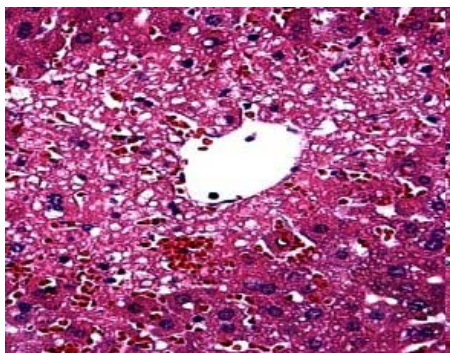
Steatosis, Macroves.



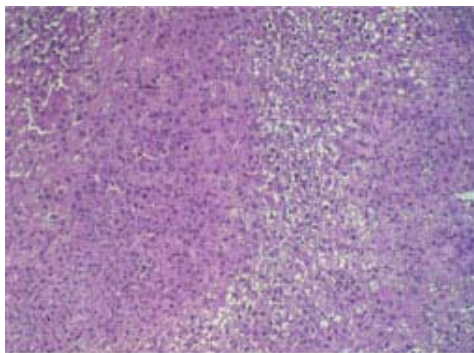
Steatosis, Microves.



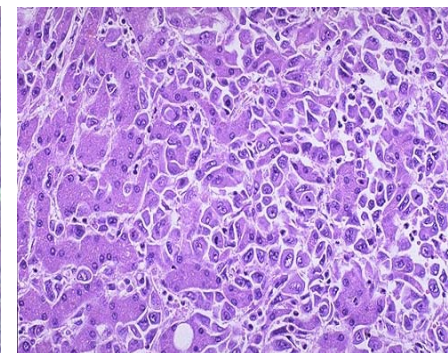
Necrosis



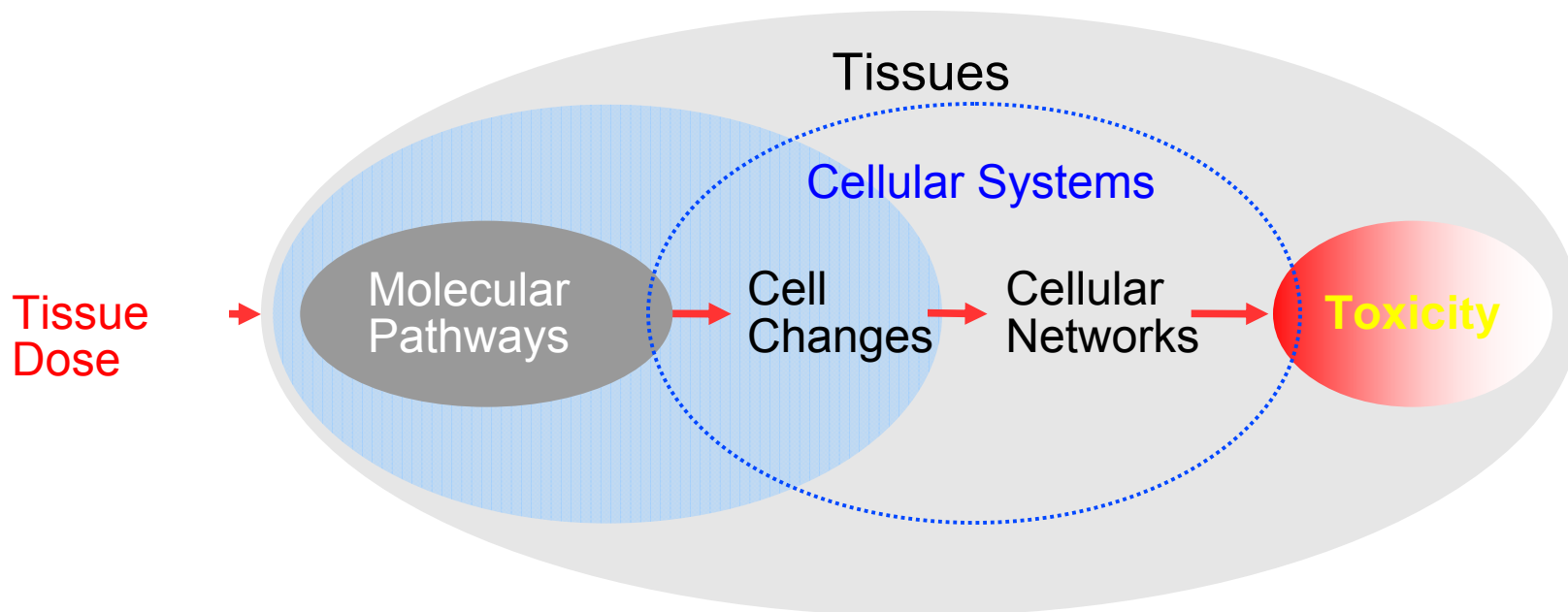
Hyperplasia



Carcinoma



# Toxicity: Cross-Scale Phenomena



Tissue lesions propagated by dynamic cellular networks

Cell changes are caused by molecular pathways

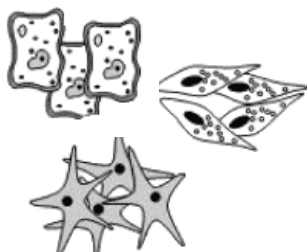
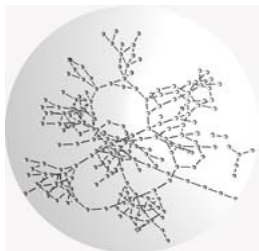
**Use a cell-oriented view to deal with complexity ...**

