

Characteristics of the ToxCast In Vitro Datasets from Biochemical and Cellular Assays

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- Data Sets: Bio-centric View
 - Assay overview
 - How were the data calculated?
 - How did the replicates perform?
 - Example results
 - What did the assay measure?
 - Potential assay artifacts

ToxCast Phase I Data Sets

- Biochemical
 - Novascreen
- Cell-Based
 - ACEA
 - Attagene
 - BioSeek
 - Cellumen
 - CellzDirect
 - Gentronix
 - NCGC
 - Solidus

Novascreen: 239 Biochemical Assays (Abstract 59)

- Protein super-families
 - GPCR
 - Kinase
 - Phosphatase
 - Protease
 - Ion channel
 - Nuclear receptor
 - Other enzyme
 - CYP P450 inhibition
- Various formats:
 - Radioligand receptor binding
 - Fluorescent receptor binding
 - Fluorescent enzyme substrate-intensity quench
 - Fluorescent enzyme substrate-mobility shift
- Initial screening:
 - 25 μ M in duplicate
 - 10 μ M in duplicate (CYPs)
- Normalize data to assay window
 - % of control activity (central reference – scalar reference)

Novascreen: What do the assays measure?

- Mainly direct effects of chemical on target protein
 - Enzyme activity
 - Ligand binding
- False positives:
 - Fluorescent compounds—fluorescing and quenching
 - Reactive compounds/covalent modification of target
 - Physical effects—colloid aggregation of target
 - Operational
- False negatives:
 - Solubility
 - Inappropriate enzymatic conditions
 - Operational
 - Target protein not physiological
 - Lack of biotransformation

Data Correction Algorithm Examples (Additive)

AChE

	1	2	3	4	5	6	7	8	9	10	11	12
A	-3	3	-29	-100	-98	-100	-68	-68	-89	-89	-98	-98
B	-4	7	-29	-12	-10	-12	-27	-22	-30	-30	-25	-16
C	8	2	-23	-23	-19	-26	-25	-26	-26	-28	-25	-13
D	-14	1	-17	-21	-23	-25	-18	-26	-26	-27	-25	-17
E	-11	-5	-23	-25	-16	-20	-26	-22	-27	-30	-26	-13
F	-16	0	-7	-7	-17	-21	-22	-24	-29	-33	-19	-17
G	-5	4	-32	-18	-25	-17	-25	-28	-25	-32	-20	-13
H	-85	-78	-22	-22	2	-14	-17	-17	NaN	NaN	NaN	NaN

normalized

	1	2	3	4	5	6	7	8	9	10	11	12
A	-1	1	-19	-100	-98	-100	-67	-67	-89	-89	-98	-98
B	0	7	-20	-10	-2	0	2	9	-15	-16	-9	-4
C	10	10	-3	-3	0	-6	-2	-5	-16	-18	-11	-1
D	4	14	7	2	2	-5	6	0	-13	-11	-9	-3
E	7	13	4	0	7	4	-2	2	-15	-18	-10	-1
F	5	22	16	16	9	6	3	3	-15	-17	-3	-2
G	12	23	-4	6	1	5	3	0	-11	-16	-1	4
H	-52	-44	7	9	20	8	4	5	NaN	NaN	NaN	NaN

Normalized
& corrected

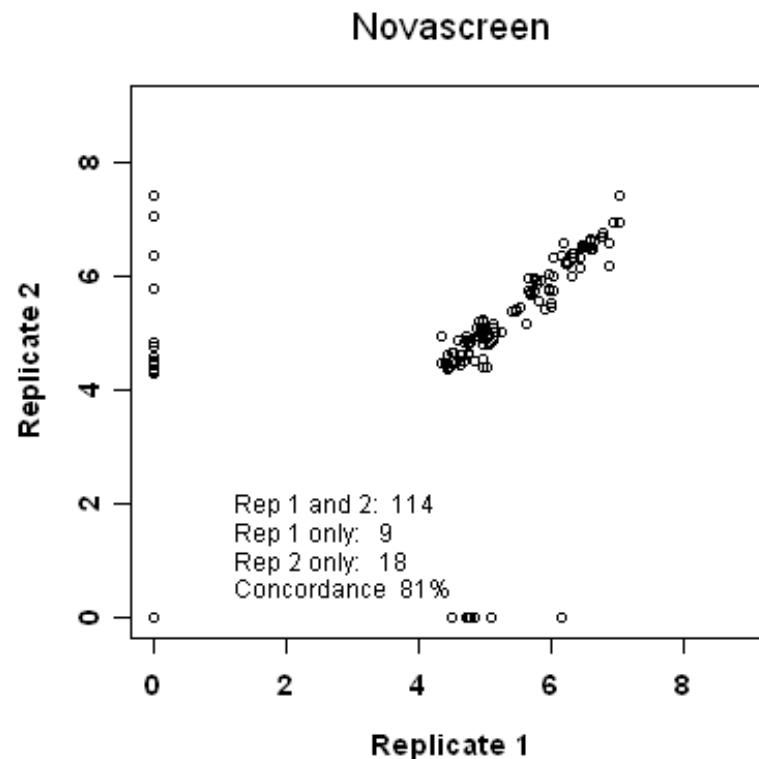
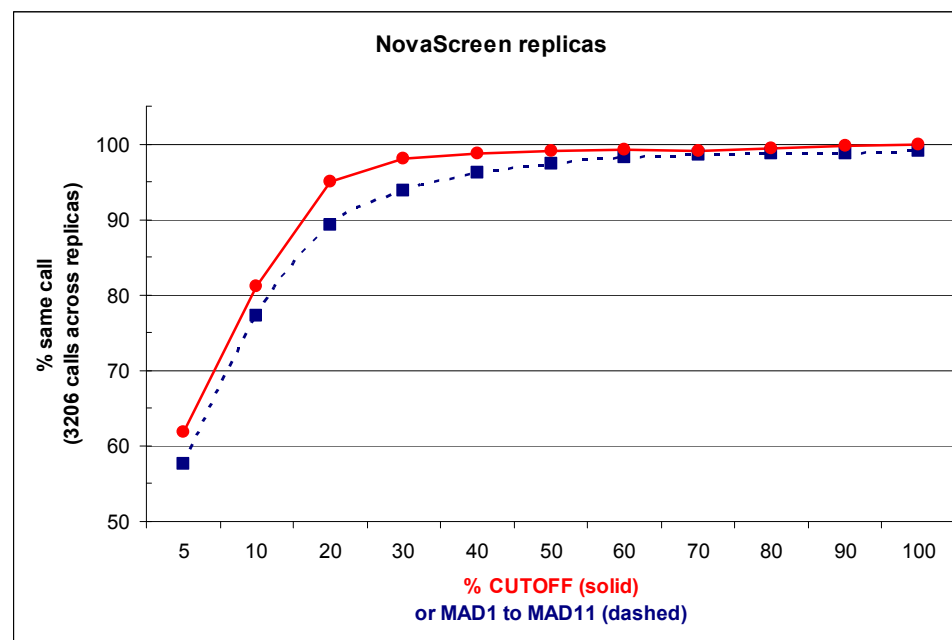
Caspase 10

	1	2	3	4	5	6	7	8	9	10	11	12
A	25	0	-3	-97	-100	-101	-35	-33	-76	-80	-100	-99
B	9	-17	-16	-23	-28	-17	-13	-16	-22	-15	-22	-3
C	3	-22	-10	-27	-24	-20	-39	-31	-24	-28	-26	-6
D	15	-22	-15	-27	-23	-27	-13	-15	-17	-18	-20	-3
E	3	-16	-16	-17	-18	-7	-6	-13	2	5	-17	12
F	-11	-34	-18	-26	-29	-21	-22	-19	-18	-19	-25	0
G	-23	-30	-27	-24	-17	-6	-8	-11	-22	-12	-9	10
H	-15	-28	-17	-20	-14	-28	-5	4	-19	-4	-18	5

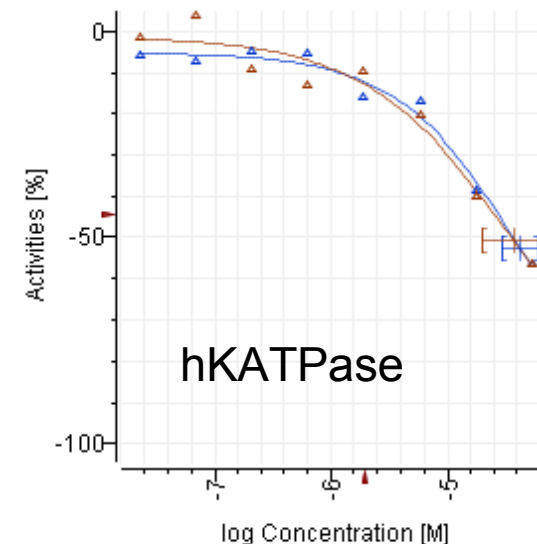
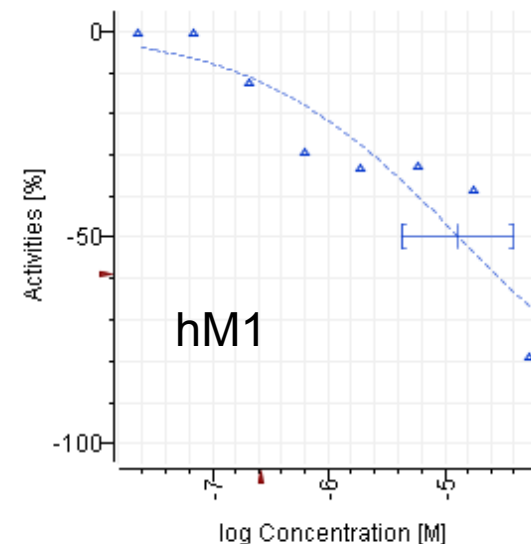
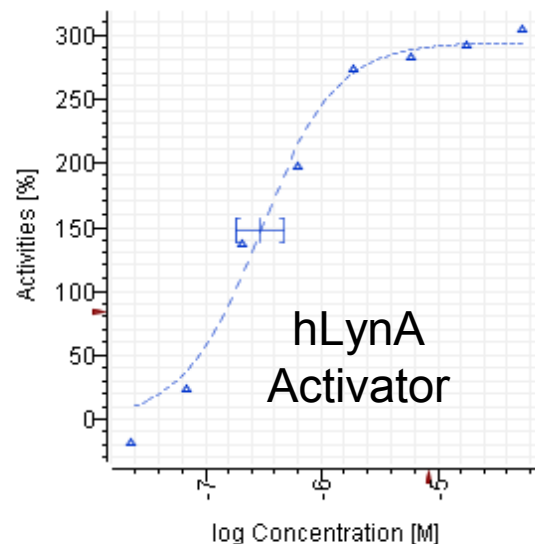
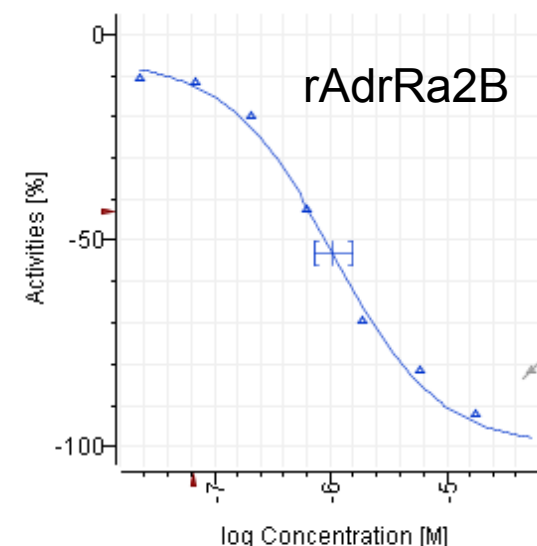
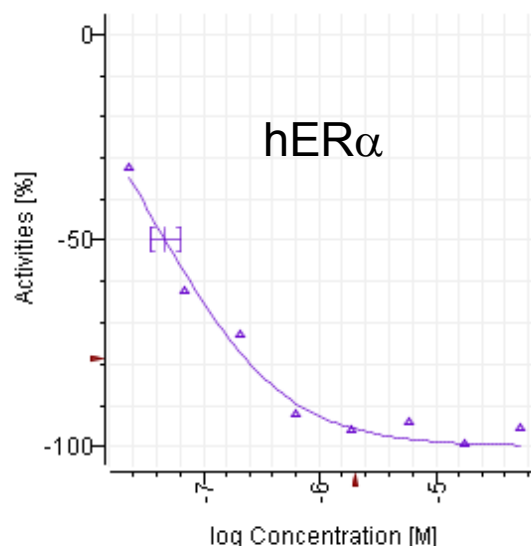
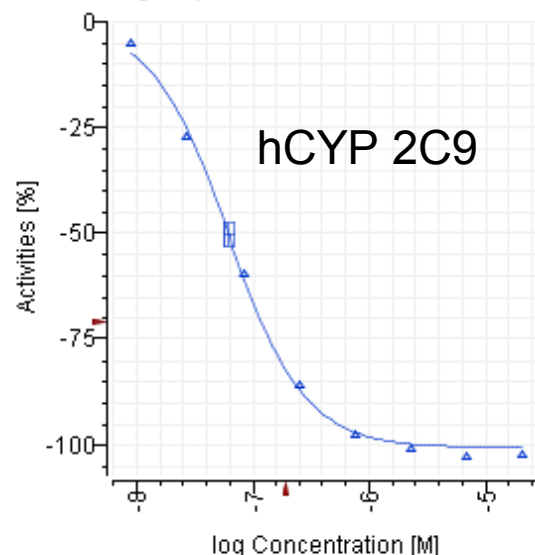
	1	2	3	4	5	6	7	8	9	10	11	12
A	23	0	-3	-97	-100	-101	-36	-34	-76	-81	-100	-99
B	13	6	7	0	-6	-3	3	2	-6	1	-5	0
C	6	-1	10	-7	-4	-5	-21	-13	-5	-10	-7	-5
D	18	0	2	-7	0	-7	6	4	0	-1	-4	1
E	1	1	2	-3	-5	-2	7	0	14	17	-6	9
F	-9	-12	4	-3	-11	-9	-4	-1	-7	-4	-11	-1
G	-17	-5	-5	-4	-1	-1	13	6	-2	3	5	8
H	-9	-11	2	0	4	-14	9	15	-4	10	-7	8

