

A Virtual Liver for Simulating Chemical-Induced Injury

John Wambaugh¹, John Jack¹, Chris Corton², and Imran Shah¹ ¹National Center for Computational Toxicology (NCCT), US EPA, RTP, NC, USA.

²National Health and Environmental Effects Research Lab (NHEERL), US EPA, RTP, NC, USA.

Introduction

Humans are exposed to over 6,000 environmental chemicals The liver is the primary organ for metabolism and often the first site of chemical-induced toxicity in animal testing. It remains difficult to translate these outcomes to humans due to uncertainties in extrapolating pathways from rodents to humans, and from in vitro to in vivo. The Virtual Liver (v-Liver[™]) is an *in* silico platform aimed at simulating clinically-relevant effects in the liver. The proof of concept is defined by 20 chemicals (VL-20) with ToxCast[™] in vitro assay data that produce a range of chronic liver lesions in rodents. The VL-20 are high-production volume (HPV) chemicals including: pesticides, persistent toxic substances, and plasticizers.

Objectives

• Develop decision support tools to extrapolate from rodents to humans and in vitro to in vivo

· Create a knowledgebase (KB) of literature-derived information on pathways to tissue lesions for VL-20

· Simulate a virtual hepatic lobule to quantitatively evaluate tissue outcomes for VL-20

Timeline

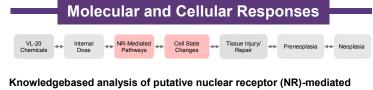
2-10

- 2009 Prototype Tissue Simulator (Draft manuscripts prepared)
- Engage with EPA Program Offices 2010 on MOA analysis of hepatocarcinogens

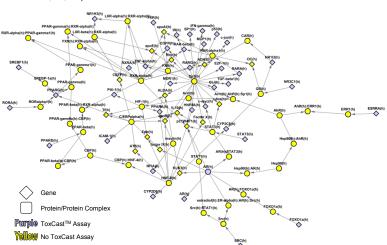
Simulate Lesion Formation for for VL-20 using ToxCast[™] data and predictions with ToxRefDB

Work with EPA Program Offices to 2011 evaluate quantitative effects of new chemicals

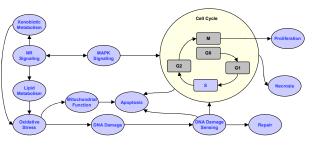
> Use system to evaluate hepatic effects across human subpopulations / genetic variation



molecular interactions. (Data on human NR activation for VL-20 from ToxCast™ shown in



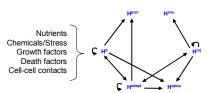
NR-mediated cellular changes. (Putative relationships between NR-activity to cellular processes using data from ToxCastTM (shown in p



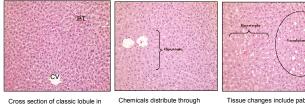
Probabilistic model of hepatocellular response to chemicals.

 Initially we are using an automaton model of cell response to microenviornment including: chemical (stress), nutrient, growth factors and cell-cell contacts.

· Based on histological observations define six hepatocellular states: normal, adapting, injured, necrotic, apoptotic or proliferating



Hepatic Microanatomy and Toxicity



Cross section of classic lobule in Untreated female rat liver: blood flows from portal triads (PT) into orally treated with 2500 ppm he central vein (CV) propiconazole for 12 weeks. hepatomegaly and centrilobula hypertrophy were observed. Images from Rockett et al. (2006)

Tissue changes include pablobula sinusoidal network. In female rate hypertrophy and cytoplasmic vacuolation for female rats orally exposed 1800 ppm triadimefon for

. Developing an agent-based spatial model of a single classical hepatic lobule in 2D

• Hepatic parenchymal and non-parenchymal cells represented as agents arranged according to lobular morphology.

· Agent responses and states defined by probabilistic hepatocellular model

 Sinusoidal network represented graphically by sinusoidal primitives. arterial and venous sources, and the central vein

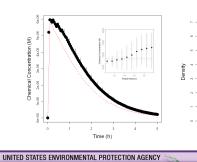
• Flow through the sinusoids modeled by ordinary differential equations (ODEs) - efficient for large systems vs. PDEs

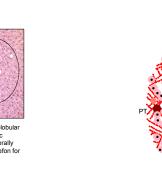
 Graphical model allows flexibility for dealing with histologic changes over doses and times

Integration with *in vitro* Data (e.g. ToxCast[™])

· Variability in predicted local concentration due to the action of hepatocytes, e.g., metabolism, and not geometry alone.

• in vitro testing can determine thresholds for endpoints such as cytotoxicity under controlled dosing conditions - we use our model to determine local conditions for individual hepatocytes





Cellular Changes to Tissue Effects

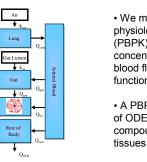
Cell State

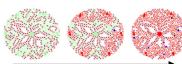
NR-Mediated

aight Aggregate Node

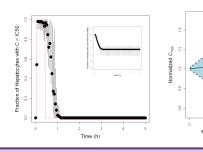
~1 mm

Relating Environmental Exposure to Local Cellular Environment for Risk Assessment





time after single exposure



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research&development

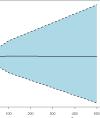


• We make use of a simple physiologically-based pharmacokinetic (PBPK) model structure to find the concentration of compound(s) in the blood flowing into the liver as a function of environmental exposure

 A PBPK model consists of a system of ODEs for the concentration of a compound or compounds in different



in this simulation exposure to enhances cytotoxicity above a threshol



Metabolic Clearance (µL/min/10⁶hep.)

Impact and Outcomes

Simulation can enhance prioritization efforts by providing context for HTS data

Modeling liver homeostasis will provide better insights into the dose response relationship for environmentally-relevant concentrations

Organizing and modeling hepatic biology can improve identification of toxic modes of action for risk assessors in NCEA and outside ORD

The flexibility of an in silico approach allows for interspecies extrapolation and novel simulated experiments

Ultimately the v-Liver[™] should reduce animal testing

Future Directions

Estimate tissue dosimetry for VL-20 to effectively utilize in vitro assay data from ToxCast

Link NR-mediated gene-networks for VL-20 to key cellular pathways involved in their MOA. Encode these pathways in Agent-Based model. Calibrate and Evaluate using in vitro assays

Enhance agent-based model to consider relevant parenchymal and non-parenchymal cell interactions to model homeostatic liver function. Calibrate and Evaluate using in vivo histomorphometry data

Augment sinusoidal network to include additional vasculatures (e.g. bile ductules and lymphatics)

References

JC Rockett, MG Narotsky, KE Thompson, . I Thillainadarajah, CR Blystone, AK Goetz, H Ren, DS Best, RN Murrell, HP Nichols, JE Schmid, DC Wolf and DJ Dix, "Effect of conazole fungicides on reproductive development in the female rat ' Reproductive Toxicology 22, 647-658 (2006)

JF Wambaugh and I Shah, "A Microdosimetry Model for Virtual Liver Simulation," in preparation.

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