

# Chemical Prioritization Methods for Nuclear Receptor Modulators at the U.S. EPA

Keith A. Houck, Ph.D.

National Center for Computational Toxicology (NCCT/ORD/EPA)

EDTA International Symposium, Dongguk University

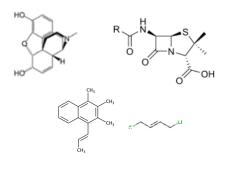
November 9, 2018

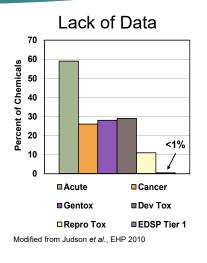
houck.keith@epa.gov

The views expressed in this presentation are those of the author and do not necessarily reflect the views or policies of the U.S. EPA Use of commercial names does not constitute endorsement of those brands U.S. Environmental Protection Agency

#### Regulatory Agencies Make a Broad Range of Decisions on Chemicals...

Number of Chemicals /Combinations

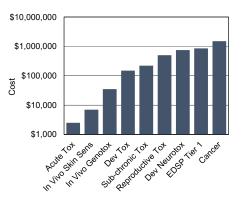




Ethics/Relevance Concerns

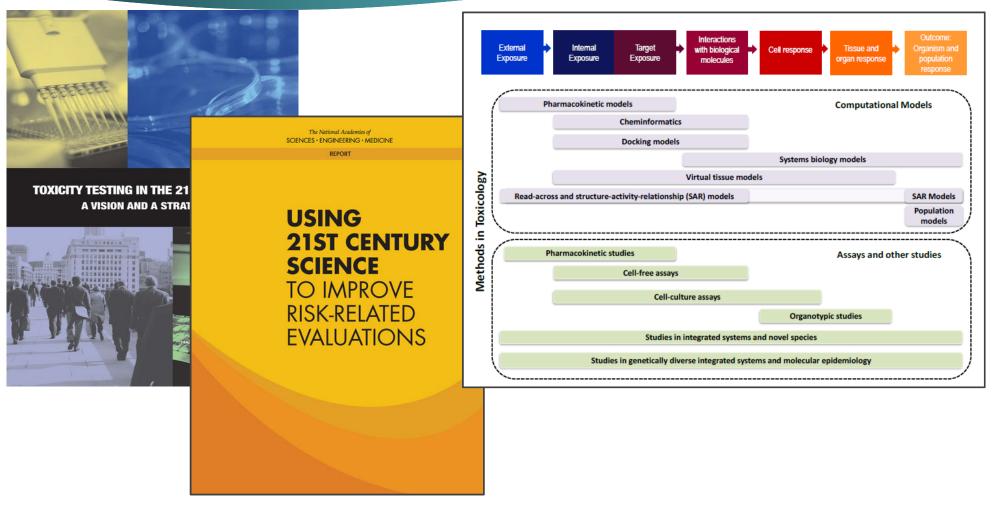


#### Economics

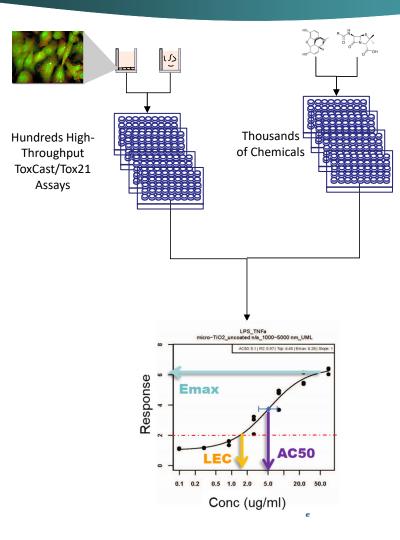


- Number of chemicals and combinations of chemicals is extremely large (>20,000 substances on active TSCA inventory)
- Due to historical regulatory requirements, most chemicals lack traditional toxicity testing data
- Traditional toxicology testing is expensive and time consuming
- Traditional animal-based testing has issues related to ethics and relevance

#### Toxicology Moving to Embrace 21<sup>st</sup> Century Methods



#### High-Throughput Assays Used to Screen Chemicals for Potential Toxicity





Interactions with biological

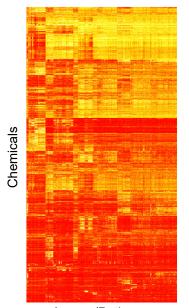
molecules

Cell response

- Understanding of what cellular processes/pathways may be perturbed by a chemical
- Understanding of what amount of a chemical causes these perturbations

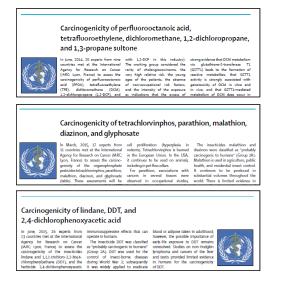
## Broad Success Derived from High-Throughput Screening Approaches

Group Chemicals by Similar Bioactivity and Predictive Modeling



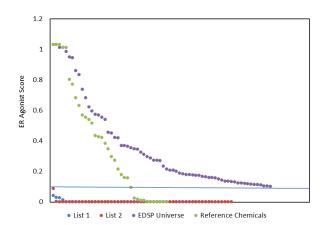
Assays/Pathways

#### Provide Mechanistic Support for Hazard ID



IARC Monographs

#### Prioritization of Chemicals for Further Testing

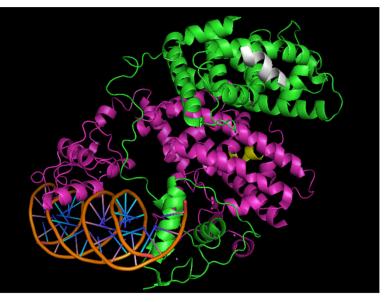


FIFRA SAP, Dec 2014

## Focus on Nuclear Receptors and Xenobiotics

- Family of ligand-regulated nuclear transcription factors (48 human)
- Conserved, modular domains
  - DNA-binding domain
  - Ligand-binding domain
    - Binds lipophilic, small molecules
    - Endogenous ligands: steroid hormones, fatty acids
- Regulates genes for key physiological processes: endocrine system, growth and differentiation, metabolism
- Endogenous ligand physicochemical properties consistent with cell permeable qualities
- Good focus for selective xenobiotic effects