Novascreen Concentration-Response Testing

- Retest actives:
 - Median absolute deviation (MAD)
median $|x - x_{med}|$
two MADs or 30% activity
 - 8 conc/3-fold serial dilutions
 - 50 μ M high conc
 - 25 μ M high conc for CYPs
- Normalize to assay window
- Fit % Activity data to 3- or 4-parameter Hill function
 - Sometimes had to fix top or bottom of curve
 - Did not extrapolate beyond testing range
 - Manual or automated removal of obvious outliers

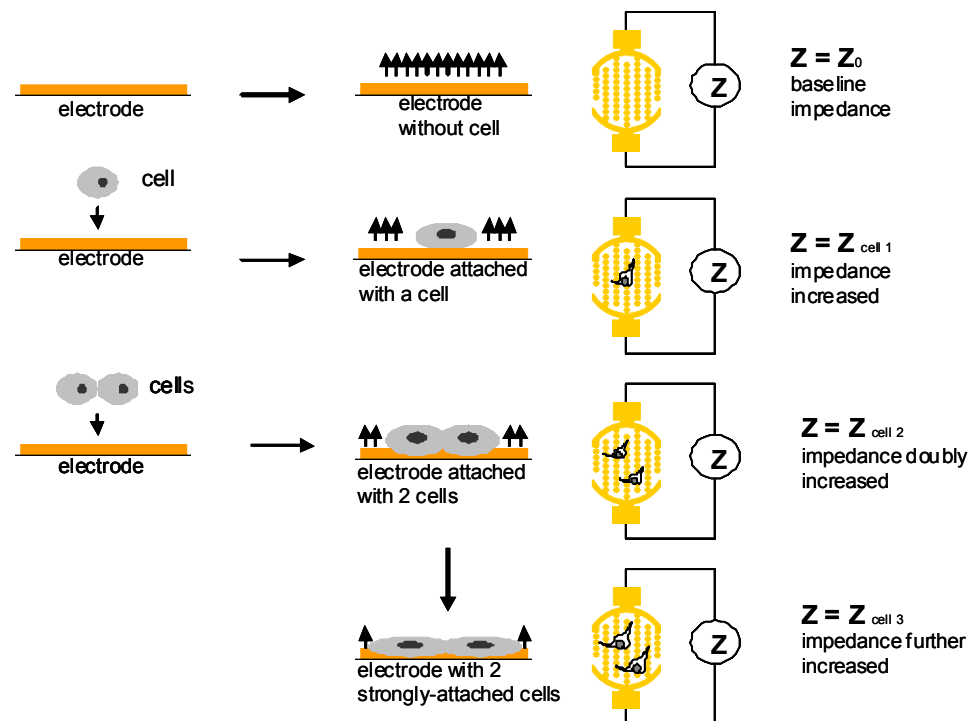


Novascreen: Example Curve Fits



ACEA: Real Time Cell Growth Kinetics

- Cytotoxicity with potential mechanistic interpretation
- Human A549 lung carcinoma cell line
 - ACEA experience with line
 - Reference compound effects
- Concentration-response testing
 - 8 conc/3-fold serial dilutions
 - Duplicate wells
- Real-time measurements during exposure (0-72 hr)
- IC50 and LECs calculated

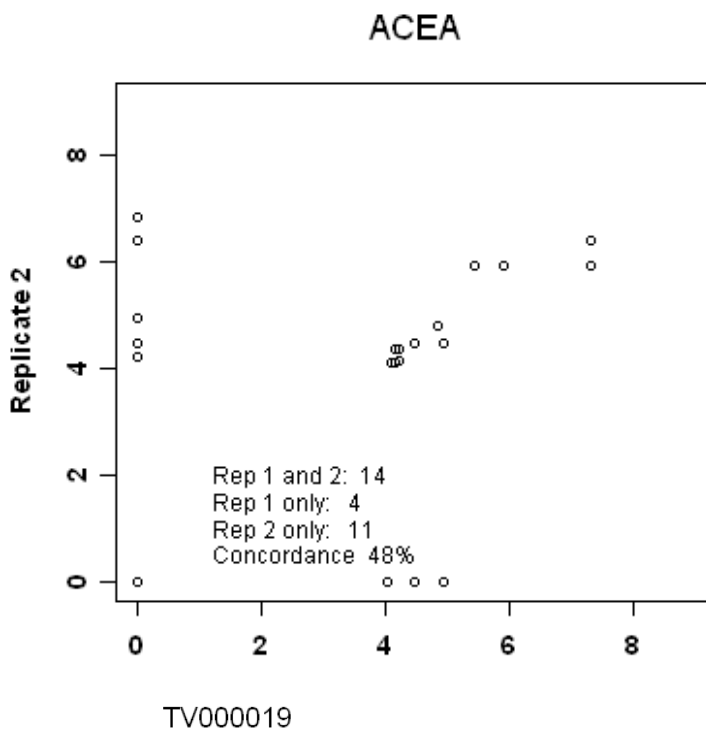


ACEA: What is measured?

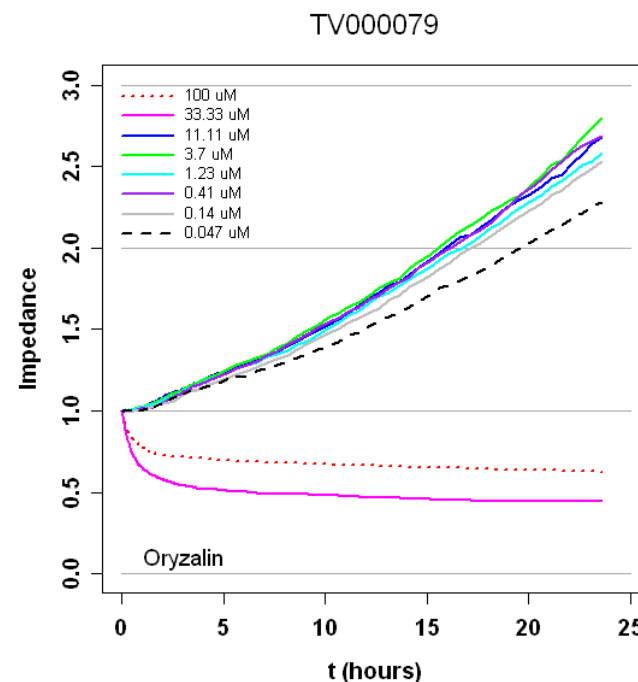
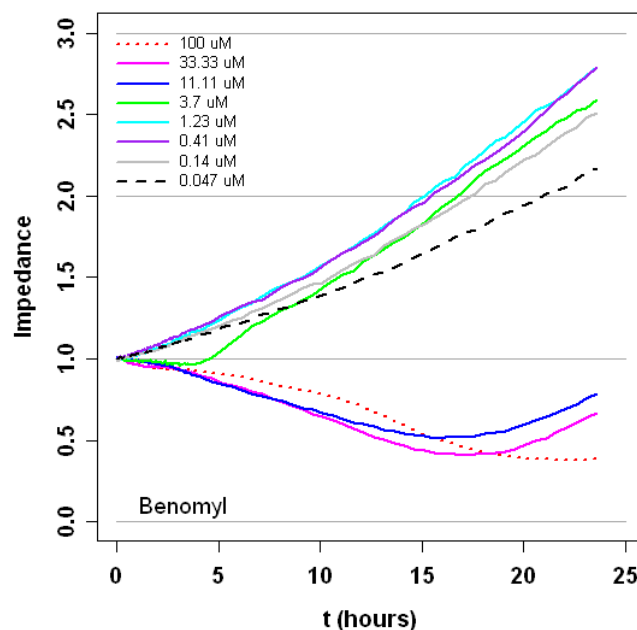
- General cytotoxicity in transformed cell line
- False positives
 - None obvious
 - Operational
- False negatives
 - Operational error
 - Solubility
 - Lack of appropriate toxicity targets (irrelevant cell line)
 - Lack of biotransformation

ACEA: Data Examples

Replicate Analysis:



Example Plots:



Attagene Multiplexed Transcription Reporter Assays

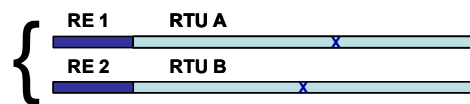
- Modulation of TF activity in human hepatoma HepG2 cells
- Multiplexed reporter gene assay
 - *cis* 52 assays (response element driving reporter)
 - *trans* 29 assays (GAL4-NR_LBD driving reporter) “ligand detection”
- IC50 for cytotoxicity measured first in HepG2
- High concentration either 100 μ M or 1/3 calculated IC50 for cytotoxicity
- Seven concentrations, 3-fold serial dilutions, 24 hr exposure
- Cells harvested, RNA isolated, processed for reporter gene quantitation
- LEC provided in data set