# Virtual Liver: v-Liver™

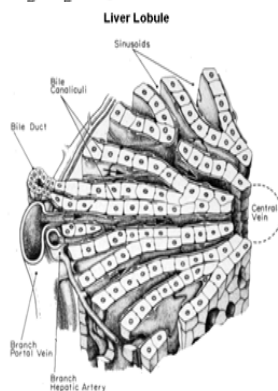
## Agent-Based Cellular Systems Model

### Tissues

Key events  
in cell  
response



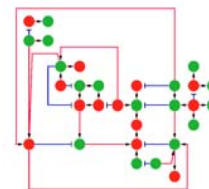
Cell  
Signaling



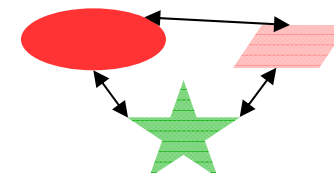
Blood  
Flow

### Virtual Tissues

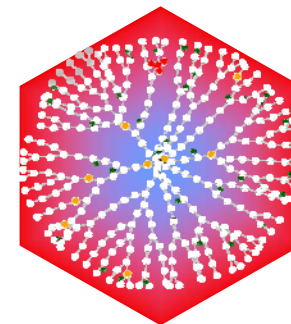
Molecular Logic  
Cell Changes



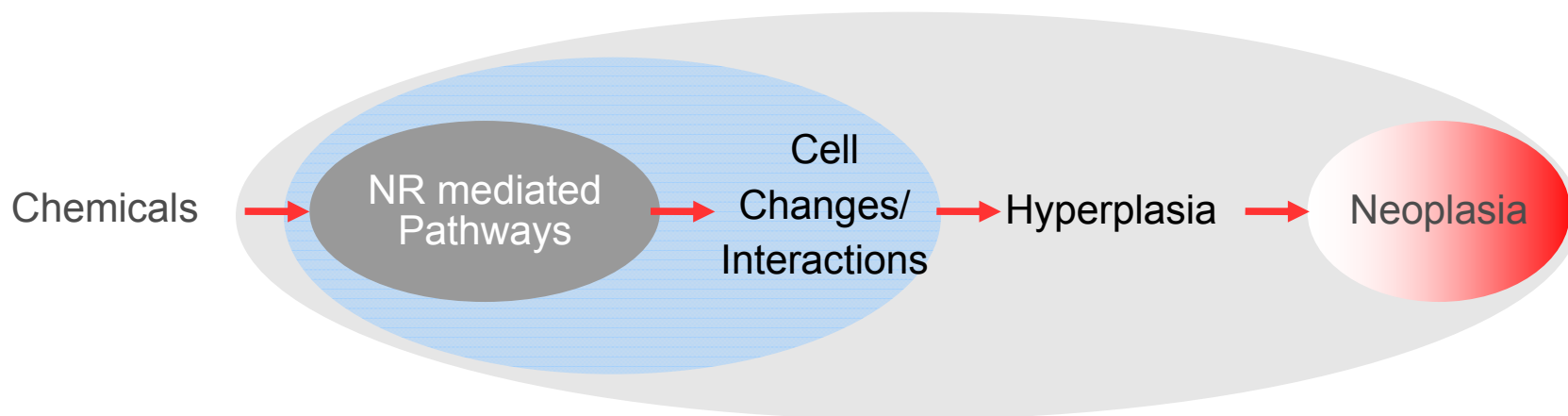
Cell  
Interactions



Spatial  
Cellular  
System



# v-Liver™: Proof of Concept



Goal: Simulate dose-dependent events in nuclear receptor-mediated proliferative lesion formation

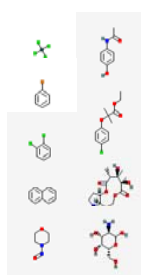
Modes-of-Action: Regenerative proliferation, Mitogen, DNA Damage

Approach: Cross-scale Cellular Systems Model

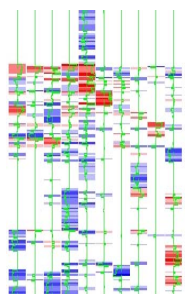
# v-Liver: Proof of Concept

- A) Select environmental chemicals from ToxCast Phase I
- B) Integrate knowledge on key cellular events in hepatocarcinogenesis
- C) Develop tissue simulation platform: key cell interactions and cell state changes
- D) Evaluate using PoC chemicals

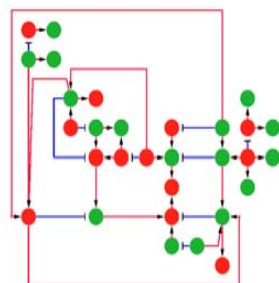
# v-Liver™ Architecture



Env.  
Chems



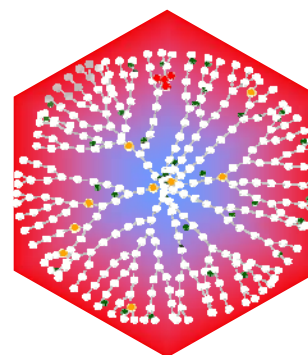
ToxCast  
HTS, HCS  
*ex vivo*



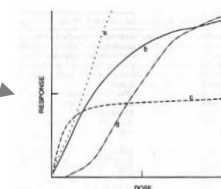
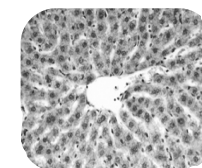
Molecular  
Events