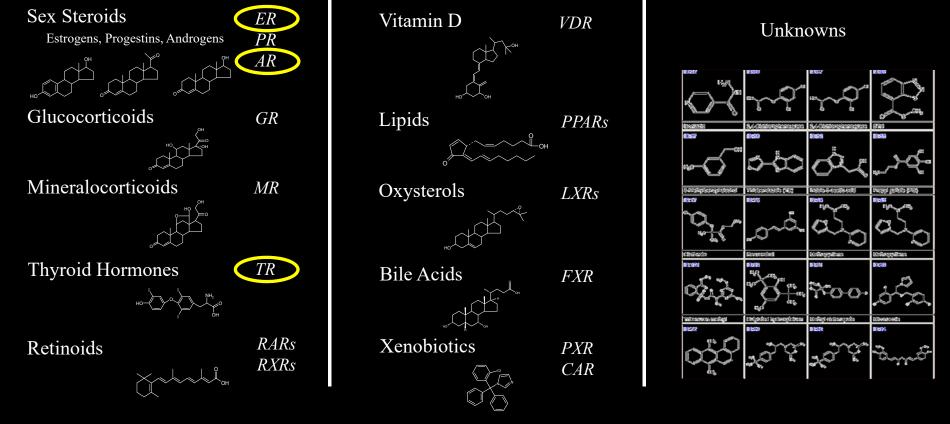




http://proteopedia.org/wiki/index.php/Image:3dzy2.png

## Ligands for Nuclear Hormone Receptors

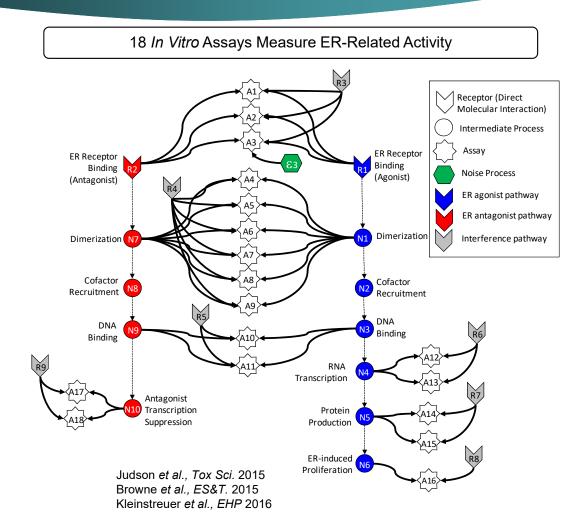
From the EPA's Endocrine Disruptor Screening Management Plan: *"Examine effects of these chemicals on estrogen, androgen and thyroid hormone-related processes"* 



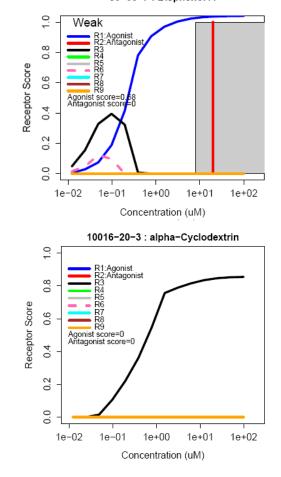
### The Estrogen Receptor Model

- Public solicitation for diverse high-throughput assays to cover broad range of bioactivity/toxicity endpoints
- Many estrogen receptor assays included
  - Binding
  - Nuclear localization
  - Transactivation
  - Cell proliferation
- No single assay perfect for a variety of reasons
- Decided to develop computational model utilizing all data

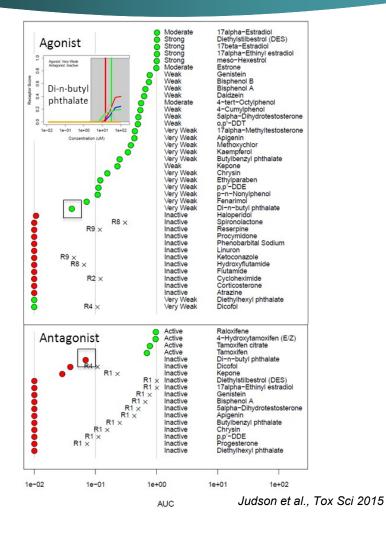
### **Targeted Pathways**

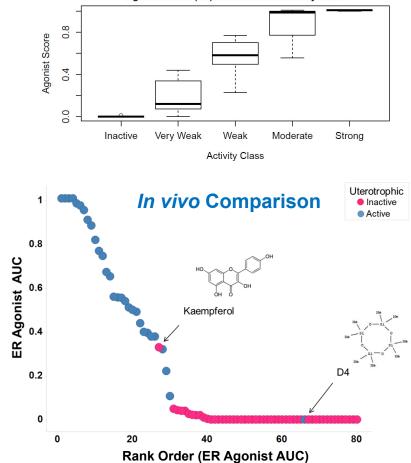






### **ER Model Performance**

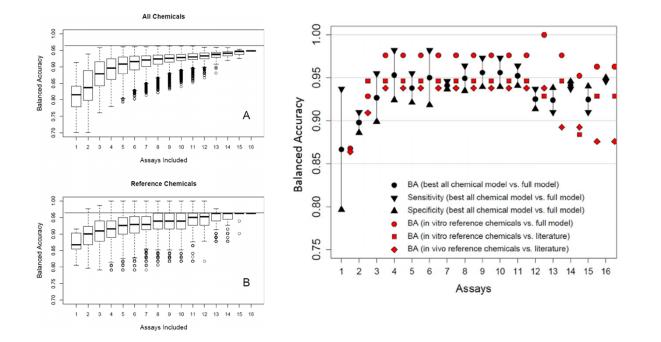




Agonist Score (R1) vs. Reference Activity Class

Browne et al., Environ. Sci. Technol., 2015

### **ER Minimal Model**



Combinations of four assays provide good balanced accuracy

R.S. Judson et al. / Regulatory Toxicology and Pharmacology 91 (2017) 39e49

## **Regulatory Applications: EDSP**



E No Sign in Sign up Search Documents

Notice

Q

Use of High Throughput Assays and Computational Tools; Endocrine Disruptor Screening Program; Notice of Availability and Opportunity for Comment

A Notice by the Environmental Protection Agency on 06/19/2015

This document has a comment period that ends in 53 days (08/18/2015)

ACTION Notice

SUMMARY

This document describes how EPA is planning to incorporate an alternative scientific approach to screen chemicals for their ability to interact with the endocrine system. This will improve the Agency's ability to fulfill its statutory mandate to screen pesticide chemicals and other substances for their ability to cause adverse effects by their interaction with the endocrine system. The approach incorporates validated high throughput assays and a computational model and, based on current research, can serve as an alternative for some of the current assays in the Endocrine Disruptor Screening Program (EDSP) Tier ( battery, EPA has partial screening results for over 1800 chemicals that have been evaluated using high throughput assays and a computational model for the estrogen receptor pathway. In the future, EPA anticipates that additional alternative methods will be available for EDSP chemical screening based on further advancements of high throughput assays and computational models for other endocrine pathways. Use of these alternative methods will accelerate the pace of screening, decrease costs, and reduce animal testing. In addition, this approach advances the goal of providing sensitive, specific, quantitative, and efficient screening using alternative test methods to some assays in the Tier 1 battery to protect human health and the environment.

TABLE OF CONTENTS • DATES: Back to Top • ADDRESSES:

• FOR FURTHER INFORMATION CONTACT:

Bade (Presious Document LEGAL DISCLAIMER Font Controls + - A PDF DEV PUBLIC INSPECTION Publication Date: Friday, June 19, 2015 Agency: Environmental Protection Agency Dates: Comments must be received on or before August 18, 2015. **Comments** Close 08/18/2015 Entry Type: Notice Action: Notice. Document Citation 80 FR 35350 Page 35350 -35355 (6 pages) Agency/Docket Numbers: EPA-HQ-OPPT-2015-0305 FRL-9928-69

> Document Number 2015-15182

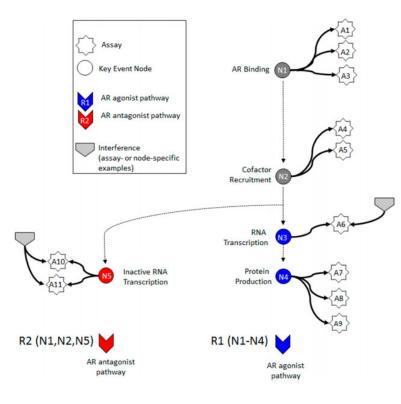
"The approach incorporates validated high-throughput assays and a computational model and, based on current research, can serve as an alternative for some of the current assays in the Endocrine Disruptor Screening Program (EDSP) Tier 1 battery."