Attagene Technology

Cis: AhR

Multiplexed Reporter Gene Assay

Library of RTUs



Cell Transfection



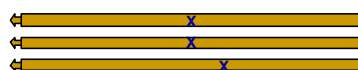
Transcription



RNA Isolation



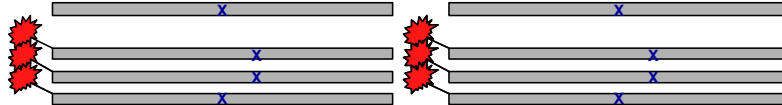
Reverse transcription



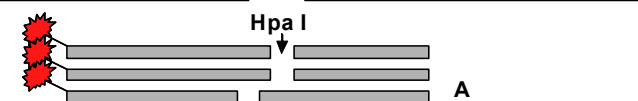
PCR amplification



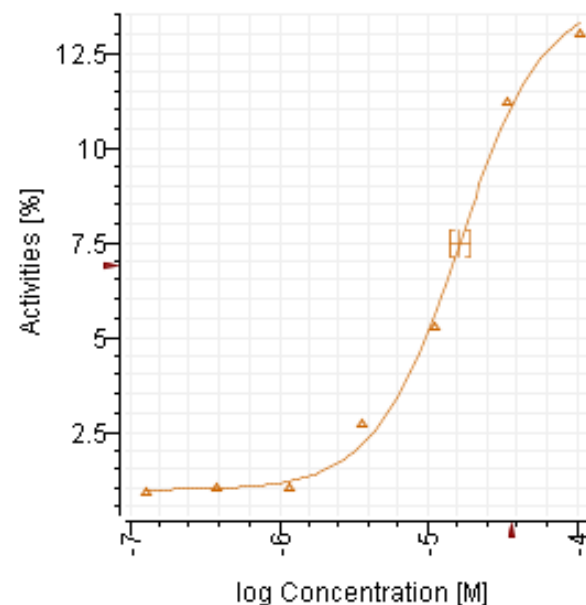
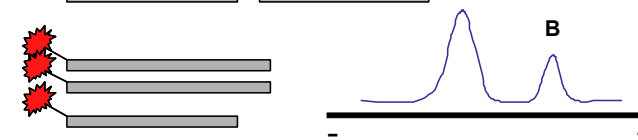
Labeling



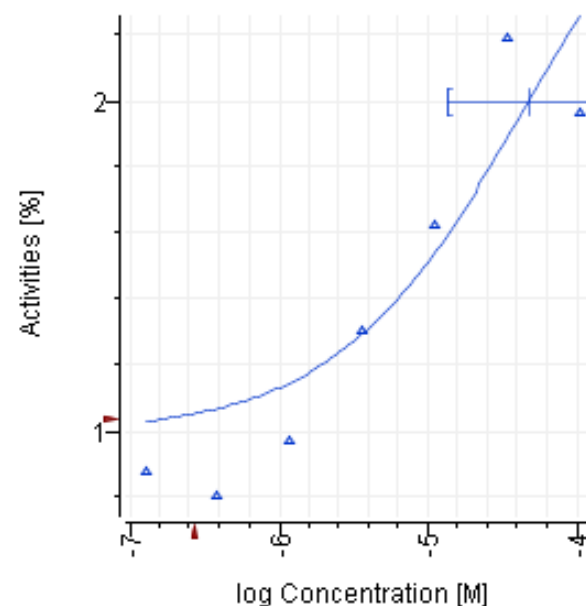
Processing (Hpa I)



Separation and detection
(capillary electrophoresis)



TV000392



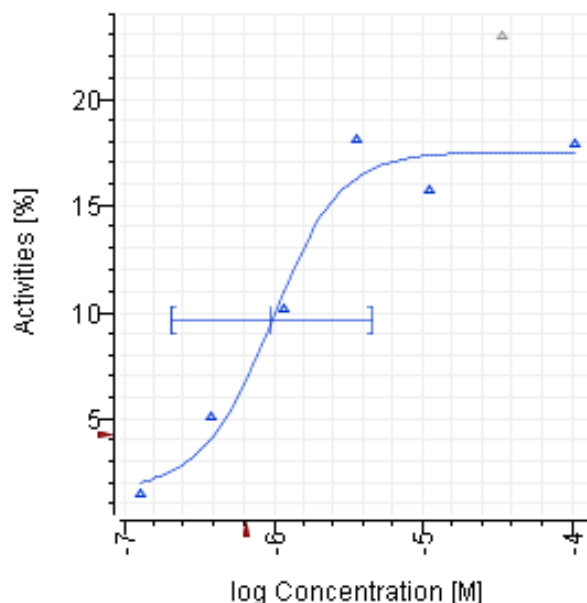
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Attagene: What Is Being Measured?

- cis Assays
 - Up/down regulation of endogenous transcription factor activity in transformed cell line
 - False positives
 - General cytotoxic response resulting in non-specific transcriptional activity
 - Promiscuity of response elements
 - Statistical, not biologically, significant response
 - Operational
 - False negatives
 - Solubility
 - Cytotoxicity
 - Operational
 - Lack of endogenous machinery
 - Lack of biotransformation
- trans Assays
 - NR agonist activity
 - False positives
 - General cytotoxic response resulting in non-specific transcriptional activity
 - Statistical, not biologically, significant response
 - Operational
 - False negatives
 - Solubility
 - Cytotoxicity
 - Operational
 - Lack of endogenous machinery
 - Lack of biotransformation

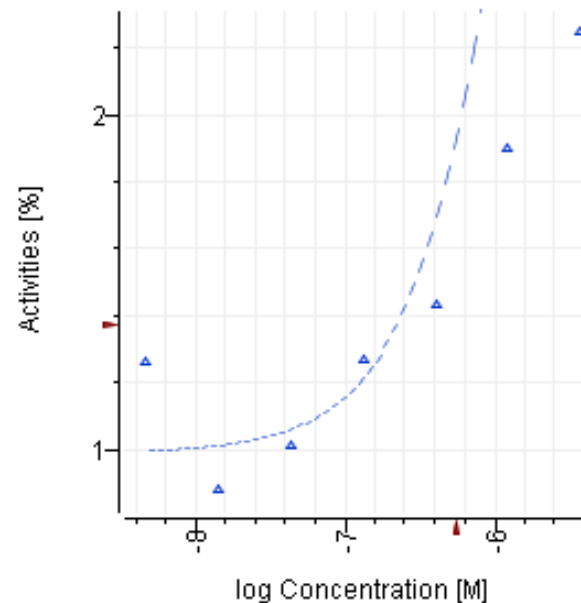
Attagene: Corresponding *cis* and *trans* Assays

Bisphenol A

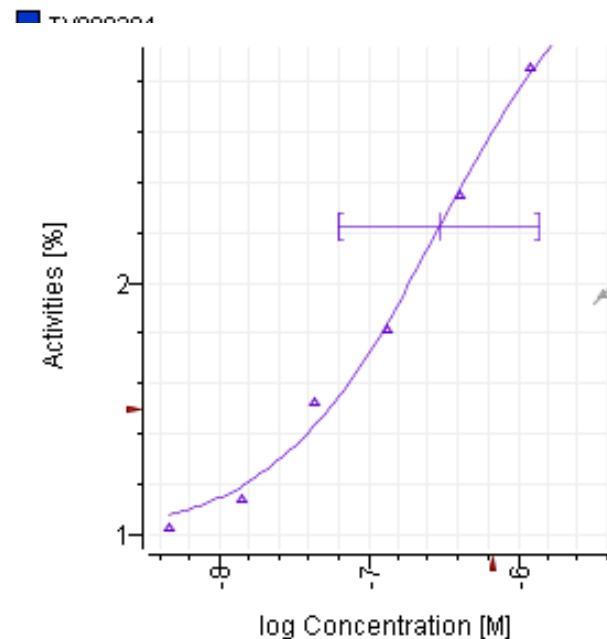
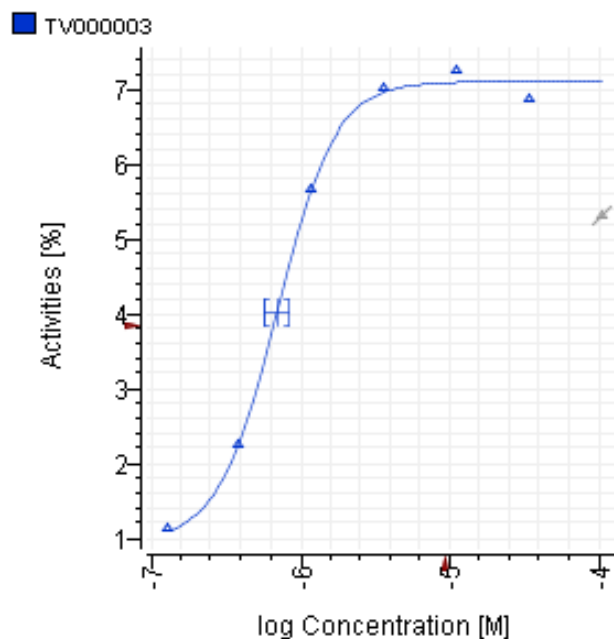