Cell-Cell  
Events

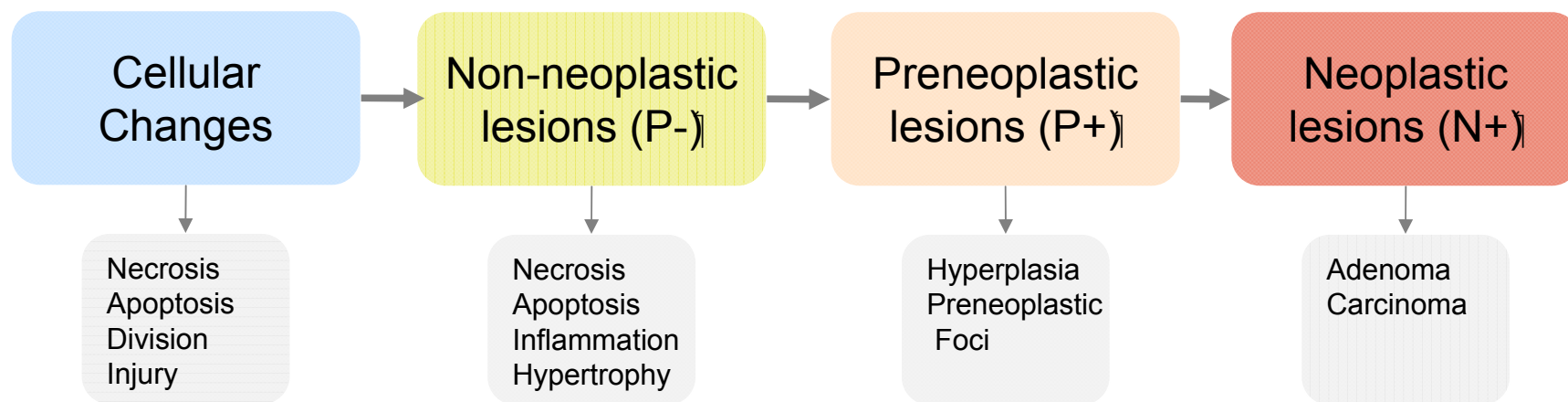


Cell Sys. &  
Blood Flow



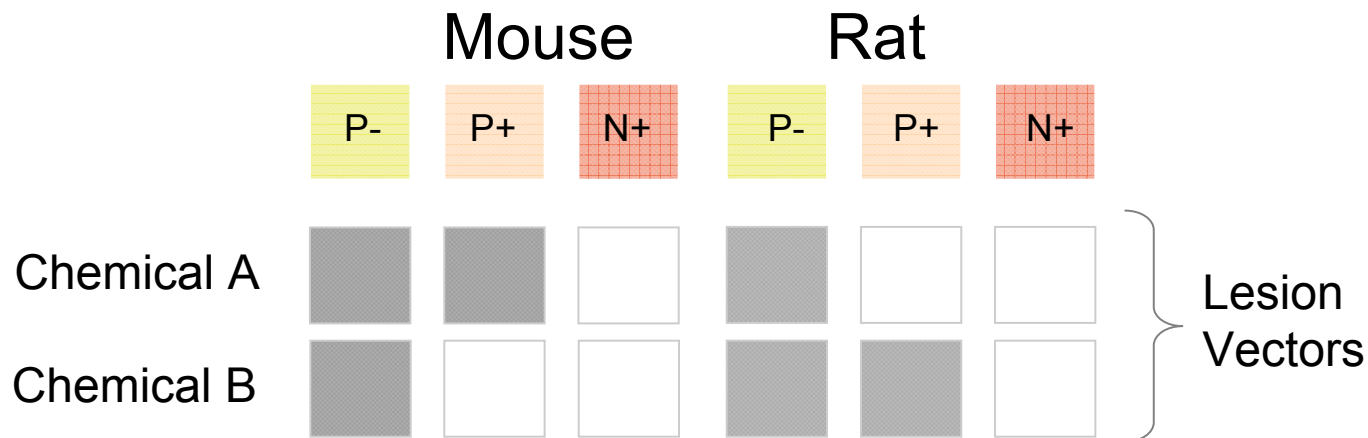
Cellular &  
Tissue Effects

# Chronic Toxicity: Cancer



- Multiple molecular and cellular pathways
- Incomplete information about mechanisms
- Carcinogenesis is chemical & dose dependent