#### Androgen Receptor Screening

- Utilize existing ToxCast/Tox21 assays to develop AR model
- Cytotoxic chemicals confounded antagonist cell-based assays
- Run additional confirmation assay for antagonists
  - Higher agonist concentration
  - Competitive antagonists show right-shift in potency

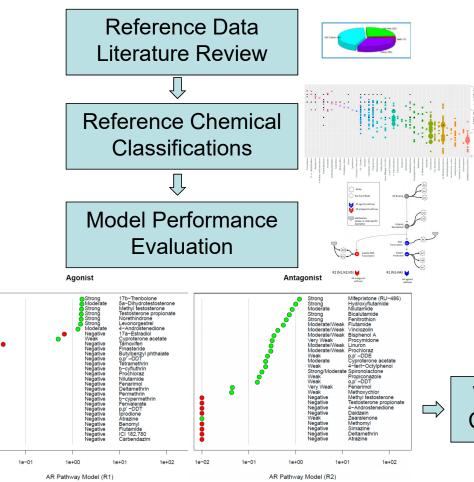
#### ToxCast/Tox21 Assays

ID	node	assay name	source	gene"	species	type	associated pathways <sup>b</sup>
A1	N1	NVS_NR_hAR	Novascreen	AR	Homo sapiens	receptor binding	R1; R2; R3
A2	N1	NVS_NR_cAR	Novascreen	AR	P. troglodytes	receptor binding	R1; R2; R3
A3	N1	NVS_NR_rAR	Novascreen	AR	Rattus norvegicus	receptor binding	R1; R2; R3
A4	N2	OT_AR_ARSRC1_0480	Odyssey Thera	AR; SRC	Homo sapiens	coregulator recruitment	R1; R2; R4
A5	N2	OT_AR_ARSRC1_0960	Odyssey Thera	AR; SRC	Homo sapiens	coregulator recruitment	R1; R2; R4
A6	N3	ATG_AR_TRANS	Attagene	AR	Homo sapiens	RNA reporter gene	R1; R5
A7	N4	OT_AR_ARELUC_AG_1440	Odyssey Thera	AR; ARE	Homo sapiens	reporter gene	R1; R6
A8	N4	Tox21_AR_BLA_Agonist_ratio	NCATS/ NCGC	AR	Homo sapi <b>e</b> ns	reporter gene	R1; R6
A9	N4	Tox21_AR_LUC_MDAKB2_Agonist	NCATS/ NCGC	AR	Homo sapiens	reporter gene	R1; R6
A10	N5	Tox21_AR_BLA_Antagonist_ratio	NCATS/ NCGC	AR	Homo sapiens	reporter gene	R2; R7
A11	N5	Tox21_AR_LUC_MDAKB2_Antagonist	NCATS/ NCGC	AR	Homo sapiens	reporter gene	R2; R7
A11'	N5	Tox21_AR_LUC_MDAKB2_Antagonist- confirmation	NCATS/ NCGC	AR	Homo sapiens	reporter gene	NA



Antagonist Mode

#### **Evaluation of AR Model**



1e-02

#### Summary

- Model has high sensitivity
- Antagonist mode specificity improved by considering antagonist assay with high agonist concentration
- Weakly active chemicals most difficult to detect
- Broad screening suggested cytotoxic compounds not all excluded

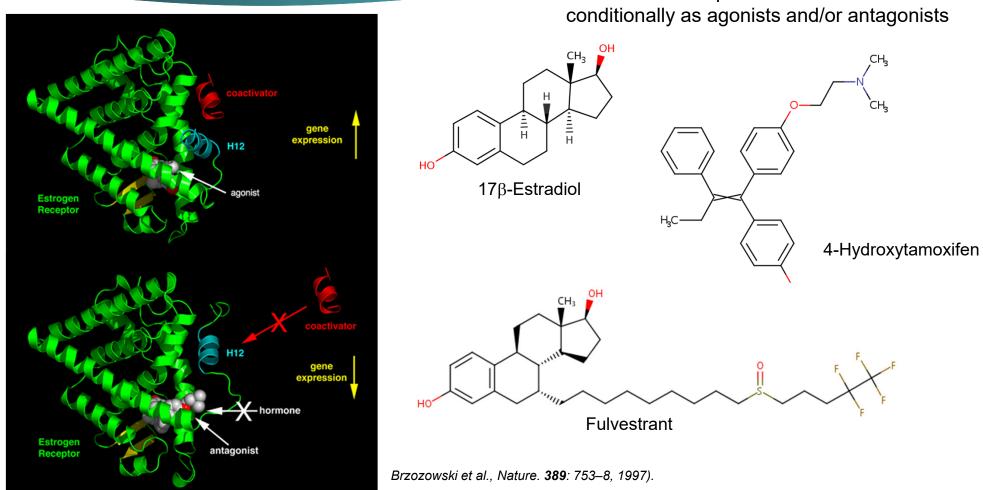
Validated Model for Chemical Screening

#### Chemical Research in Toxicology

Development and Validation of a Computational Model for Androgen Receptor Activity

Nicole C. Kleinstreuer,\*<sup>1</sup><sup>©</sup> Patricia Ceger,<sup>‡</sup> Eric D. Watt,<sup>§</sup><sup>©</sup> Matthew Martin,<sup>§</sup> Keith Houck,<sup>§</sup> Patience Browne,<sup>II</sup> Russell S. Thomas,<sup>§</sup> Warren M. Casey,<sup>†</sup> David J. Dix,<sup>⊥</sup> David Allen,<sup>‡</sup> Srilatha Sakamuru,<sup>#</sup> Menghang Xia,<sup>#</sup> Ruili Huang,<sup>#</sup> and Richard Judson<sup>§</sup> Chem Res Toxicol. 30:946-964, 2017.

### Agonists versus Antagonist



Selective receptor modulators behave

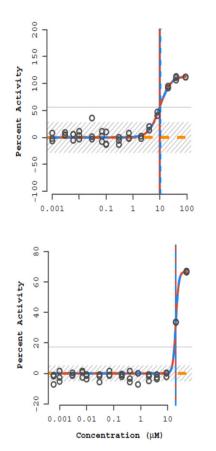
#### **Tox21 AR Screening Results**

 $\sim 8,000$  chemicals

- Only 102 chemicals positive using strictest criteria
- Expanding criteria allows for ranking of chemicals based on strength of evidence
- Chemicals that are confounded by cytotoxicity are not eliminated but evidence is weaker
- Potency not currently considered but is another important factor

# Challenges with assessing NR antagonism *in vitro*

- Measuring loss of signalconfounded by cytotoxicity
- To address:
  - Two different assay platforms
  - Use bootstrapping techniques to determine effect of cytotoxicity
  - Two concentrations of agonist R1881
  - MARCoNI assay for corepressor/activator recruitment



ASSAY: AEID1816 (TOX21\_AR\_LUC\_MDAKB2\_Antagonist2)

 NAME:
 Benzethonium chloride

 CHID:
 23810
 CASRN: 121-54-0

 SPID(S):
 Tox21\_202488

 M4ID:
 4754104

HILL MODEL (in red): tp ga gw val: 114 0.996 2.14 sd: 2.71 0.0262 0.208 GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	117	1.01	2.04	2.13	8.85
sd:	NaN	NaN	NaN	NaN	NaN
	CNST	HIL	6	GNLS	
AIC:	441.3	291	.95	295.54	
PROB:	0	0.8	6	0.14	
RMSE:	50.36	8.1	1	8.1	

MAX MEAN: 112 MAX MED: 112 BMAD: 9.44 ASSAY: AEID1017 (TOX21\_AR\_LUC\_MDAKB2\_Antagonist2\_viab:

 NAME:
 Benzethonium chloride

 CHID:
 23810
 CASRN:
 121-54-0

 SPID(S):
 Tox21\_110403\_1

 M4ID:
 10861077

HILL MODEL (in red): tp ga gw val: 66.8 1.3 8 sd: 1.03 0.00424 4.68

GAIN-LOSS MODEL (in blue): lw 1a val: 66.8 1.3 3.49 6.26 sd: NA NA NA NA NA CNST HILL GNLS AIC: 345.27 225.7 229.7 PROB: 0 0.12 0.88 RMSE: 19.49 2.82 2.82 MAX MEAN: 66.7 MAX MED: 67 BMAD: 1.74