***trans*: ERa**

HPTE



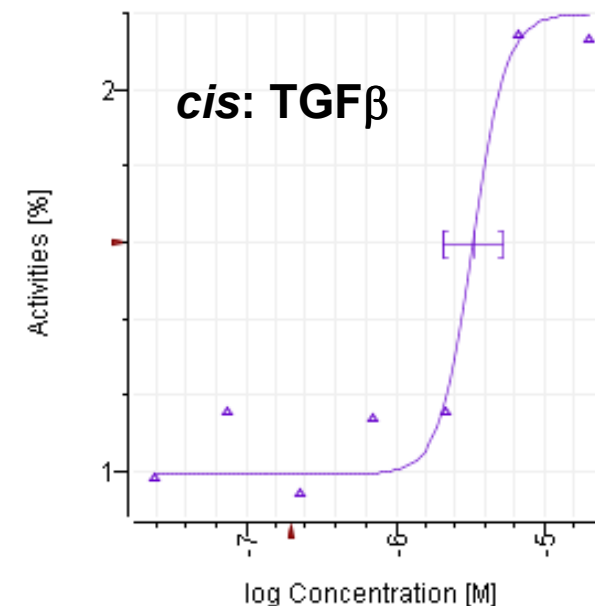
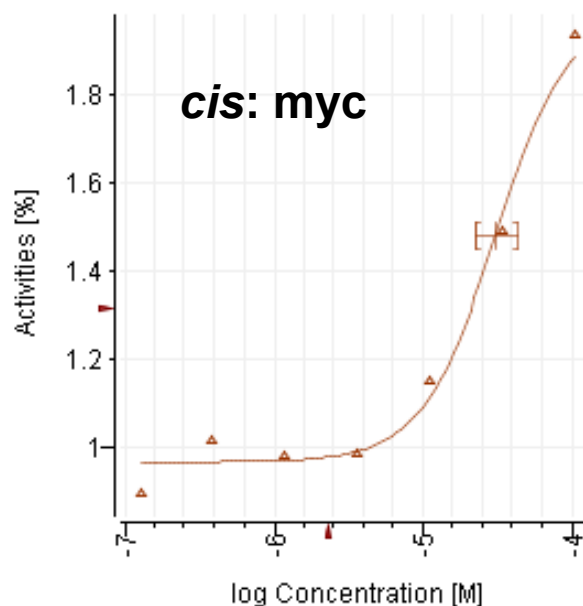
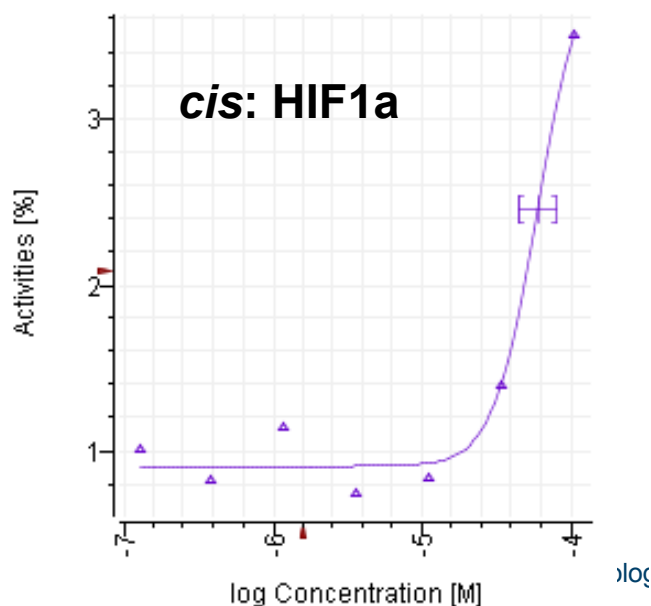
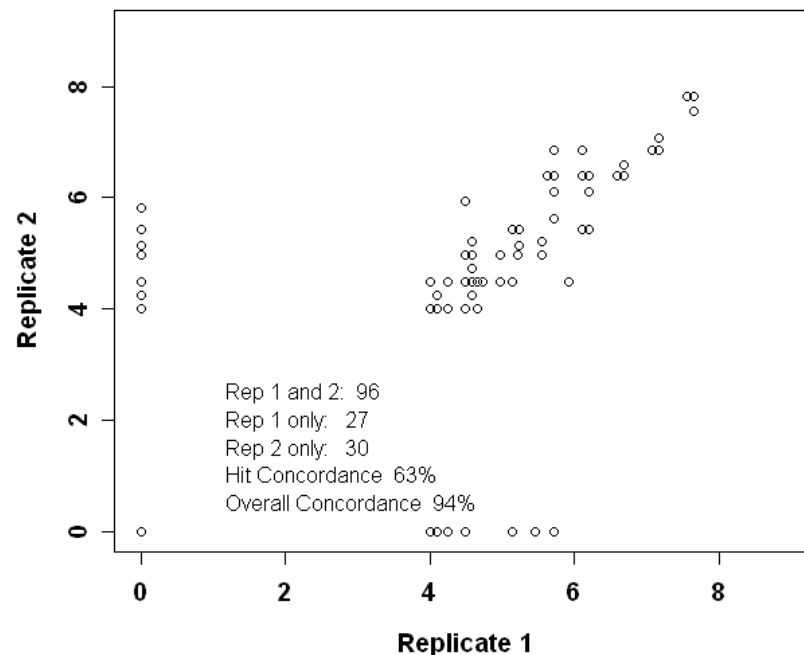
***cis*: ERE**



Attagene: Data Calculation Challenges

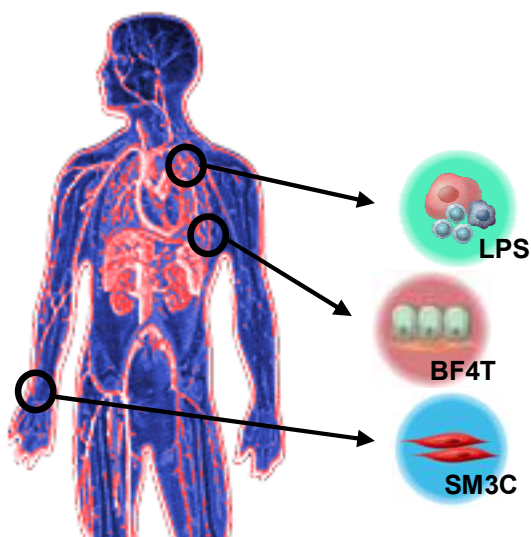
- No positive reference compound for each endpoint
- Responses, especially for *cis* assays, tended to be monotonic so potency value difficult to derive
- Biological vs statistical significance for LEC

Attagene Replicate Analysis



BioSeek: BioMAP® Technology Platform (Abstract 24)

Assays



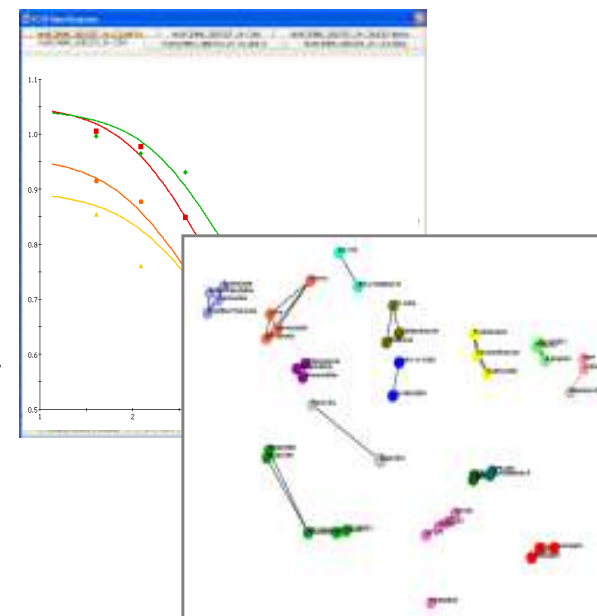
Human primary cells
Disease-like culture
conditions

Profile Database











Biological responses to
drugs and stored in the
database

Informatics

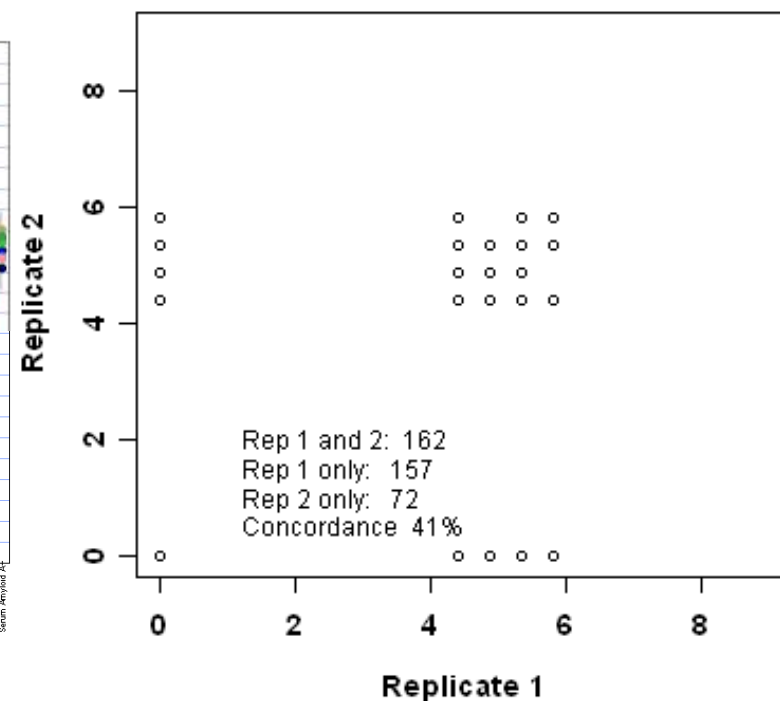


Specialized informatics tools
are used to mine and analyze
biological data

Primary Human Cell-Based Assay Platform for Human Pharmacology

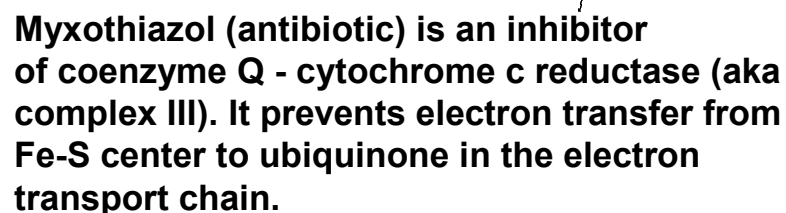
System		Cell Types	Environment	Readouts
3C		Endothelial cells	IL-1 β +TNF- α +IFN- γ	MCP-1, VCAM-1, ICAM-1, Thrombomodulin, Tissue Factor, E-selectin, uPAR, IL-8, MIG, HLA-DR, Prolif., Vis., SRB (13)
4H		Endothelial cells	IL-4+histamine	VEGFRII, P-selectin, VCAM-1, uPAR, Eotaxin-3, MCP-1, SRB (7)
LPS		Peripheral Blood Mononuclear Cells + Endothelial cells	TLR4	CD40, VCAM-1, Tissue Factor, MCP-1, E-selectin, IL-1 α , IL-8, M-CSF, TNF- α , PGE2, SRB (11)
SAg		Peripheral Blood Mononuclear Cells + Endothelial cells	TCR	MCP-1, CD38, CD40, CD69, E-selectin, IL-8, MIG, PBMC Cytotox., SRB, Proliferation (10)
BE3C		Bronchial epithelial cells	IL-1 β +TNF- α +IFN- γ	uPAR, IP-10, MIG, HLA-DR, IL-1 α , MMP-1, PAI-1, SRB, TGF- β 1, tPA, uPA (11)
HDF3CGF		Fibroblasts	IL-1 β +TNF- α +IFN- γ +bFGF+EGF+PDGF-BB	VCAM-1, IP-10, IL-8, MIG, Collagen III, M-CSF, MMP-1, PAI-1, Proliferation, TIMP-1, EGFR, SRB (12)
KF3CT		Keratinocytes + Fibroblasts	IL-1 β +TNF- α +IFN- γ +TGF- β	MCP-1, ICAM-1, IP-10, IL-1 α , MMP-9, TGF- β 1, TIMP-2, uPA, SRB (9)
SM3C		Vascular smooth muscle cells	IL-1 β +TNF- α +IFN- γ	MCP-1, VCAM-1, Thrombomodulin, Tissue Factor, IL-6, LDLR, SAA, uPAR, IL-8, MIG, HLA-DR, M-CSF, Prolif., SRB (14)

- BioSeek**



BioSeek Assays: What is Being Measured?