# Data on Lesion Progression



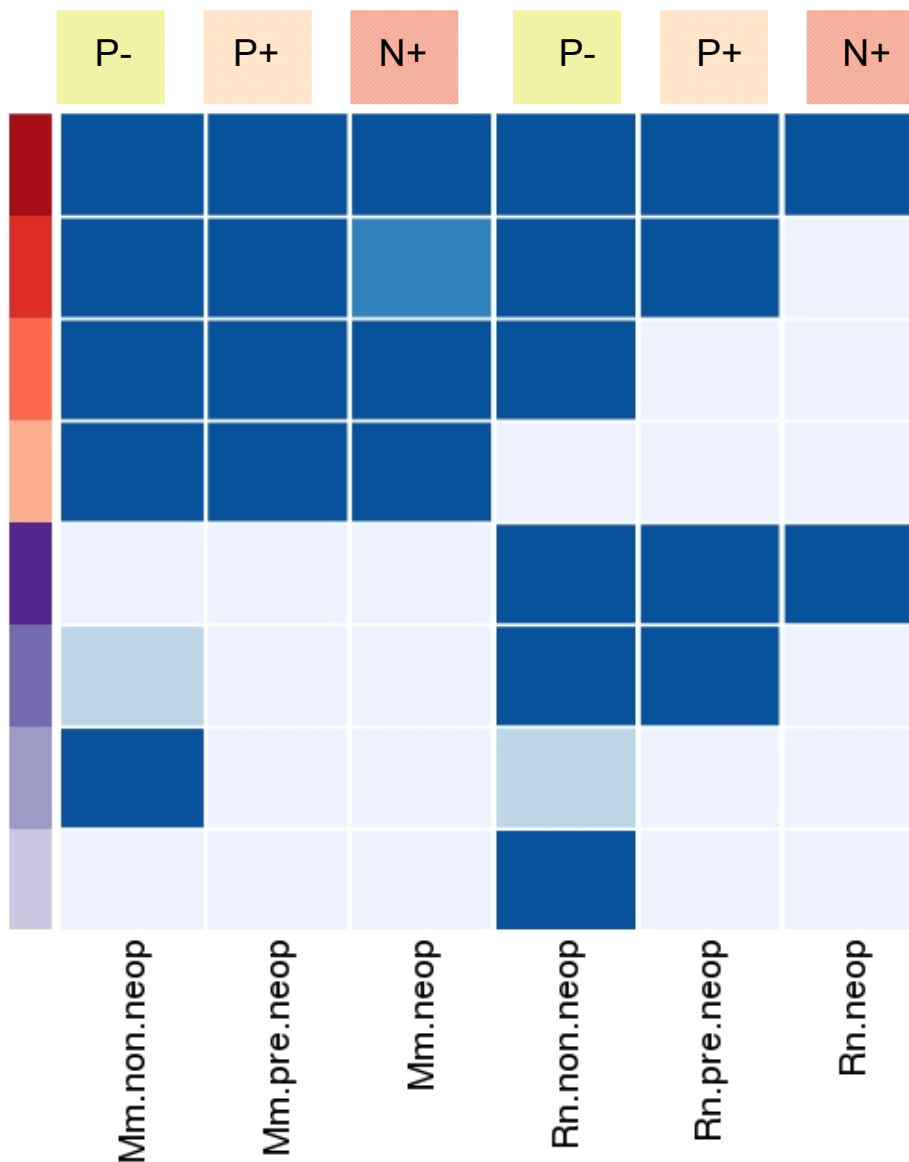
- Lesion progression can be species-dependent
- Summarize chemical-induced tissue effects across rat and mouse