COFF: 17.4 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS :

## Antagonist Reference Chemical Results

#### Antagonist Screening

- LUC: R1881 = 0.5 nM
- LUC\_counterscreen: R1881 = 10 nM

Criteria	Sensitivity	Specificity	Balanced Accuracy
Active in BLA	0.9	0.88	0.89
Active in LUC_counterscreen	0.8	1	0.9
Active in LUC	0.9	0.88	0.89
Active in at least 2 assays	0.9	1	0.95
Active in all three assays	0.7	1	0.85
Active in all three assays, LUC vs.			
LUCcs difference, not confounded	!	[ ]	i
by cytotoxicity	0.5	1	0.75
Active in LUC, LUC vs. LUCcs			
difference, not confounded by		1 1	i
cytotoxicity	0.7	1	0.85

			LUC vs	LUC vs
Chemical	Designation	Assay Hitcalls	LUC_counterscreen	LUC_viability
Procymidone	Very Weak Antagonist	BLA, LUC, LUCcs	Yes	Yes
Fenarimol	Very Weak Antagonist	BLA, LUC, LUCcs	Yes	Yes
4-(1,1,3,3-				
Tetramethylbutyl)phenol	Weak Antagonist	LUC	Yes	Yes
o,p'-DDT		BLA, LUC	Yes	Yes
p,p'-DDE		LUC	Yes	Yes
Propiconazole		BLA, LUC, LUCcs	Yes	No
Zearalenone		BLA, LUC, LUCcs	No	No
Methoxychlor	Weak Antagonist	BLA, LUC, LUC2	No	No
Linuron	Moderate/Weak Antagonist	BLA, LUC	Yes	No
Vinclozolin	Moderate/Weak Antagonist	BLA, LUC, LUCcs	Yes	Yes
Flutamide	Moderate/Weak Antagonist	BLA, LUC, LUCcs	Yes	Yes
Bisphenol A	Moderate/Weak Antagonist	BLA, LUC, LUCcs	Yes	Yes
Prochloraz	Moderate/Weak Antagonist	BLA, LUC, LUCcs	Yes	Yes
Cyproterone acetate	Moderate Antagonist	BLA, LUC	Yes	Yes
Nilutamide	Moderate Antagonist	BLA, LUC, LUCcs	Yes	Yes
Spironolactone	Strong/Moderate Antagonist	BLA, LUC	No	Yes
Mifepristone	Strong/Moderate Antagonist	BLA, LUC, LUCcs	No	Yes
Fenitrothion	Strong Antagonist	BLA, LUC, LUCcs	Yes	Yes
Hydroxyflutamide	Strong Antagonist	BLA, LUC, LUCcs	Yes	Yes
Bicalutamide	Strong Antagonist	BLA, LUC, LUCcs	Yes	Yes
17-Methyltestosterone		NA	NA	NA
4-Androstene-3,17-dione		NA	NA	No
Atrazine		NA	NA	NA
Daidzein		BLA	NA	NA
Deltamethrin		NA	NA	NA
Methomyl	Negative Antagonist	LUCcs	NA	No
Simazine		NA	NA	NA

### MARCoNI assay

Microarray Assay for Real-time Coregulator-Nuclear receptor

Interaction

- Cell-free assay measuring co-regulator recruitment to AR-LBD
  - 154 co-regulators
  - 3 concentrations (1, 10, 100 uM)
  - log fold-change of binding compared to DMSO
- Tested 318 suspected AR antagonists
- Reduced variables (co-regulators) to 28 most affected
- Goal: to see if patterns of coregulatory recruitment can distinguish between true antagonists and false antagonists (cytotoxicity/artifacts)

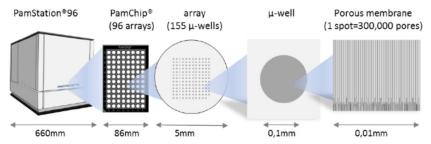
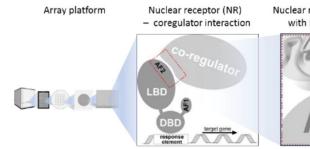


Figure 1: Outline of PamStation®96 instrument and PamChip®96 microarray, each array consisting of 155 μ-wells, (The PamChip®96 is therefore a 14880 μ-well plate)



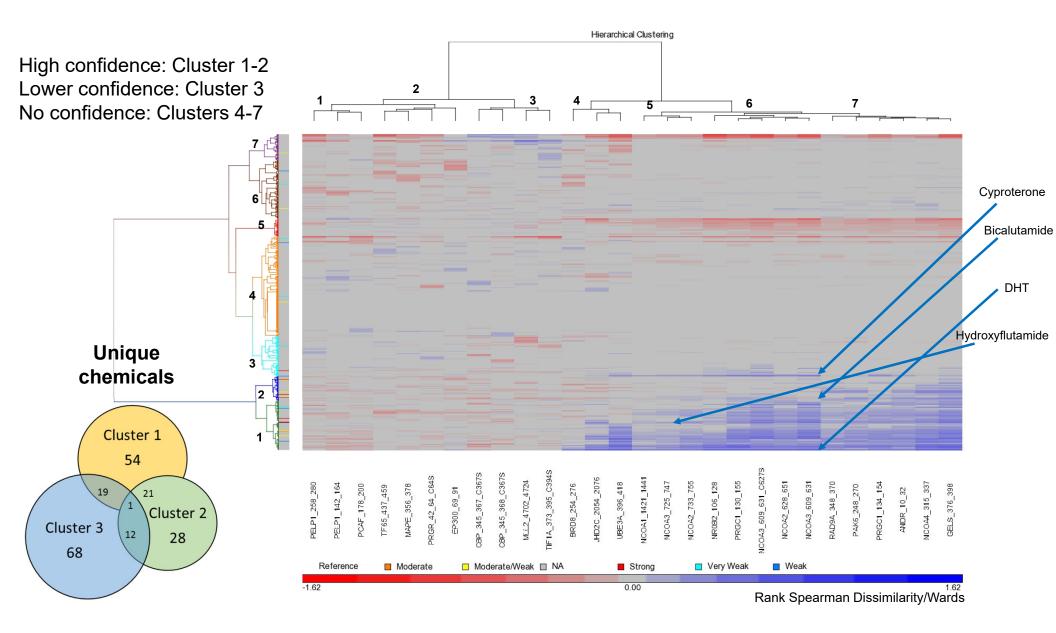
Nuclear receptor(AF2) domain interacts with LxxLL motifs of coregulators



LBD; Ligand binding domain DBD; DNA binding domain AF2; Activation function LxxLL; peptide binding sequence

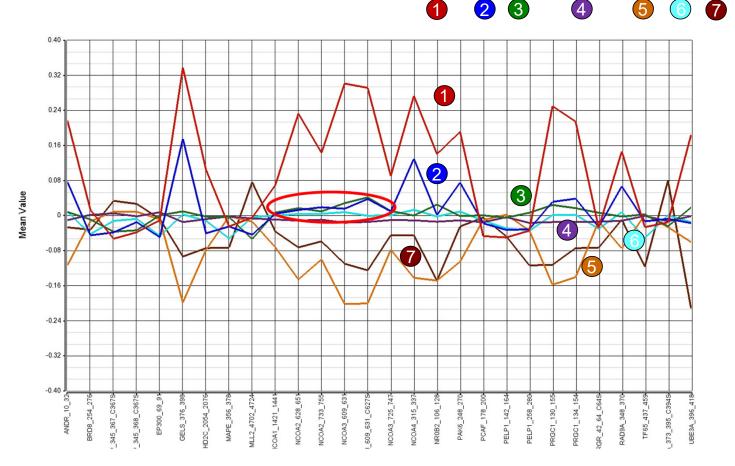
Figure 2: Outline of the nuclear receptor - coregulator interactions in the µ-wells