- Effects of chemicals on signaling pathways in primary human cells
- False positives
 - Cytotoxicity (down-regulation in particular)
 - **Statistical vs biological significance**
 - Operational
- False negatives
 - Solubility
 - Lack of biotransformation
 - Cytotoxicity
 - Operational

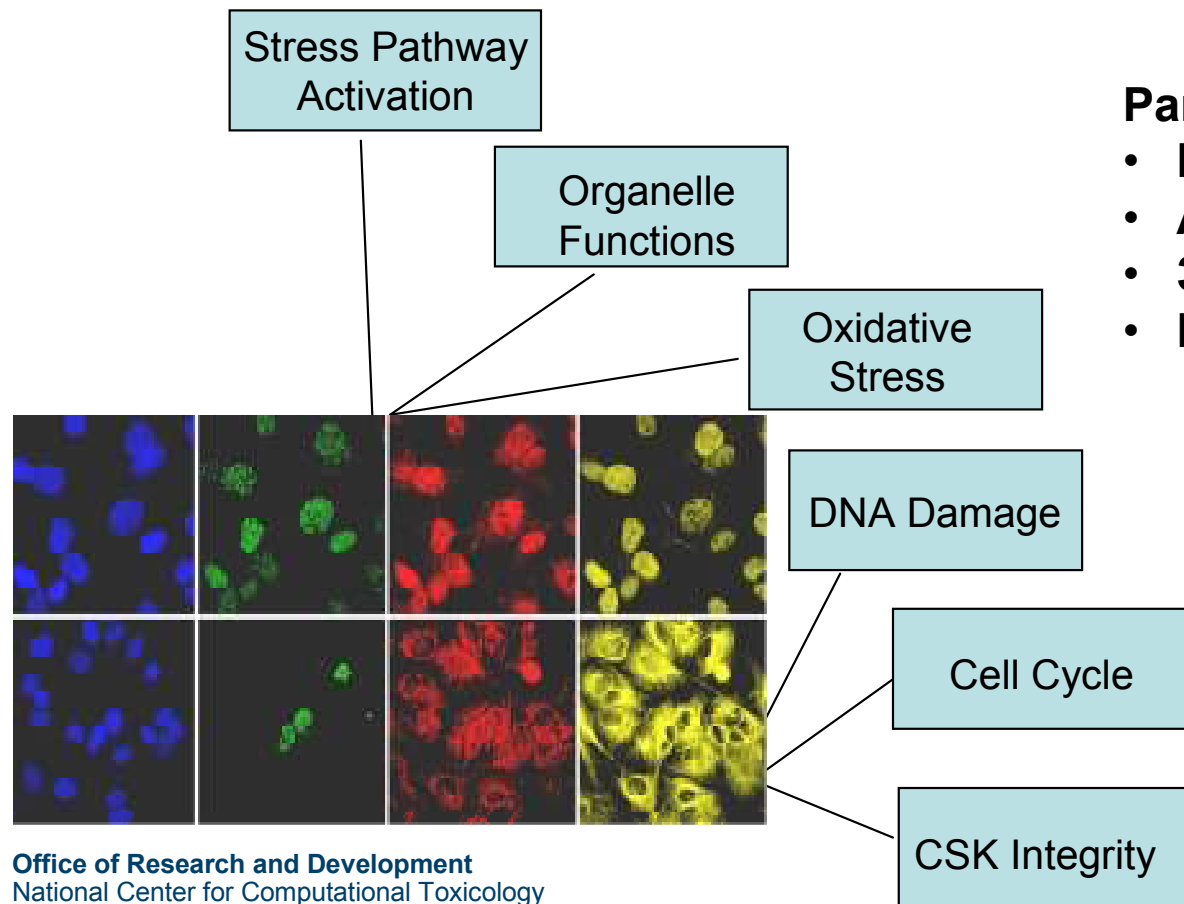


**Strobilurin fungicides: MOA--
blocks electron transfer within
respiratory chain**



Cellumen: High-Content Screening of Cellular Phenotypic Toxicity Parameters (Abstract 38)

- Technology: automated fluorescent microscopy
- Objective: Determine effects of chemicals on toxicity biomarkers in a cell culture of HepG2 and primary rat hepatocytes



Panel 1 design*:

- Multiple mechanisms of toxicity
- Acute, early & chronic exposure
- 384-well capacity
- HepG2

Cellumen: CellCiphr™ Cytotoxicity Panel

- 10-point conc-response (200 μ M-39 nM) in duplicate
- Three time points (1 hr, 24 hr, 72 hr)
- 11 endpoints per assay

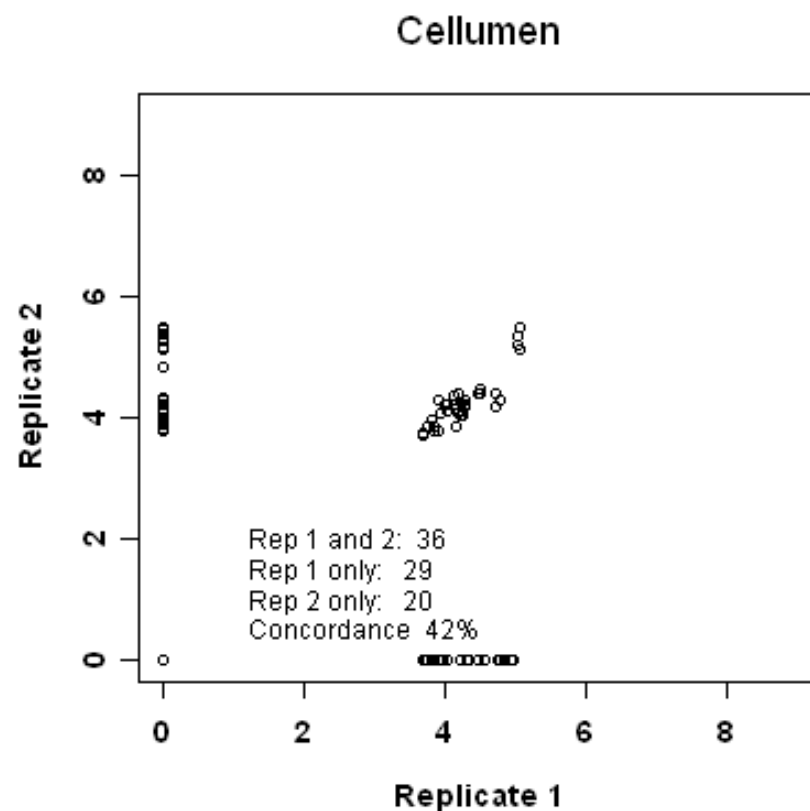
Biomarker	Measurement	Positive Control	Z'
Stress Pathway	Phospho-c-jun	Anisomycin	0.63
Oxidative Stress	Phospho-Histone H2A.X	Camptothecin	0.7
Mitochondrial Function	Mitochondrial membrane potential	CCCP	0.55
Mitochondrial Mass	Mitochondrial mass	CCCP	0.35
Cell Loss	Cell number	Camptothecin	0.56
Cell Cycle	DNA content	Paclitaxel	0.54
DNA Degradation	DNA structure	Paclitaxel	0.6
Nuclear Size	Area of nuclear region	Paclitaxel	0.63
DNA Damage	Detection of p53	Camptothecin	0.43
Mitotic Arrest	Phospho-Histone-H3	Paclitaxel	0.63
Cytoskeletal Integrity	Detection of α -tubulin	Paclitaxel	0.3

Cellumen: What is Being Measured?

- Cellular toxicity phenotypes in a transformed cell line
- False positives
 - Imaging artifacts
 - Fluorescent compounds
 - Statistical vs biological significance
 - Operational
- False negatives
 - Solubility
 - Cytotoxicity
 - Lack of biotransformation
 - Operational

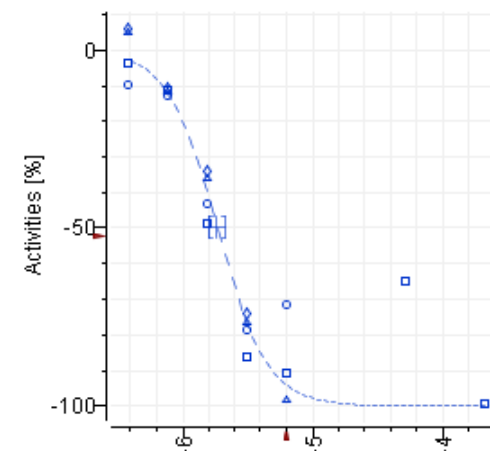
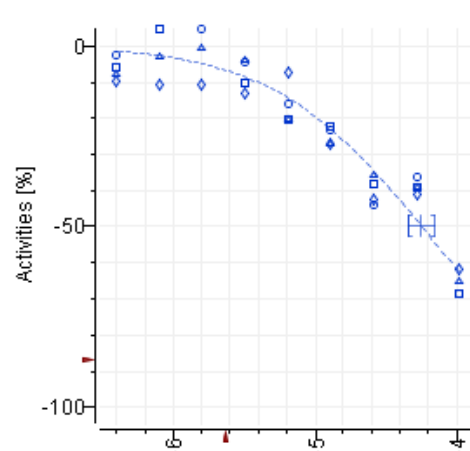
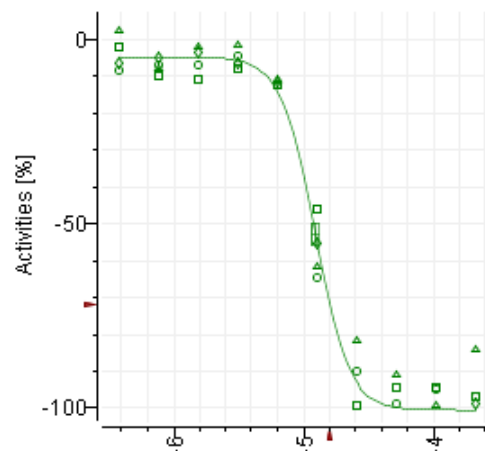
Cellumen: Data Calculation

- Data not normalized to controls (with a few exceptions)
- Fit to 3- or 4-parameter Hill equation
- AC50 reported with these rules:
 - For Cell Loss, AC50 is reported as calculated
 - For other endpoints, if AC50 for endpoint is > AC50 for Cell Loss at the corresponding exposure time, AC50 for endpoint is set to 100 μ M (to account for imaging artifacts of cytotoxicity)
- Issues:
 - Lack of positive controls for all endpoint/time combinations
 - Large differences in maximal response
 - noisy curves due in part to effects of cytotoxicity

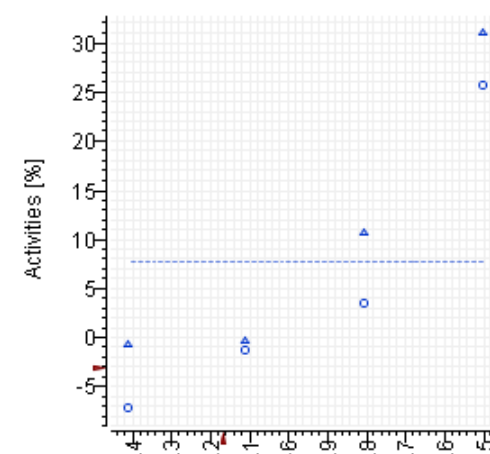
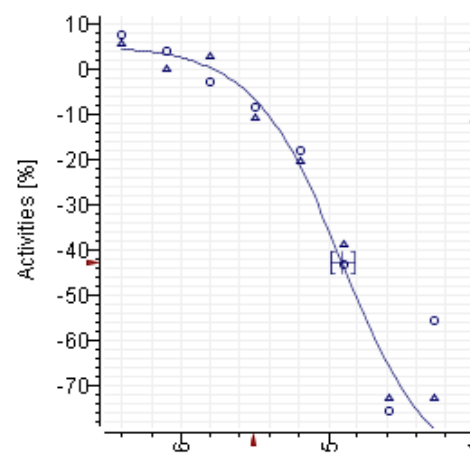
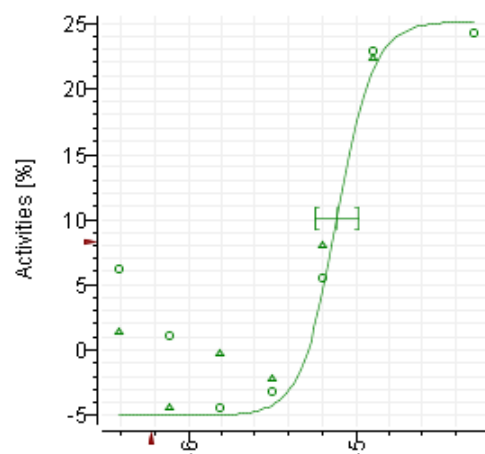