# Cancer Lesion Progression

**ToxRefDB:**  
220 Chemicals

Rodent  
testing studies

Detailed  
histopathology



# Rodent Cancer & NRs

ToxRefDB  
*in vivo*

ToxCast™  
*in vitro*

Mouse Chronic Pathology  
Rat Chronic Pathology

Nuclear Receptor  
Activity

## ToxCast:

309 Chemicals

600 Assays

Chronic Pathology

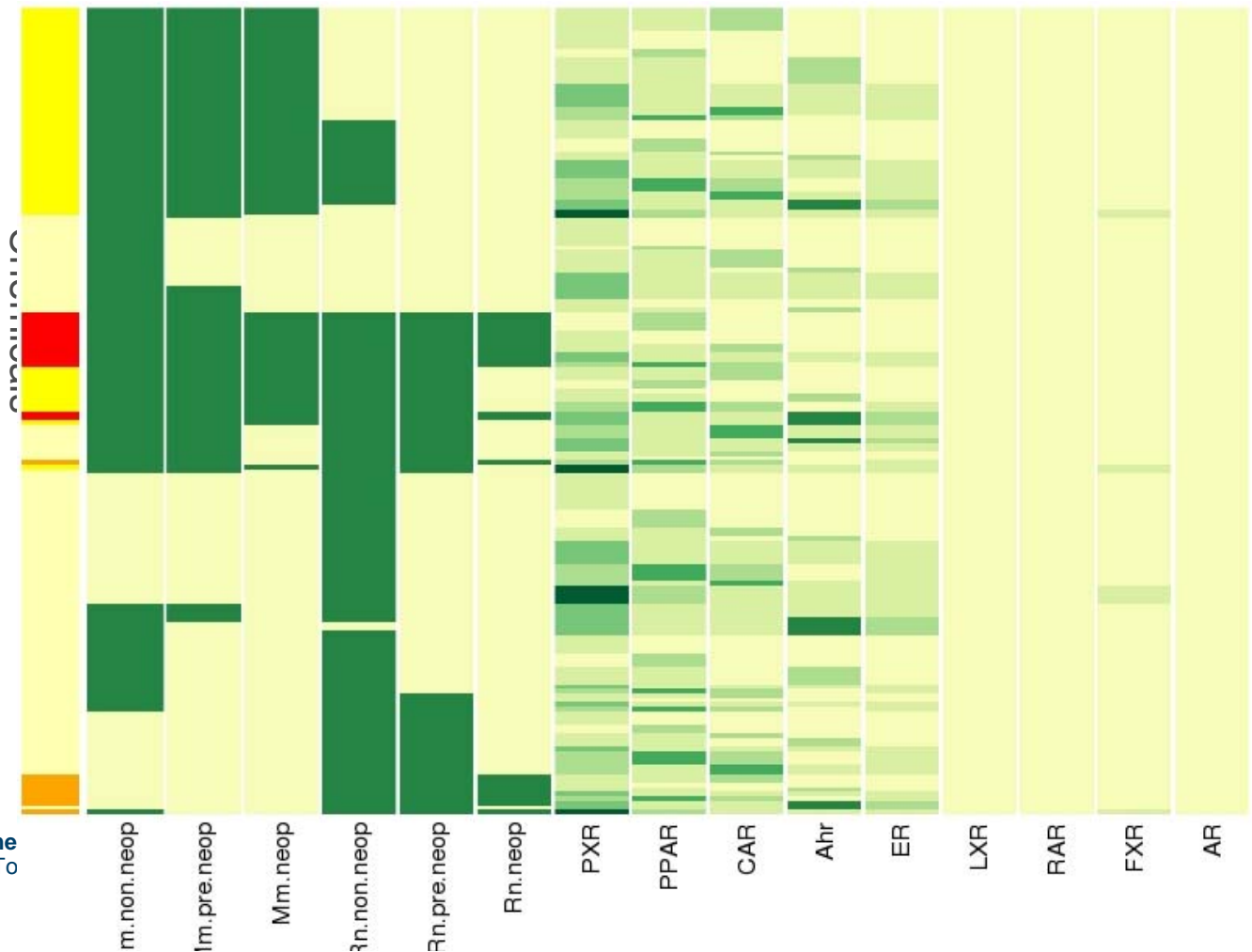
20 PoC Chemicals

Nuclear receptor +  
Chronic pathology +/-

Conazoles  
Pyrethroids

Phthalates

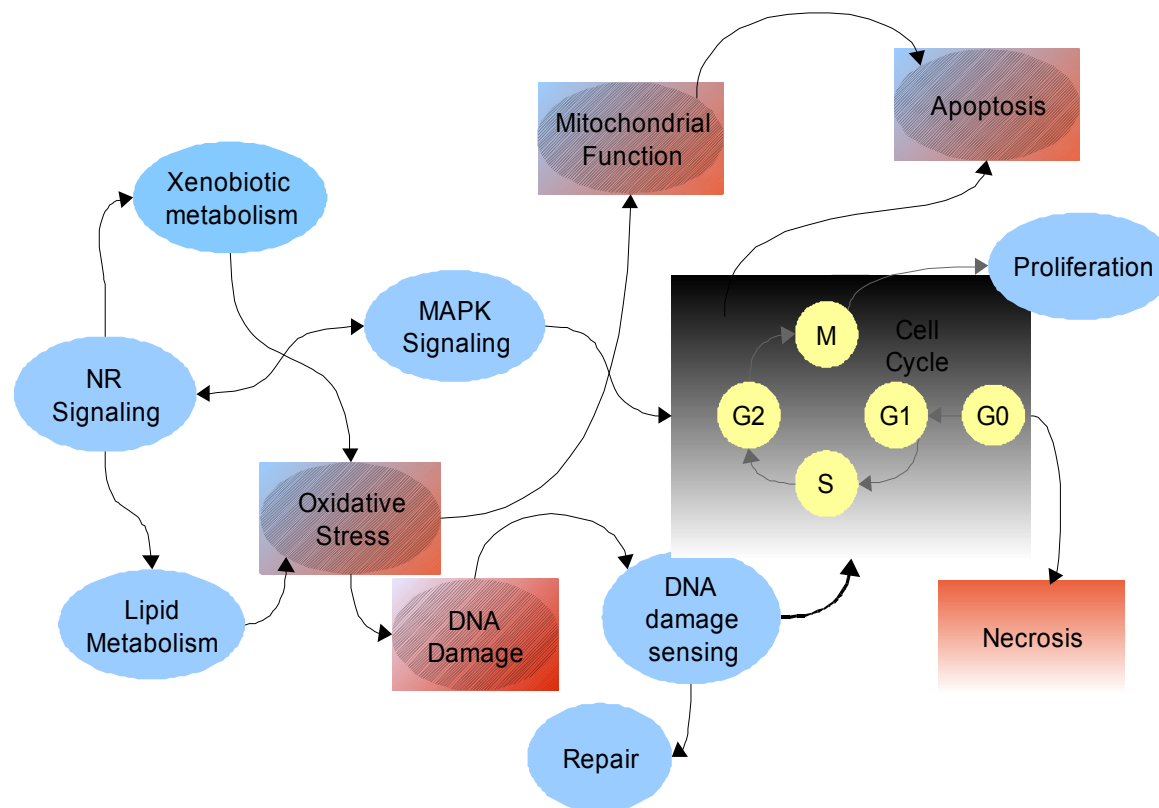
Perfluorinated



# Cellular Pathways: *Macrostates*

Mine literature,  
DBs and prior  
data:

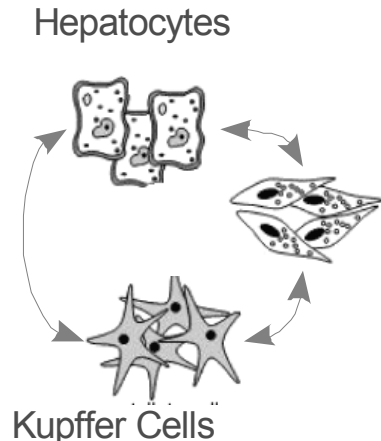
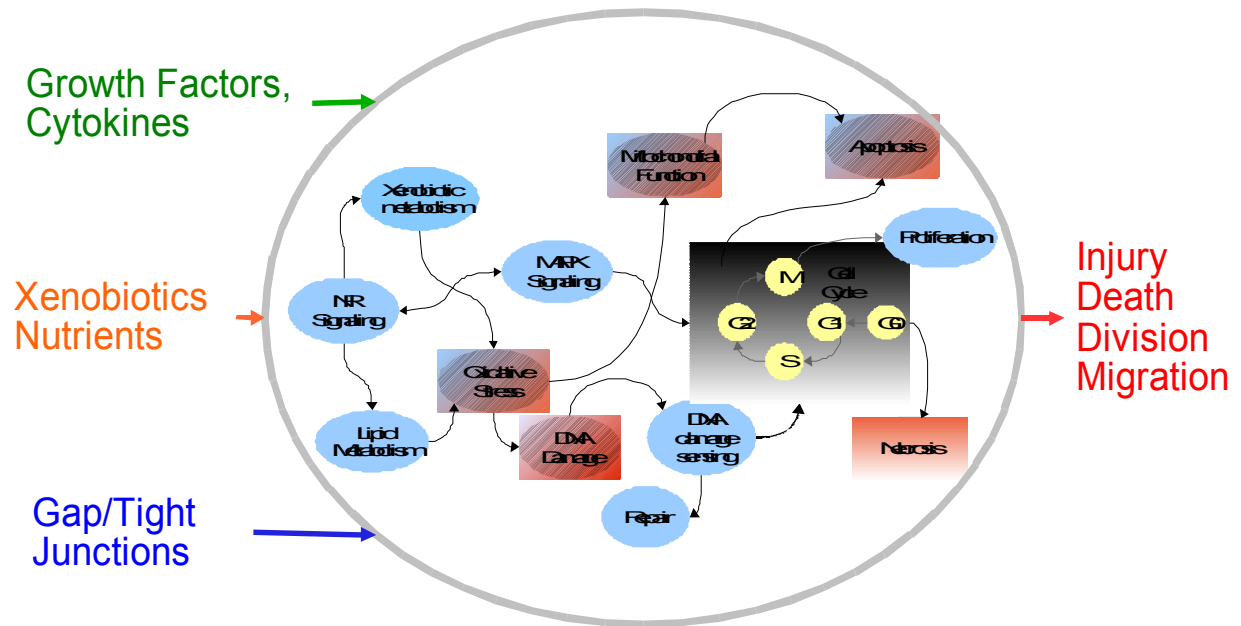
Cellular  
**pathways**  
involved in  
**death, division,**  
**initiation &**  
**promotion**



# Cellular Interactions

Mine literature,  
DBs and prior  
data

Cells &  
interactions  
involved in  
cancer  
progression



# KB: Cytoscape Plugin

Configure Database

Enter a SPARQL query service address.

URL:

Save

Specify namespaces

Namespace	URI
owl	http://www.w3.org/2002/07/ontology
ti	http://www.epa.gov/ncct/oliver
bp2	http://www.biopax.org/release/20090101
bp	http://www.biopax.org/release/20090101
rdfs	http://www.w3.org/2000/01/rdf-schema
rdf	http://www.w3.org/1999/02/22-rdf-syntax-ns
xsd	http://www.w3.org/2001/XMLSchema
tp	http://www.epa.gov/ncct/oliver
nci	http://pid.nci.nih.gov/biopax#
rx	http://www.reactome.org/biopax#
bn3	http://www.biopax.org/release/20090101

Apply QNames to Search Results

Add New Remove Selected Save

Cytoscape Desktop (Session: net-02-19-09.cys)

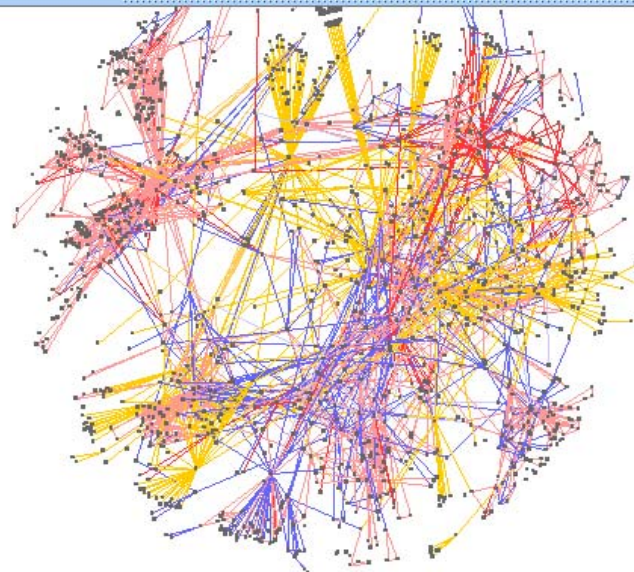
File Edit View Select Layout Plugins Help

Search:

Control Panel

Network	Nodes	Edges
Mantic Network	24(0)	39(0)

HepatoCyc-mol-int-1



C-Mantic Plug-in

Create Network

Use SPARQL queries to create Cytoscape networks.

Specify a data source

Define namespace prefixes

Enter and execute a query

Save Network

Save a Cytoscape network as an RDF file.