Image: pamgene.com

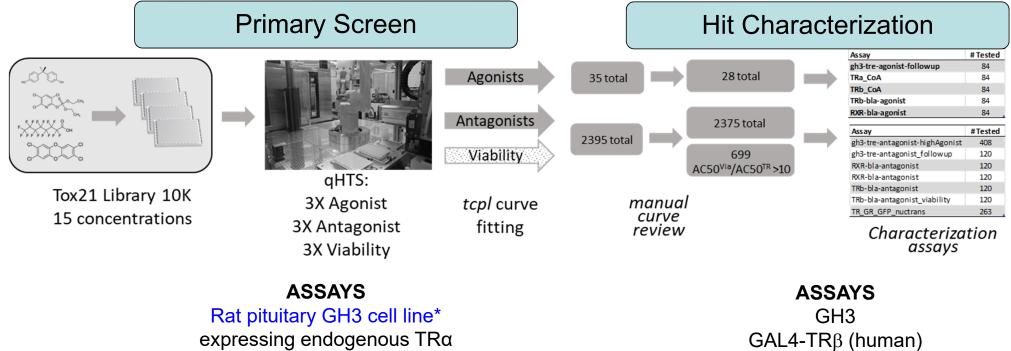


### **Co-regulator Recruitment Patterns**

- Mean value of cluster plotted per coregulator
- Loss of binding seen in cluster 2&3 versus 1 (red oval)
- These represent SRC coactivators that have histone acetyl transferase activity
- Selective receptor modulators; likely would influence biological response



## Thyroid Hormone Receptor Modulators: Tox21 qHTS Campaign

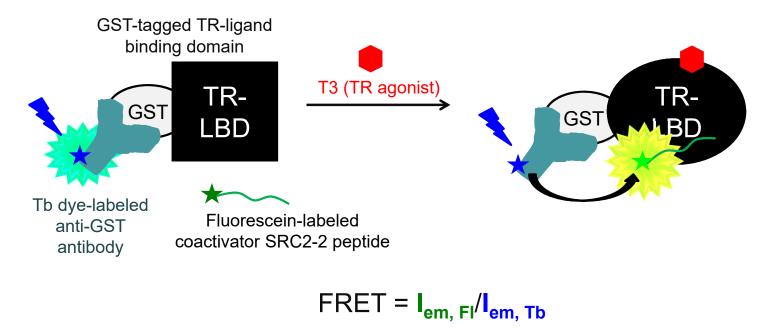


and TRβ, with TRE regulating luciferase expression Cell viability GH3 GAL4-TRβ (human) GAL4-RXRα (human) TRα/TRβ coactivator recruitment TR nuclear translocation

\* Developed by Albertinka Murk, Wageningen University, the Netherlands

#### TR Modulator Hit Characterization: TR Coactivator Assay (Invitrogen):

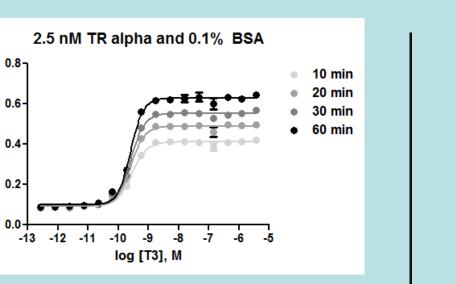
- Direct-acting modulators should regulate coactivator recruitment
- Test in both agonist (recruitment) and antagonist (disassociation) format



High FRET

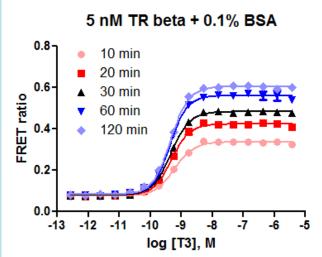
Low FRET

#### **Optimization of TR Coactivator Assay**



2.5 nM TRα-LBD	S/B	CV%	EC50 (M)	
10 min	5.0 3.6		2.786E-10	
20 min	5.9	3.3	2.605E-10	
30 min	6.7	4.2	2.629E-10	
60 min	7.0	4.6	2.571E-10	

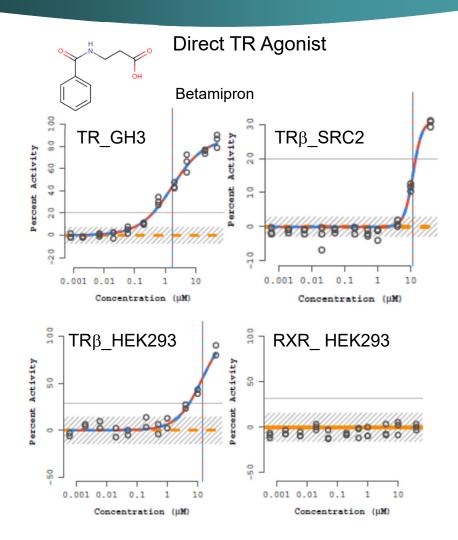
FRET ratio



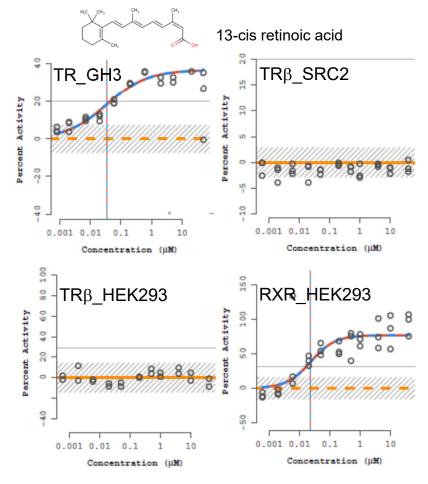
TRβ-LBD	S/B	CV%	EC50 (M)	
10 min	4.4	3.9	5.911E-10	
20 min	5.3	4.7	4.786E-10	
30 min	6.1	4.6	4.661E-10	
60 min	6.8	4.8	4.380E-10	
120 min	7.4	5.7	4.507E-10	

U.S. Environmental Protection Agenc

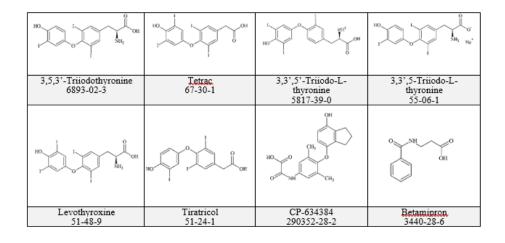
#### **Example Agonists**



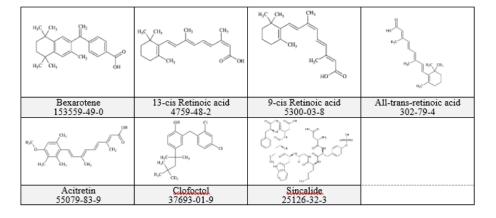
Indirect TR Agonist/RXR Agonist







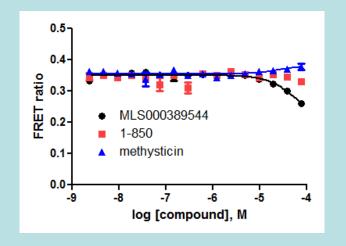
Direct



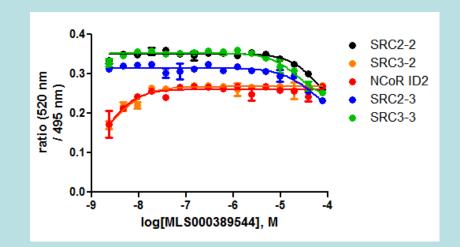
Indirect

#### TR-FRET TRβ Coactivator Assay, Antagonist Mode

Three known TR antagonists tested with the SRC2-2 peptide:



MLS000389544 tested on various coactivator peptides:

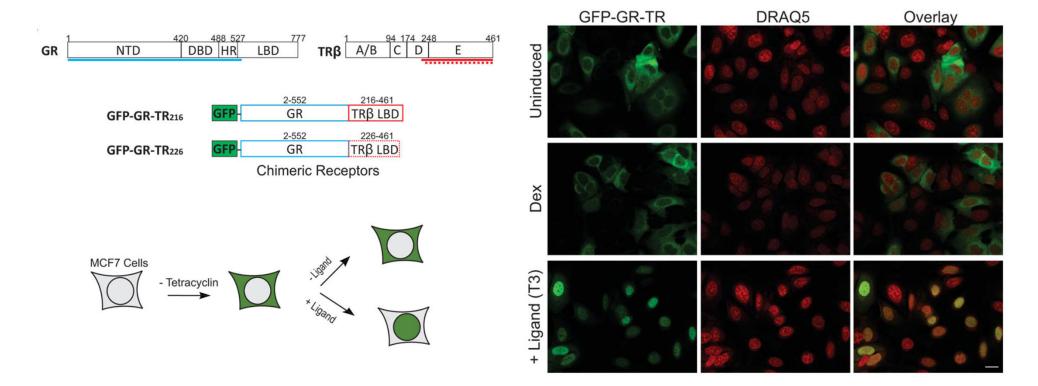


No understanding of why this assay failed.

U.S. Environmental Protection Agency

Reference for MLS000389544: J Biomol Screen. 2011 Jul;16(6):618-27.

## Development of a TR Nuclear Localization Assay

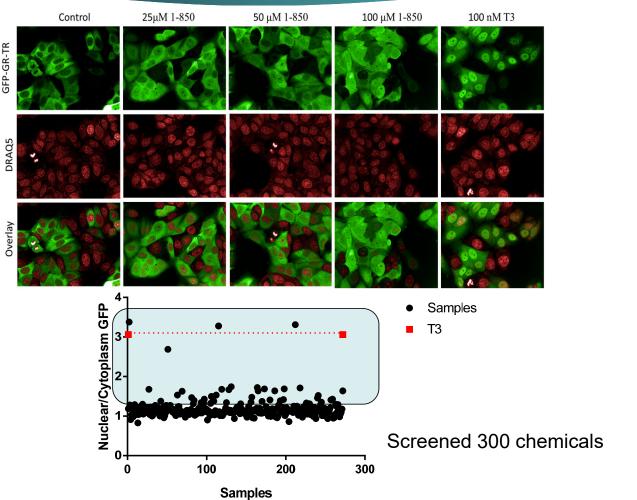


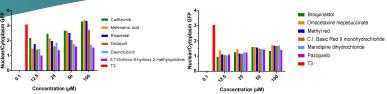
Stavreva et al., Toxicology 368–369: 69-79, 2016.

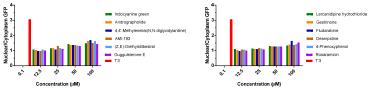
### **TR Nuclear Translocation Assay**

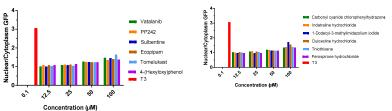
GFP-GR-TR

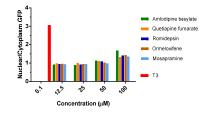
Overlay



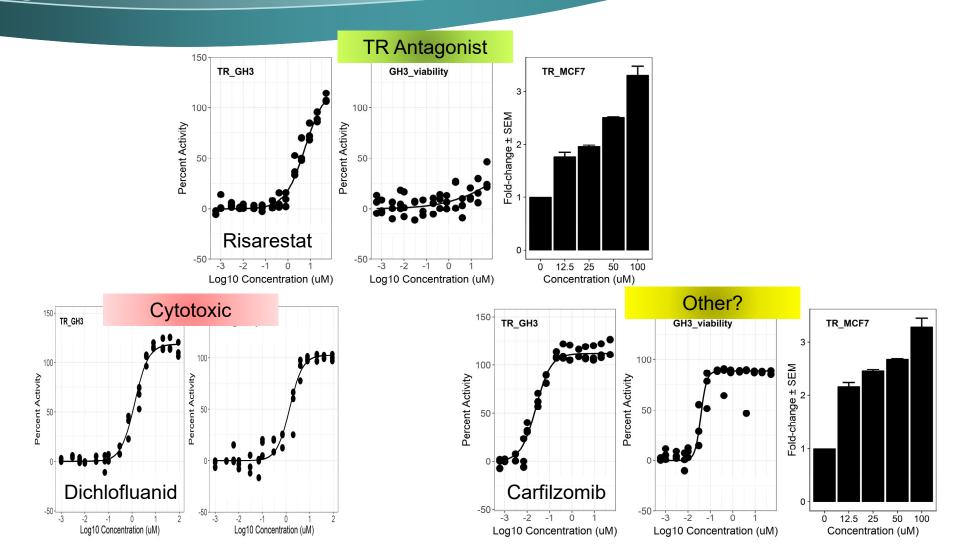








### **Antagonist Characterization Examples**

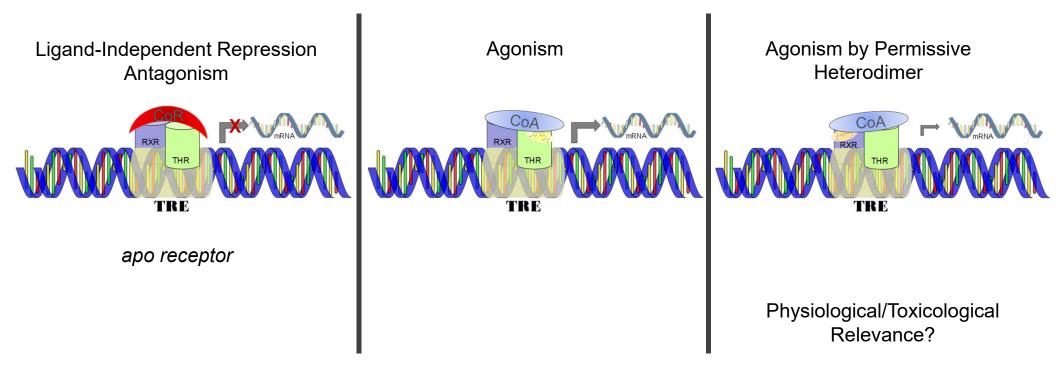


# TR Antagonist Candidates

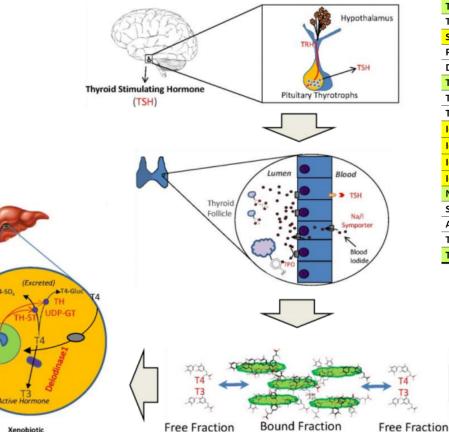
Diclazuril	Mefenamic acid	Risarestat
101831-37-2	61-68-7	79714-31-1