Cellumen: Data Examples

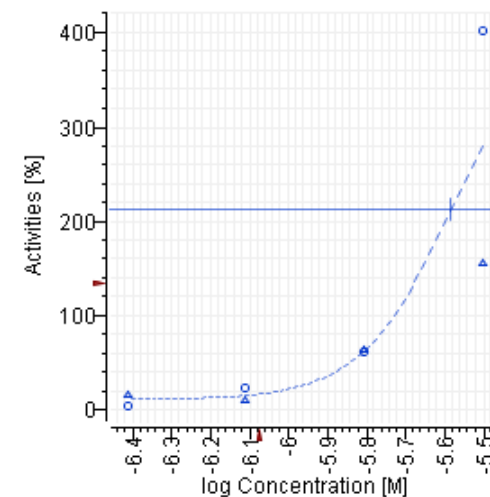
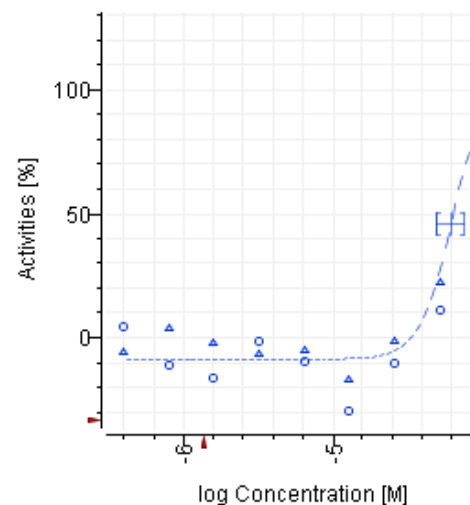
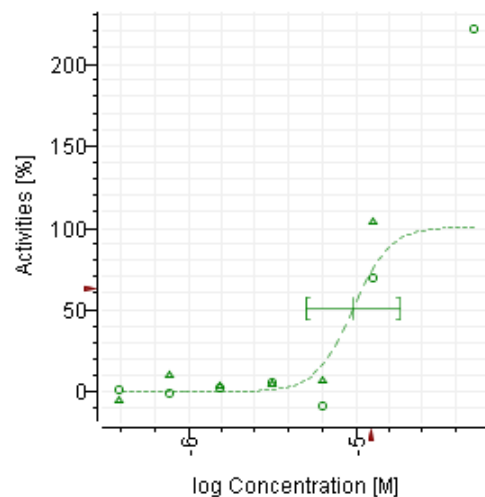
Cell Loss



Mitochondrial Membrane Potential

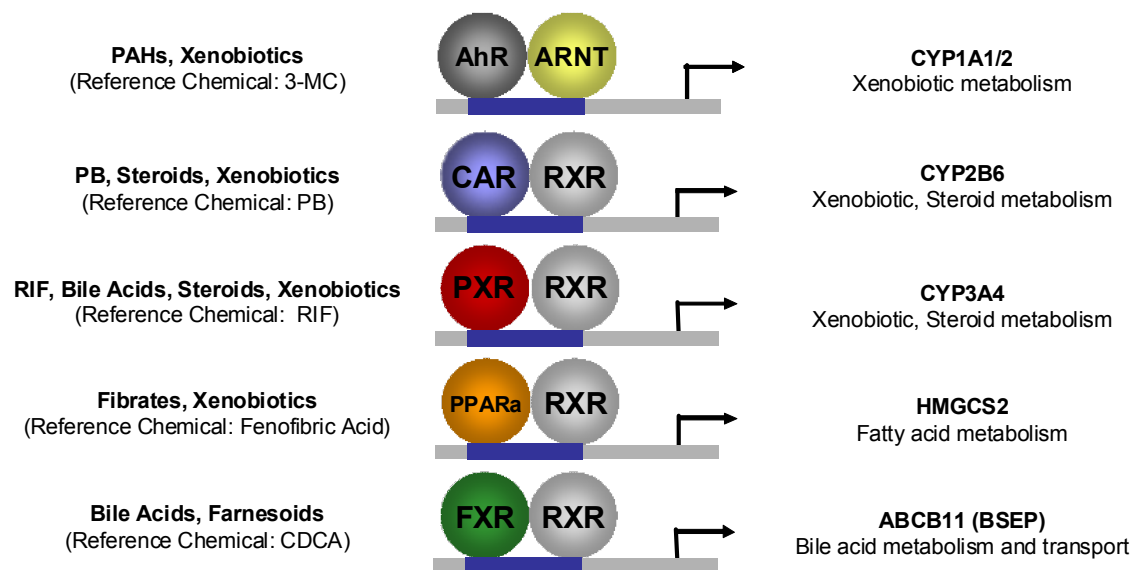


DNA Damage



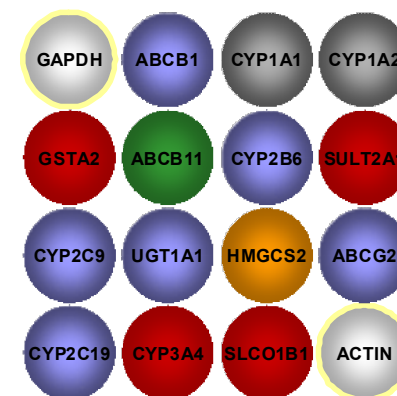
CellzDirect: XME Gene Expression in Primary Human Hepatocytes (Abstract 22)

- Primary human hepatocytes from two donors used
- Cells exposed for 6, 24, and 48 hr; medium/chemical refreshed daily
- Concentrations tested: 40, 4, 0.4, 0.04, and 0.004 μM
- 16 Genes measured in multiplexed RNase protection assay (qNPA)
- Genes targeted XME and transporters



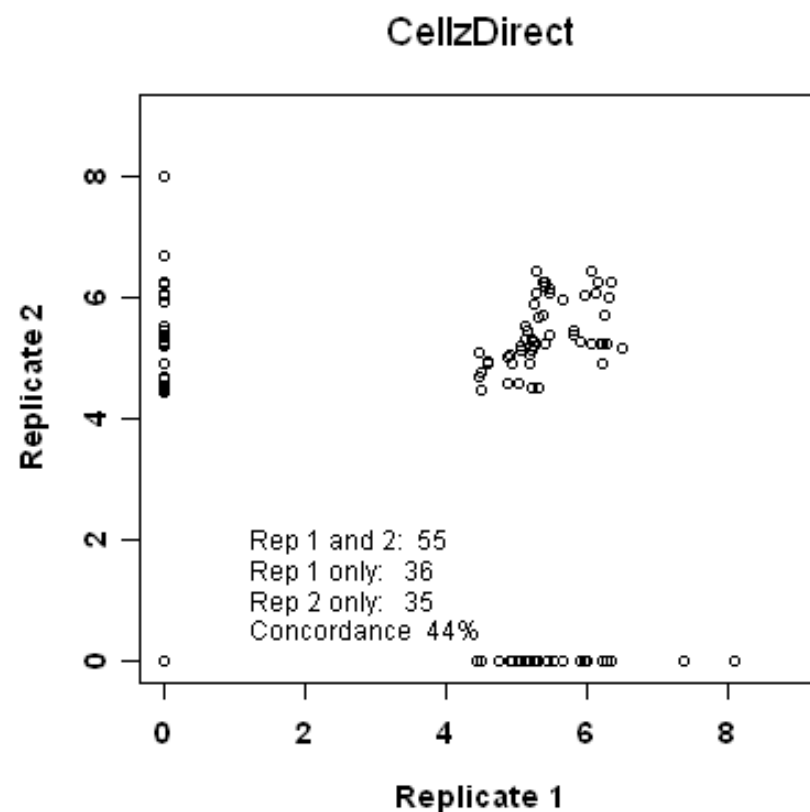
Target Gene Categories

CYP450 (6)
Transporter (4)
Phase II Metabolism (3)
Cholesterol Synthesis (1)
Endogenous control (2)



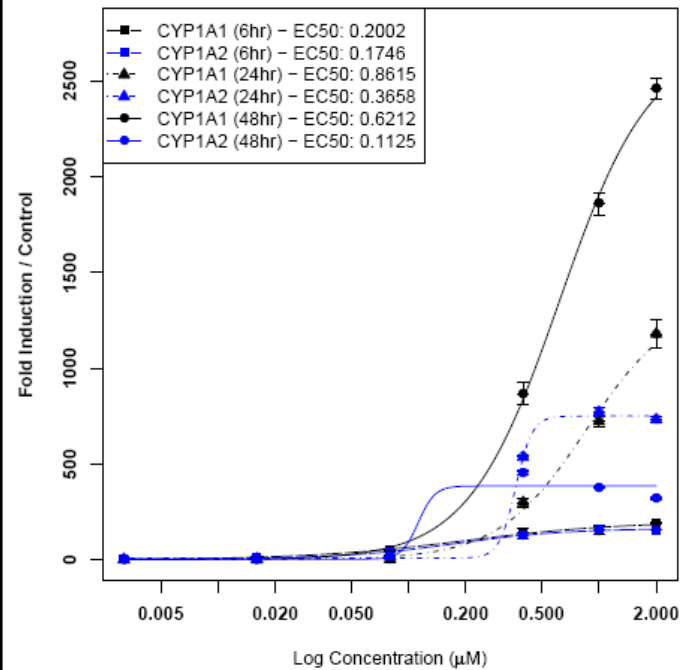
CellzDirect: Data Calculation

- Normalized to solvent control; expressed as fold-change
- Curves fit to Hill equation
 - Upper and lower limits defined by minimum and maximum responses observed over dataset of a particular gene/donor/time
- LECs determined
- Large variations in maximal responses
 - Biological vs statistical significance
- Measuring endogenous promoter activity reflects complex, multifactorial regulation of gene expression
- 6 hr exposure data not provided due to high variability associated with limited time for gene induction