Query

Query File:

```
construct
{?a ?b ?x .?x bp2:ID ?z . ?x rdf:type ?y }
where
{ ?a ?b ?x .
?x rdf:type ?y .
?x bp2:ID ?z .
filter(str(?z)="200958") }
```

Subject	Predicate	Object
rx:UniProt_P20815_RecName_Full_Cyto...	bp2:SYNONYMS	"CYP3A5"^^<http://www.w3.org/2000/01/rdf-schema#type>
rx:UniProt_P20815_RecName_Full_Cyto...	bp2:SYNONYMS	"CYP3A5"^^<http://www.w3.org/2000/01/rdf-schema#type>
rx:UniProt_P20815_RecName_Full_Cyto...	bp2:SYNONYMS	"CYP3A5"^^<http://www.w3.org/2000/01/rdf-schema#type>
rx:UniProt_P20815_RecName_Full_Cyto...	bp2:SYNONYMS	"CYP3A5"^^<http://www.w3.org/2000/01/rdf-schema#type>
rx:UniProt_P20815_RecName_Full_Cyto...	bp2:SYNONYMS	"CYP3A5"^^<http://www.w3.org/2000/01/rdf-schema#type>
rx:UniProt_P24462_RecName_Full_Cyto...	bp2:SYNONYMS	"CYP3A7"^^<http://www.w3.org/2000/01/rdf-schema#type>
rx:UniProt_P24462_RecName_Full_Cyto...	bp2:SYNONYMS	"CYP3A7"^^<http://www.w3.org/2000/01/rdf-schema#type>

Rotate in Degrees:

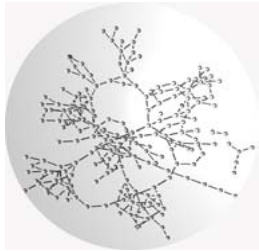
0 90 180 270 360

Rotate Selected Nodes Only

Node Attribute Browser Edge Attribute Browser Network Attribute Browser C-Mantic Results

# Virtual Hepatic Lobule

## Knowledgebase

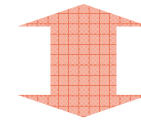
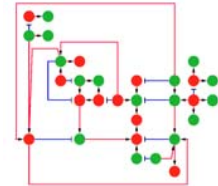


Key events  
in cell  
response

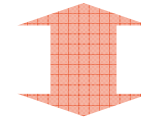
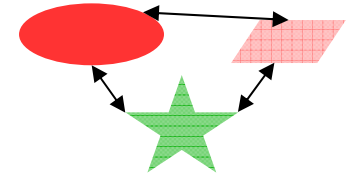
*In vitro*  
data

## Tissue Simulator

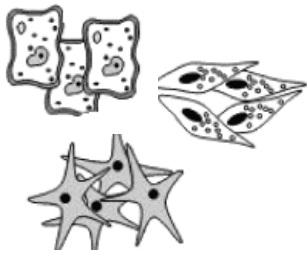
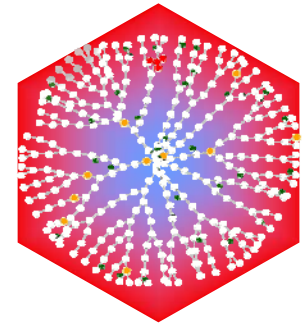
Molecular Logic  
**Agent Decisions**