( + ) + ) + ( + ) +	NH <sub>2</sub>	HO CH <sub>3</sub>		Carfilzomib	PP242		Ecopipam
				868540-17-4	1092351-67-1	20830-81-3	112108-01-7
Omacetaxine mepesuccinate 26833-87-4	C.I. Basic Red 9 monohydrochloride 569-61-9	Methyl red 493-52-7	Lercanidipine hydrochloride 132866-11-6	n de la companya. De la companya.			
				Pazopanib 444731-52-6			

## Modes of TR Modulation



### **Thyroid Axis Targets**



Screening Assay Status **Molecular Target** Existing In Development **TRH Receptor** X (ToxCast) **TSH Receptor** Х Sodium-Iodide Symporter (NIS) Х Pendrin Dual Oxidase (DUOX) Thyroperoxidase (TPO) Х TH Serum Transport Proteins Х **TH Membrane Transporters** Iodothyronine Deiodinase Type I Х Х Iodothyronine Deiodinase Type II Iodothyronine Deiodinase Type III Х **Iodotyrosine Deiodinase** Х **Nuclear Receptors** X (ToxCast) Sulfation and Glucuronidation **Alanine Side Chain Activation TH Receptor Binding** TH Transcription (Agonist/Antagonist) X (ToxCast)

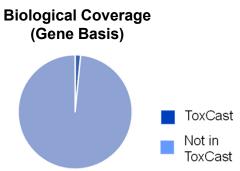
TH Transmembrane Transporters (MCT8) T2 Transcriptio

Xenobiotic (activates CAR, PXR or AhR)

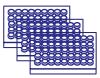
T4-50,

### Some Existing Limitations in High-Throughput and *In Vitro* Test Systems

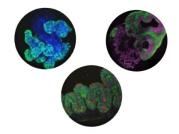




Chemical Coverage and Specific Chemical Types (e.g., VOCs)



Organ and Tissue Responses





Metabolic

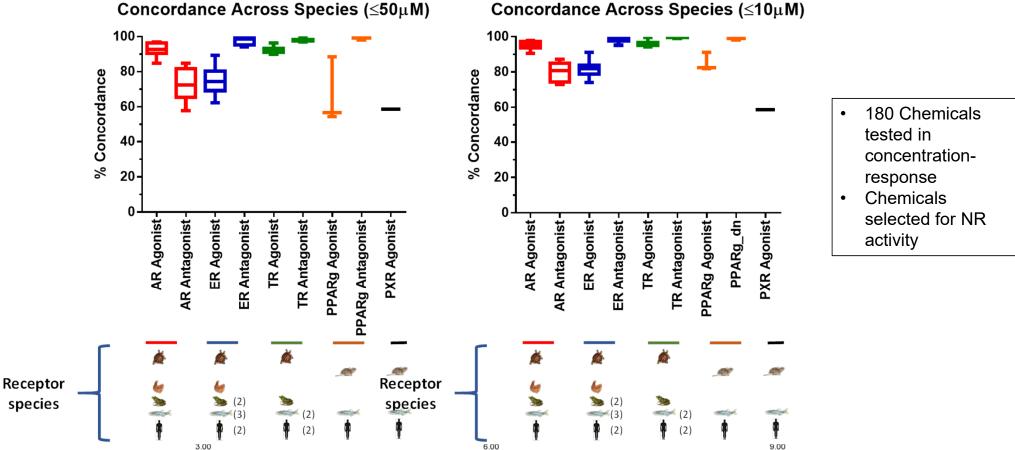
## Assessing Cross-Species Differences in Response

#### Multispecies Attagene Trans Reporter Assav Gal4 promoter RTU A RTU A RTU B RTU B RTU C RTU C 23 GAL4-NR RTUs ERa LBD transfect Gal4 DBD $\overline{\mathbf{O}}$ 4xUAS TATA pool ERα LBD 9 Estrogen 兦 Gal4 DBD 10/10 Highly multiplexed GAL4-NRs 4XUAS TATA FACTORIAL MS (human, reporter gene bird, fish. turtle, frog) 卫 assay ERa Gal Profile reporter RTU RNAs DBD 0 4xUAS TATA 仑 5 Androgen (human, bird, fish, turtle, frog) -0 NR activity profile GAL4-NRs 6 Thyroid (human, fish, frog, turtle) 0 GAL4-NRs in the first of the first firs 3 PPAR (human, mouse, fish) -0 GAL4-NRs Houck et al., unpublished

NR family NR Class Species Sequence ID ER1 Danio rerio BC162466 ER2a Fish Danio rerio BC044349 ER2b BC086848 Danio rerio ER1 Xenopus laevis NM 001089617 Amphibian ER2 Estrogen Xenopus laevis NM 001130954 ER1 Reptilian Chrysemys picta NM 001282246 ER1 Avian Gallus gallus NM 205183 ERa Homo Sapiens NM 000125 Mammalian ERb Homo Sapiens NM 001437 AR Fish Danio rerio NM 001083123 AR Amphibian Xenopus laevis NM 001090884 Androgen Chrysemys picta XM 005279527 AR Reptilian NM 001040090 AR Avian Gallus gallus AR Mammalian NM 000044 Homo Sapiens TRa Danio rerio BC096778 Fish TRb BC163114 Danio rerio TRa Amphibian Xenopus laevis NM 001088126 Thyroid XM 005294120 TRa Reptilian Chrvsemvs picta NM 199334 THRa Homo Sapiens Mammalian THRb Homo Sapiens NM 000461 PPARg NM\_131467 Fish Danio rerio PPAR PPARg Mus musculus NM 001127330 Mammalian PPARg BC006811 Homo Sapiens

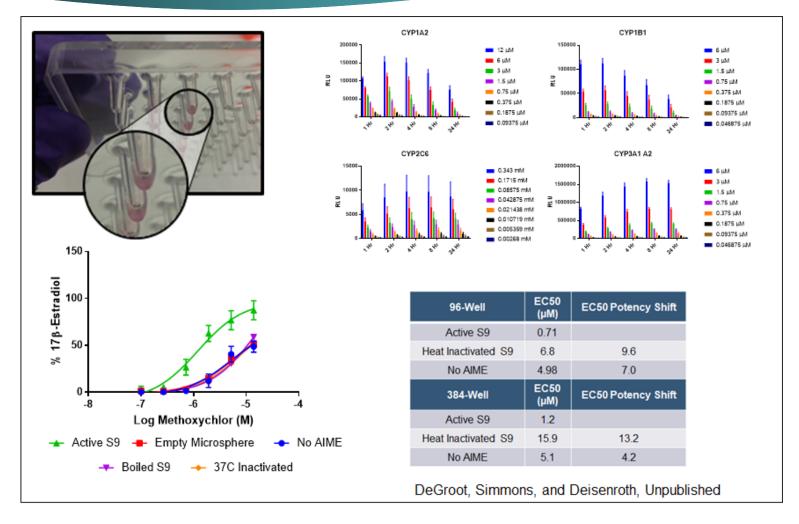
- Host cell: human HepG2
- Agonist mode for all receptors
- Antagonist for ER and AR

## **Cross-Species Differences in Nuclear Receptor Responses**

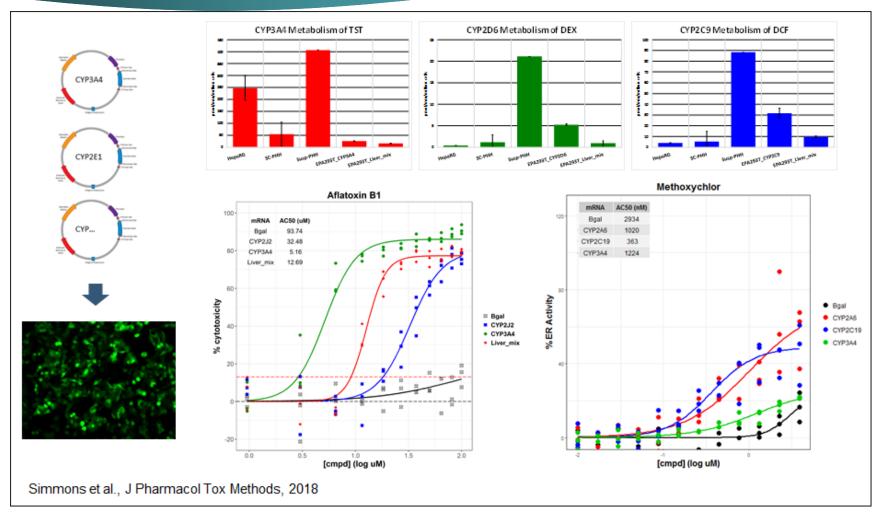


Concordance Across Species (≤10µM)