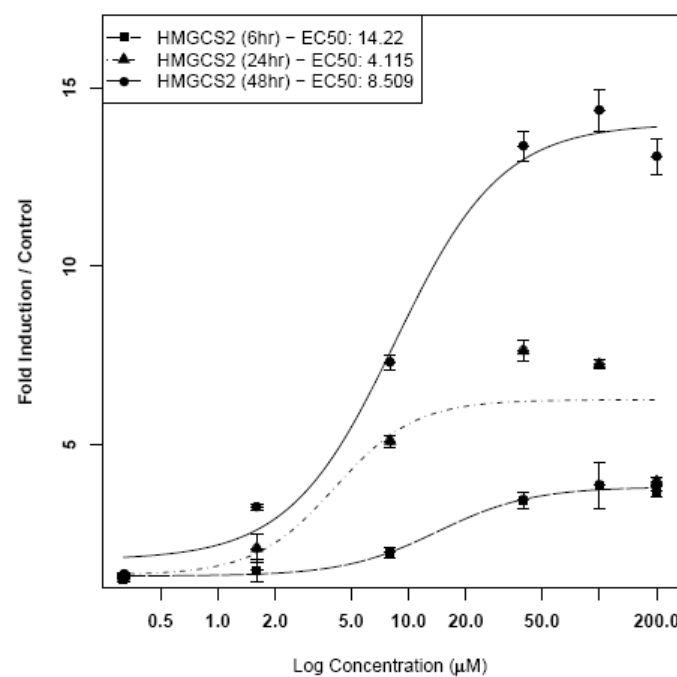


CellzDirect: Data Examples

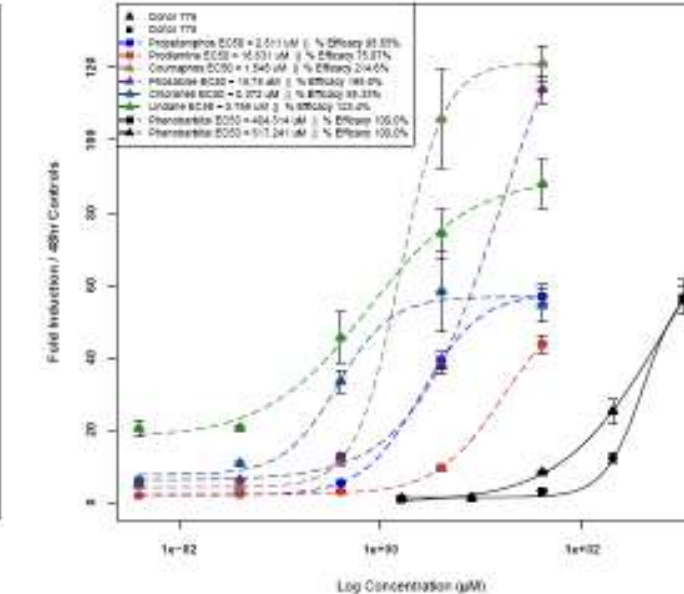
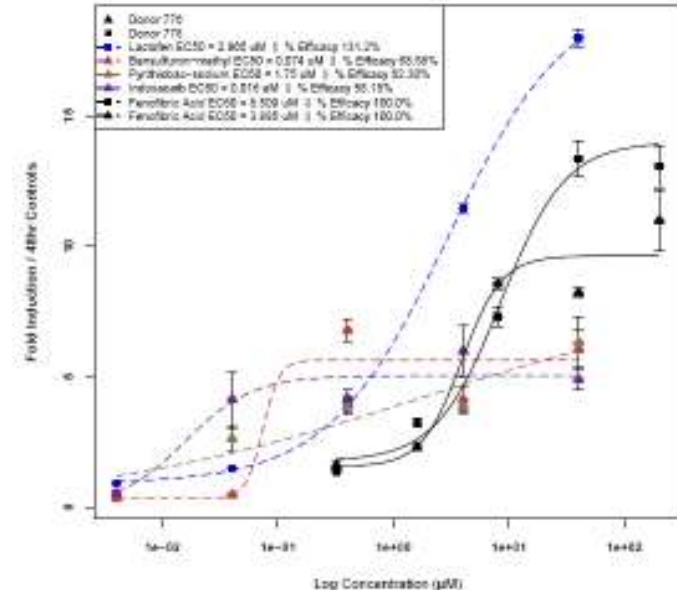
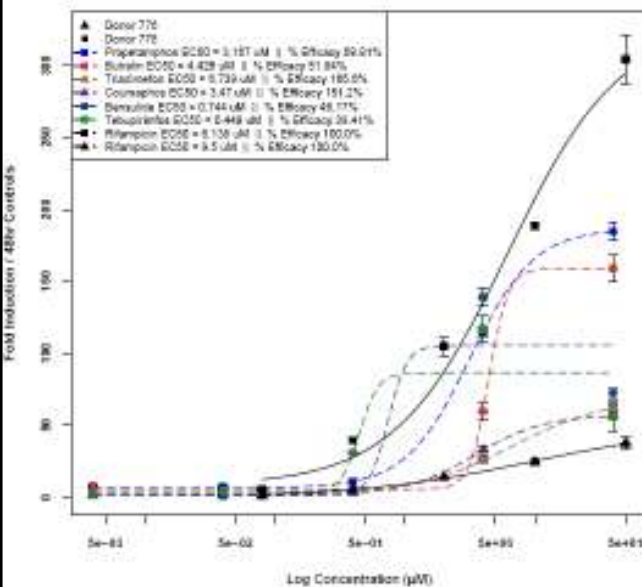
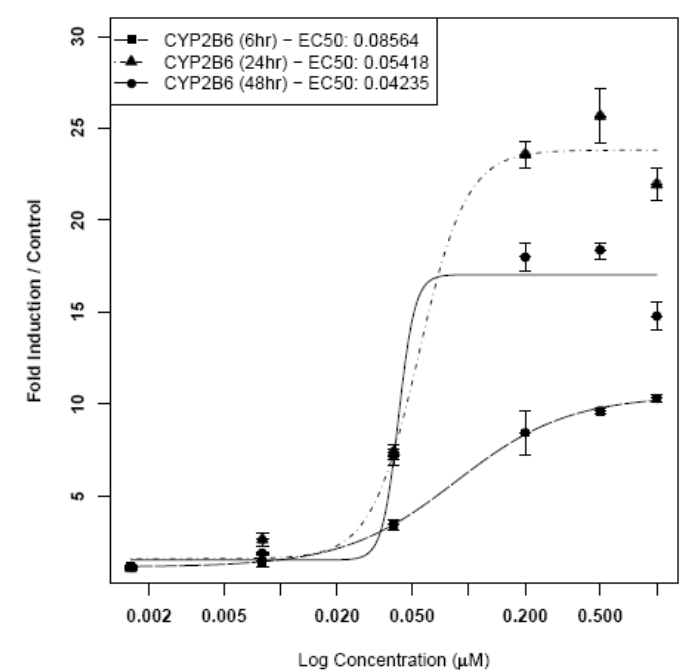
CYP1A1-AhR



HMGCS2-PPAR α



CYP2B6-CAR

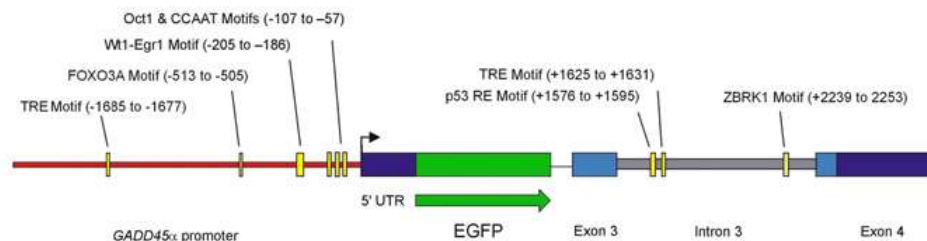
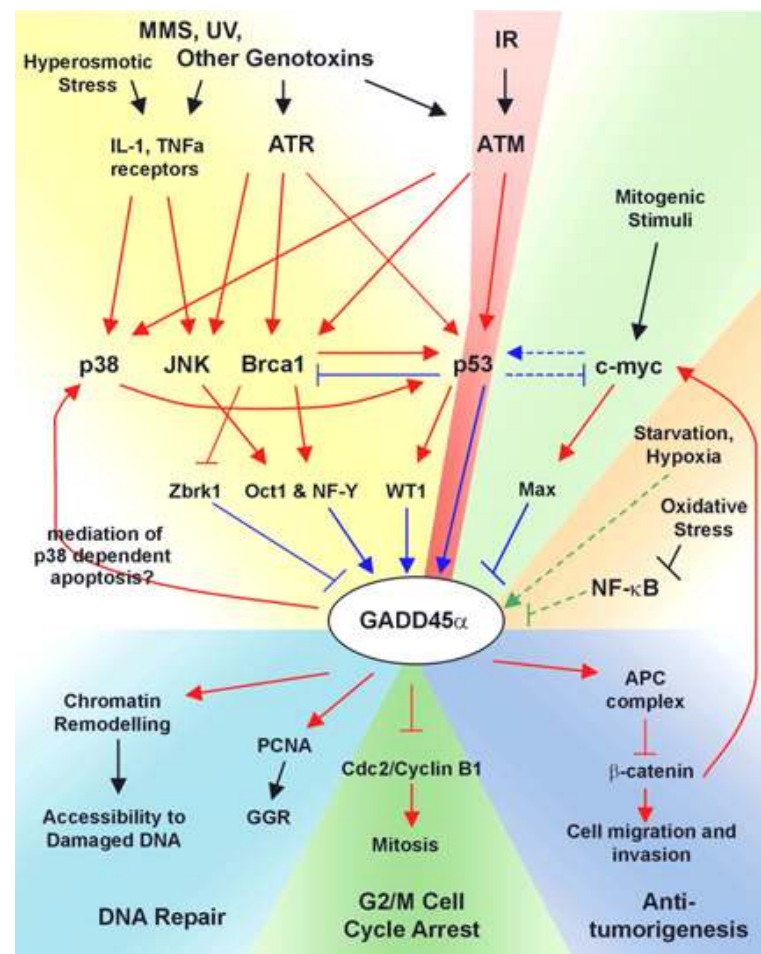


CellzDirect: What is Being Measured?

- Up/down regulation of mRNA for XME and transporters in primary human hepatocytes
- False positives:
 - General effect of cytotoxicity on transcriptional activity
 - Statistical vs biological significance
 - Operational
- False negatives:
 - Solubility
 - Cytotoxicity
 - Operational
 - Lack of biotransformation
 - Inter-individual donor variation

Gentronix: GADD45a Reporter Gene Assay for DNA Damage (Abstract 41)

- TK6 cell line expressing GFP under control of GADD45a promoter
- Cells exposed at 200, 100, 50 μ M for 24 and 48 hr
- Cytotox assay to discount artifacts
- Retested at lower conc if cytotoxic

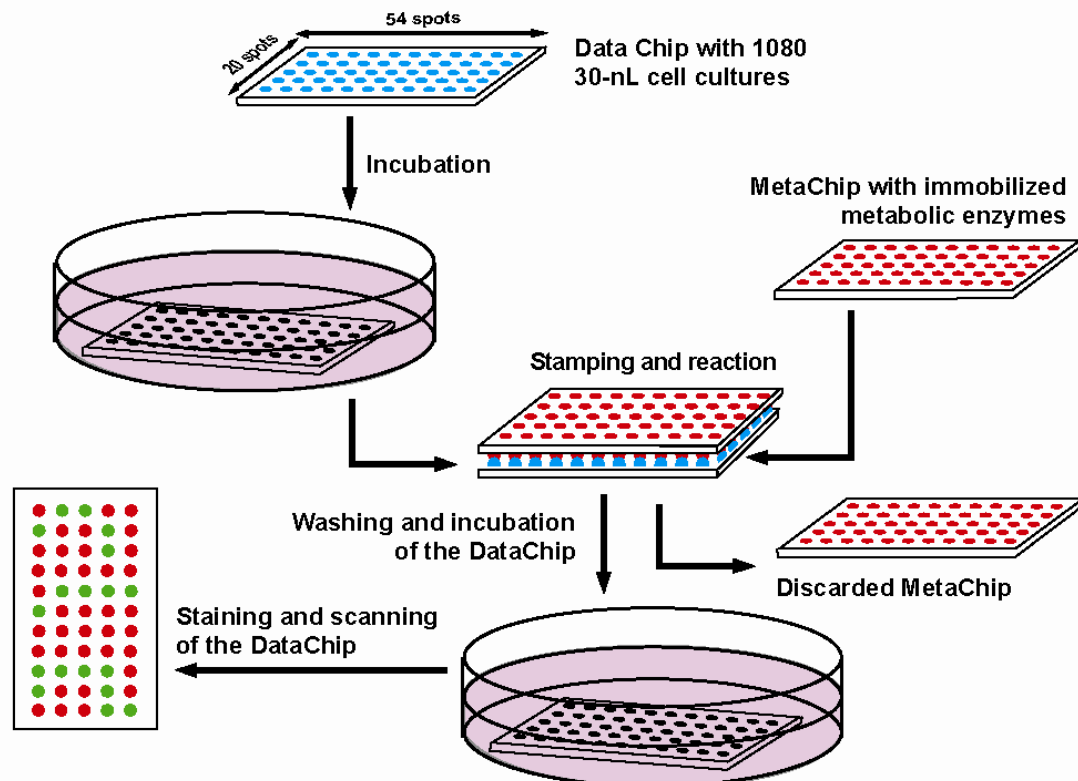


Gentronix: Data Calculations

- Induction of GFP fluorescence
>50% = genotoxic
- If over 2 or 3 concentrations,
strongly genotoxic
- LELs calculated
- Replicate analysis: no actives
among replicates
- False positives:
 - Cytotoxicity resulting in general
transcriptional activity
 - Cellular stress other than DNA
damage
 - Statistical vs biological
validation
 - Operational
- False negatives:
 - Solubility
 - Cytotoxicity
 - Lack of biotransformation
 - Operational