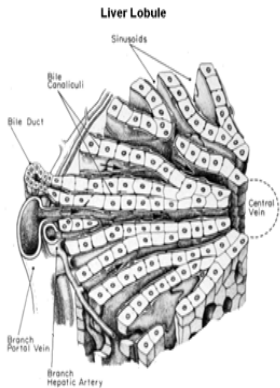
**Agent  
Communication**



**Multi-Agent  
System**

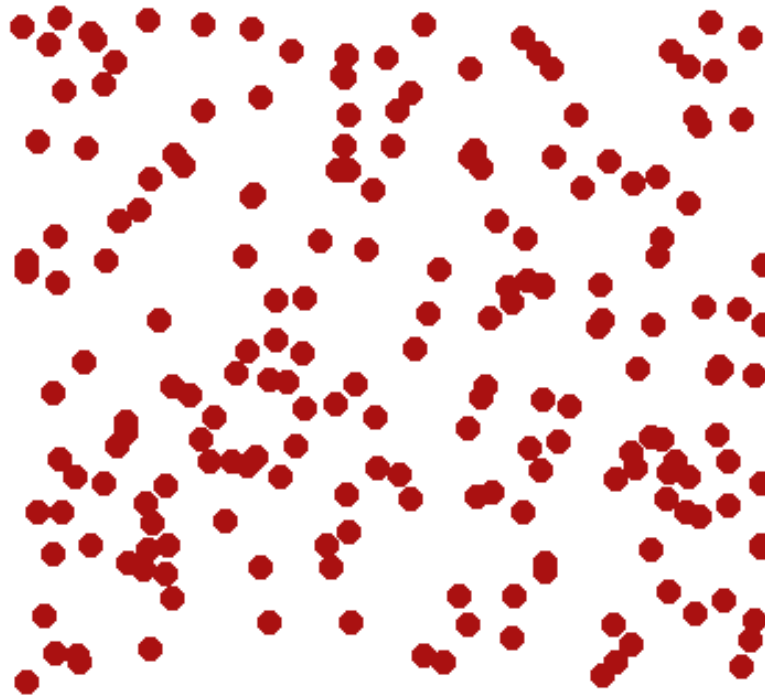


Cell  
Signaling

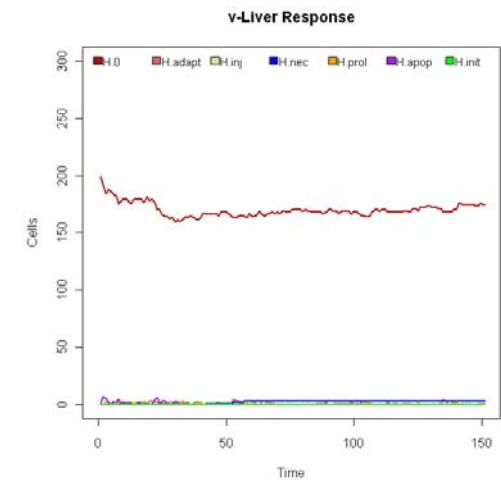
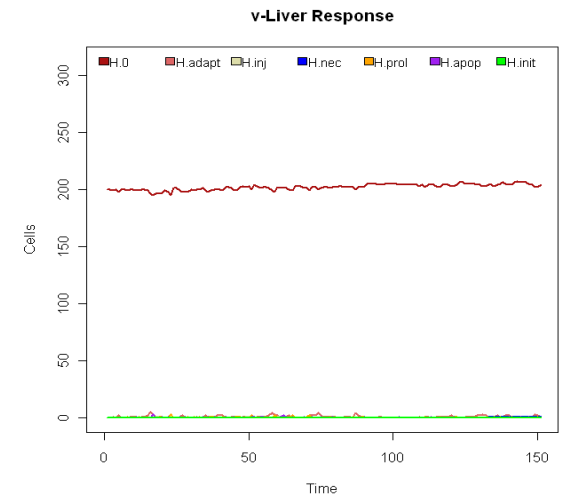


Blood  
Flow

# Simulate *in vitro* Hepatocytes

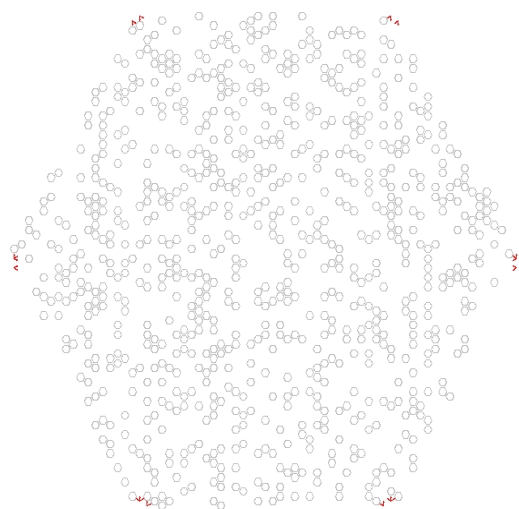


- Normal/Quiescent
- Stressed/Adaptive
- Stressed/Injured
- Necrotic
- Proliferative
- Apoptotic

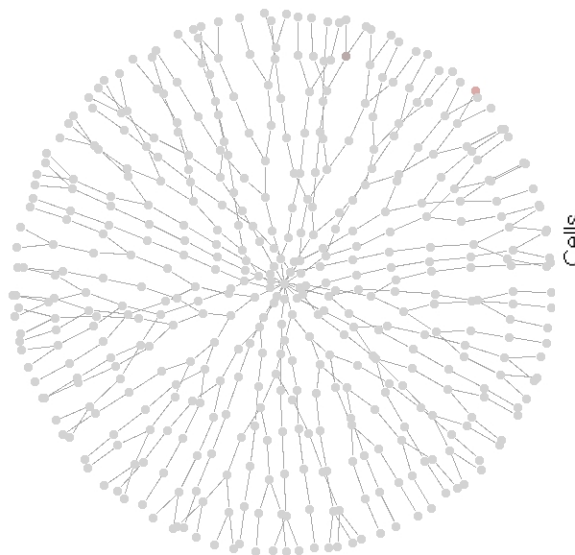


# Simulating Chemical Effects

## Microdosimetry

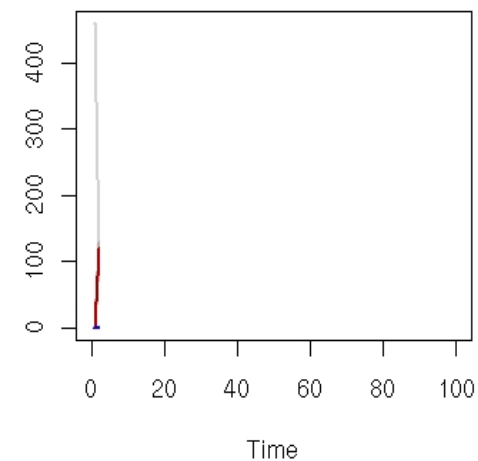


## “Virtual Tissue”



Multi-agent system:  
spatial distribution of  
agents with tight  
junctions

## Simulated Outcome



Dynamics of multi-  
agent system: temporal  
changes in cell death/  
proliferation

# Challenges

- Dealing with incomplete knowledge
- Using *in vitro* data in an *in vivo* context
- Dealing with time across molecular and cellular scales
- Evaluating *in vivo* predictions – how do we calibrate and evaluate tissue models ?



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

John Wambaugh  
 Jason Pirone  
 Mike DeVito  
 Nicholas Luke  
 Rory Conolly  
 Thomas Knudsen  
 David Dix  
 Matt Martin  
 Keith Houck  
 Richard Judson  
 David Reif  
 Woody Setzer  
 Amar Singh  
 Lockheed Martin

Risk Assessors:  
 Clients