#### Assays Retrofit for Xenobiotic Metabolism: Extracellular

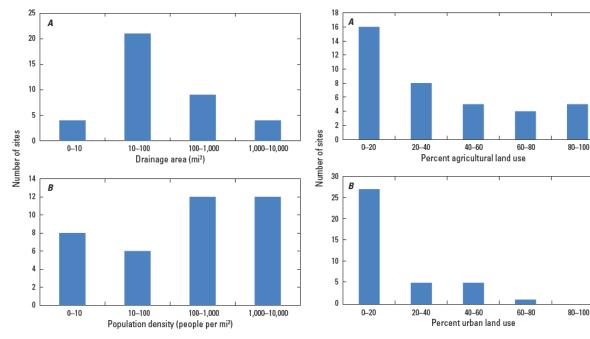


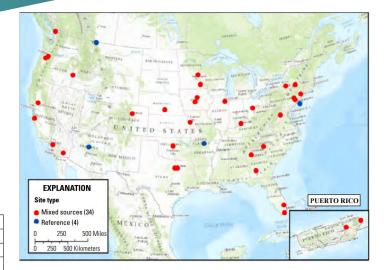
#### Assays Retrofit for Xenobiotic Metabolism: Intracellular



#### Environmental Monitoring Application: Nationwide Streams Surveillance

- 38 total sites (4 reference sites) across US and PR
- Water samples collected 2012-2014
- Locations varied by watershed drainage

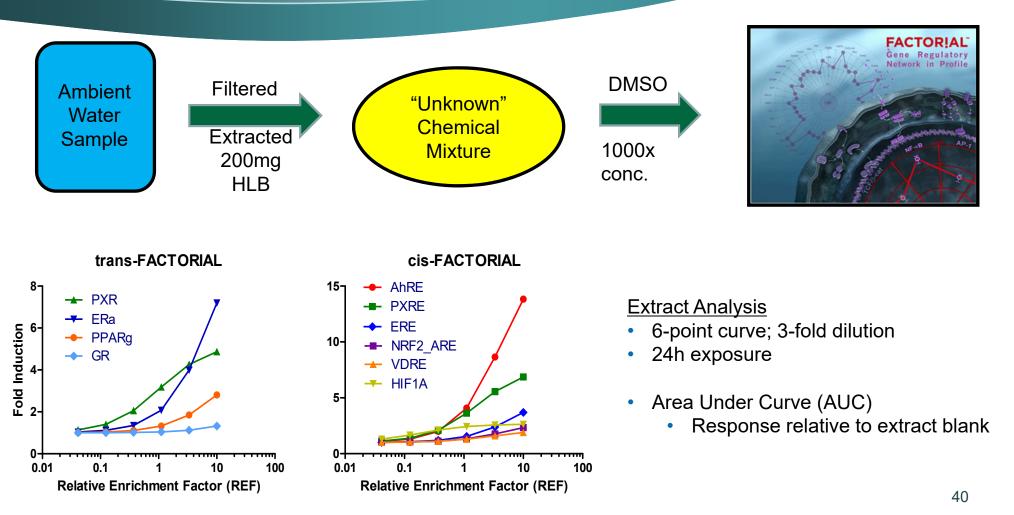








## **Bioassay Analysis Workflow**

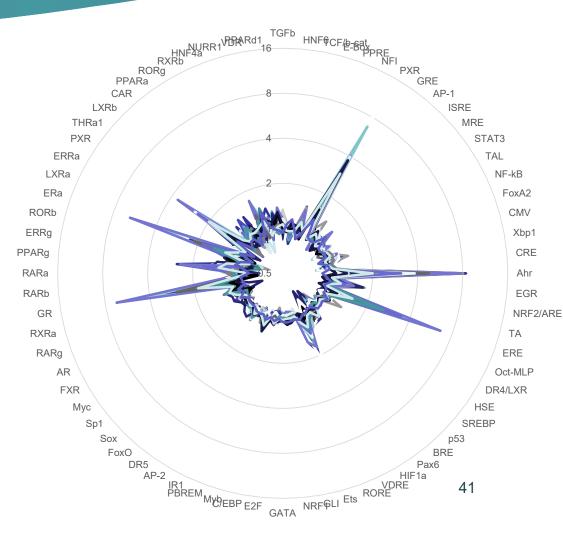


### **Bioassay Results**

- 26/70 endpoints AUC >1.25-fold (borderline active)
- 11/70 endpoints AUC >1.5-fold (active)

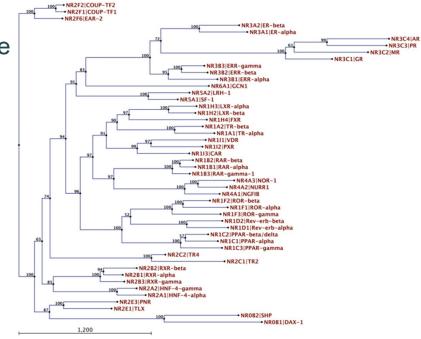
#### Active Endpoints

- PXRE, PXR, AhRE 30-36 sites
- ERE 17 sites
- ER $\alpha$ , PPAR $\gamma$  10 sites
- GR, VDRE, NRF2 6-8 sites
- RORE, RXR $\beta$  2 sites



### **Concluding Remarks**

- In vitro/alternative and computational approaches are valuable for chemical prioritization, especially where we understand the targets and toxicity pathways
- Nuclear receptors are an inherently important target of environmental chemicals
- Endocrine disruption is one important mode of action mediated by NR's but there are many more receptors with varied, important physiology
- Using high-throughput approaches will require systematically addressing key technical and data analysis challenges
  - assay interferences
  - selective receptor modulators
  - cytotoxicity



# **Thank You for Your Attention!**

#### ICCVAM

- Nichole Kleinstreuer
- Patricia Ceger
  - Warren Casey
  - David Allen

#### EPA/ORD/NHEERL

- Mike Hornung
- Susan Laws
- Tammy Stoker
- Jun Wang
- Daniel Hallinger
- Ashley Murr
- Angela Buckalew
- Joseph Korte
- Jennifer Olker
- Jeff Denny
- Carsten Knutsen
- Phillip Hartig
- Mary Cardon
- Sigmund Degitz
- Dan Villeneuve
- Anthony Schroeder
- Gerald Ankley
- Brett R Blackwell

EPA's National Center for Computational Toxicology

#### NCATS/NIH

- Menghang Xia
- Ruili Wang
- Chia-Wen Hsu
- OECD
- Patience Browne

#### NCI/NIH

- Diana Stareva
- Vikas Soni
- Lyuba Varticovski
- Razi Raziuddin
- Gordon Hager

#### NCCT/ORD/EPA

- Katie Paul-Friedman
- Richard Judson
- Kevin Crofton
- Rusty Thomas
- Audrey Bone
- Ann Richard
- Matt Martin
- Tom Knudsen
- Nisha Sipes
- Eric Watt
- David Dix
- Woody Setzer