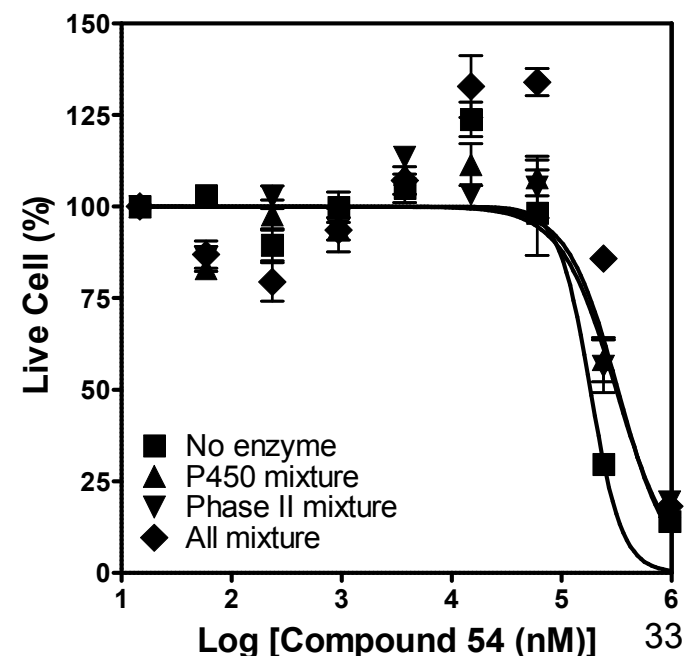
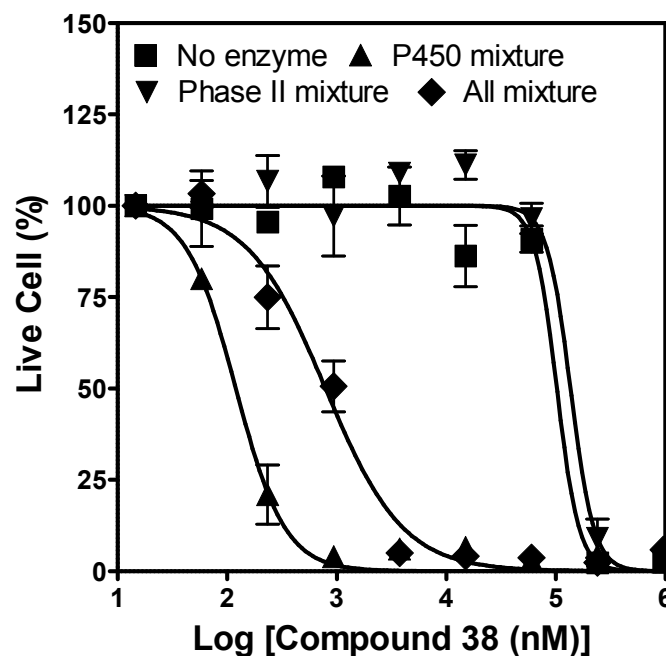
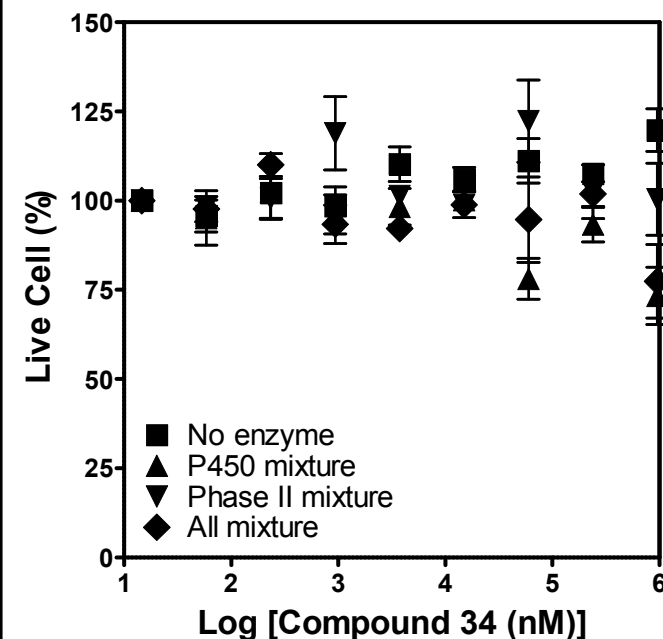
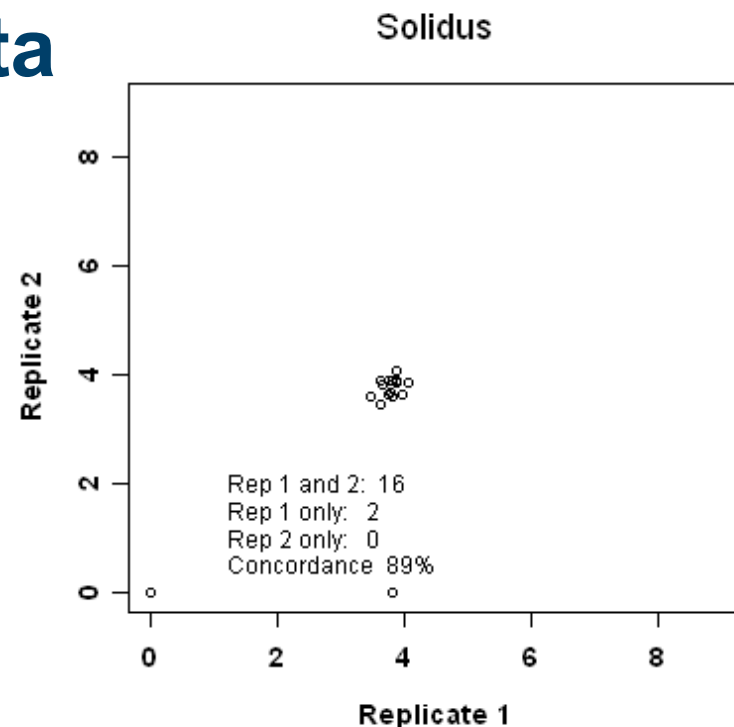
Solidus: Biotransformation Chip and Effect of Cytotoxicity (Abstract 30)

- Alginate-immobilized Phase I and Phase II enzymes
- ToxCast_320 exposed 6 hr to:
 - Control
 - Ph I
 - Ph II
 - Ph I and Ph II
- 960 μM high conc/4-fold serial dilutions/9 concentrations/5 replicates
- Cytotoxicity in Hep3B measured 48 hr later



Solidus: Example Data

- Data normalized to control values
- Concentration-response data fit to Hill equation
- LC50 determined for each assay condition



Solidus: What is Being Measured?

- Effect of Phase I and Phase II enzymes on cytotoxicity activity of chemicals against a transformed cell line
- False positives:
 - Possibly not optimized Phase I and/or Phase II mix
 - Operational
- False negatives
 - Solubility
 - Possibly not optimized Phase I and/or Phase II mix
 - Availability of compound from alginate-immobilized enzyme matrix
 - Operational

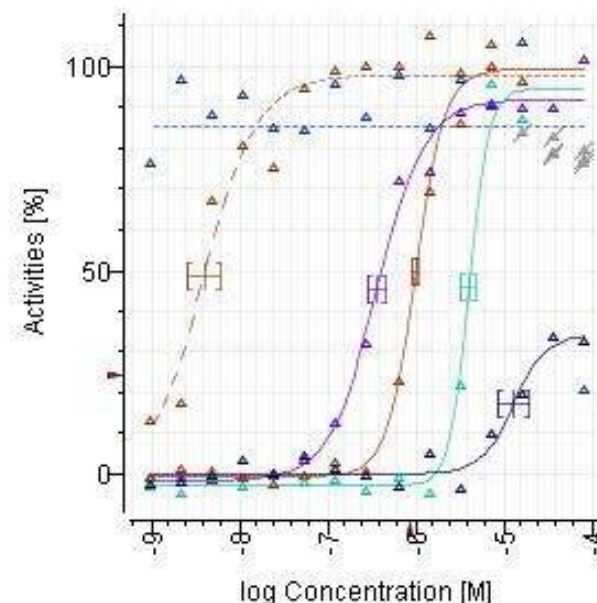
NCGC Reporter Gene Assays

- Nuclear Receptors
 - GAL4 System (ligand detection assay)
 - 11 human receptors
 - 1 rat (PXR)
 - β -lactamase reporter gene assays except:
 - PXR assays are luciferase reporter gene assays
- p53 Reporter Gene assay
 - β -lactamase reporter gene assay
- Parental cell lines mostly HEK293 (also HeLa and DPX-2)
- 12-15 point concentration-response curves (single replicate)

NCGC: Data Calculations

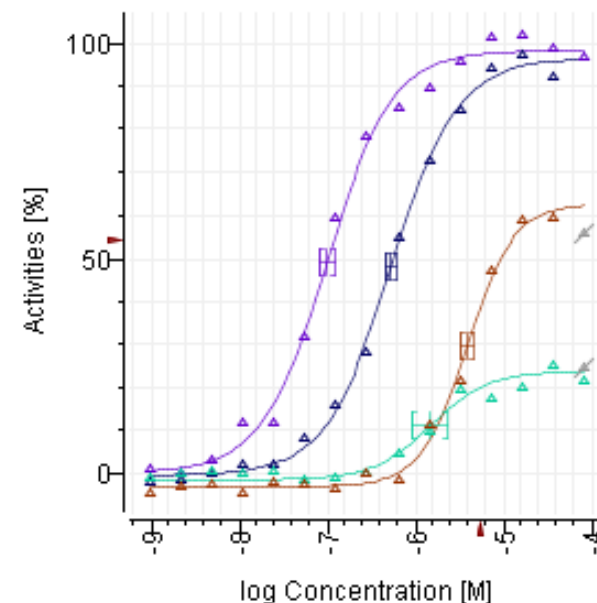
- Data normalized to reference compound effect
- Curves fit to 3- or 4-parameter Hill equation
- Artifacts removed where obvious fluorescence or cytotoxicity detected
- Required at least 25% efficacy of control compound to calculate AC50
- AC50 values provided
- Antagonist format assays challenging due to effects of cytotoxicity
- LXR assay problematic—contaminated with GR reporter line?

ER α



■ NCGC00090749-04
■ NCGC00161666-02

PPAR γ



■ NCGC00164420-01
■ NCGC00093991-03
■ NCGC00164230-01
■ NCGC00022570-07

NCGC Assays: What is Being Measured?

- NR assays are ligand-detection assays
- False positives
 - Fluorescent compounds
 - Statistical vs biological significance
 - Gal4_NR-LBD not physiological
 - Cytotoxicity (antagonist format)
 - Operational
- False negatives
 - Fluorescent compounds
 - Cytotoxicity
 - Gal4_NR-LBD not physiological
 - Operational
 - Lack of biotransformation

Additional Data Sets To Be Added Soon:

- NHEERL
 - Zebrafish developmental toxicity (Padilla) Poster
 - Neurite outgrowth and neuronal proliferation (Mundy and Shafer)
 - Stress gene reporter assays (Simmons)
 - ES cell differentiation (Hunter)
- Plasma protein binding and hepatocyte clearance (Thomas)
- PPAR α and AhR (NCGC)
- Primary rat hepatocyte HCS Cell Health (Cellumen)