

## Imaging-based phenotypic profiling for high-throughput chemical screening at the U.S. Environmental Protection Agency

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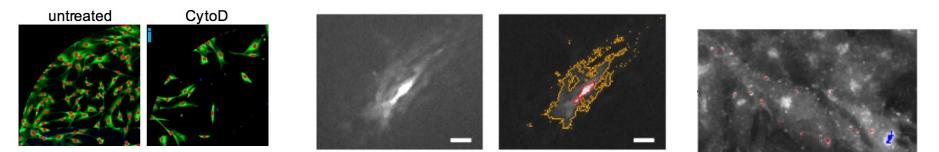
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Presentation for UFZ December 7<sup>th</sup>, 2021

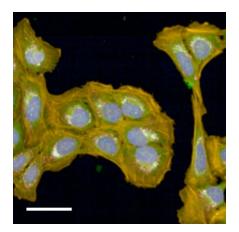


## Introduction: Dr. Johanna Nyffeler

- BSc in Biochemistry, MSc in Genetics
- PhD at University of Konstanz, Germany
  - group of Dr. Marcel Leist
  - development of high-content assays for *in vitro* developmental neurotoxicology



- PostDoc at Center for Computational Toxicology & Exposure (CCTE), US EPA
  - group of Dr. Joshua Harrill
  - high-throughput image-based profiling ('Cell Painting'), computational toxicology







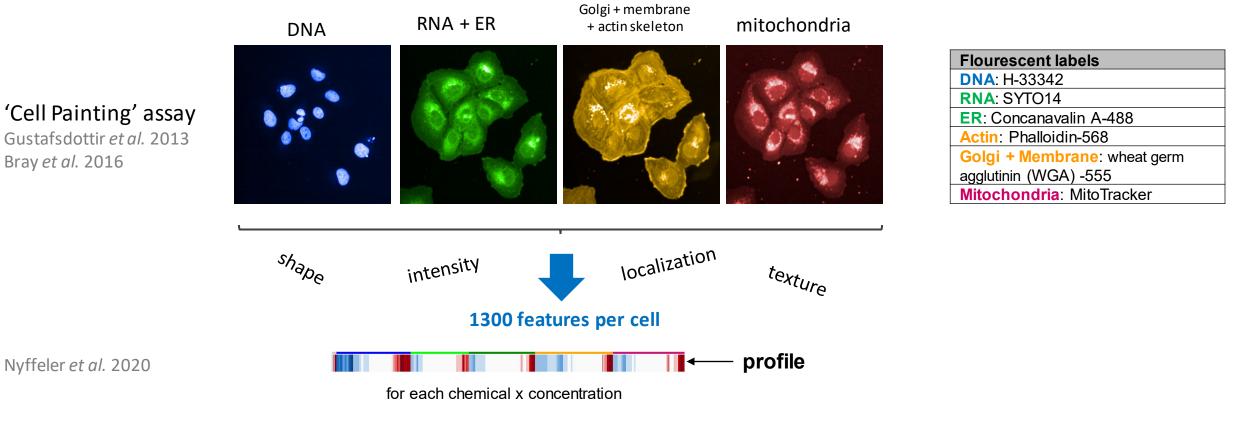
### **Overview**

- 1. What is imaging-based phenotypic profiling?
- 2. Implementation at CCTE/EPA
- 3. Aims/Focus for CCTE/EPA
- 4. Application 1: Potency estimation
- 5. Application 2: Mechanistic prediction



## What is Imaging-Based Phenotypic Profiling?

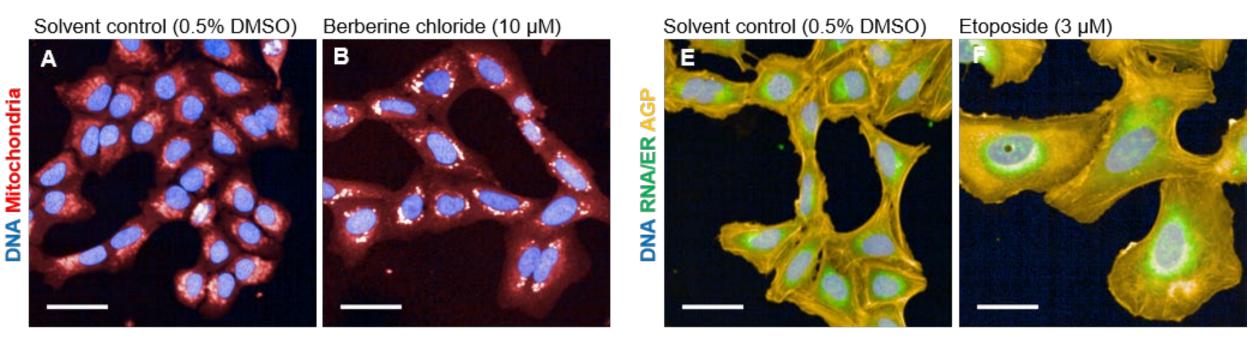
- labeling of various cell organelles with fluorescent probes in *in vitro* cultures
- assessing a large variety of morphological features on individual cells



Cell Painting = Phenotypic Profiling High-Throughput Phenotypic Profiling = HTPP



## **Example Chemicals: Qualitative Observation**



 $\rightarrow$  Cells are larger

Strong phenotypes are observable qualitatively

 $\rightarrow$  Mitochondrial

compactness/texture

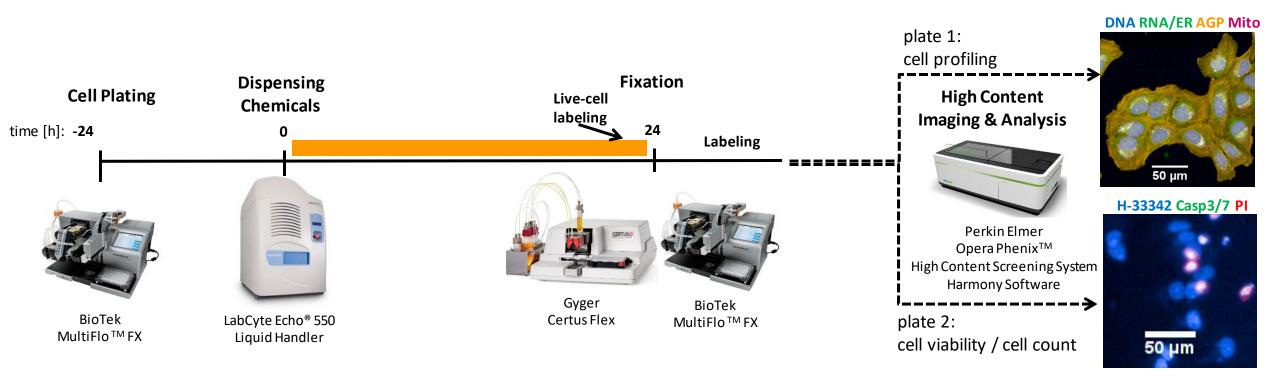
adapted from Nyffeler *et al.* (2020) Toxicol Appl Pharmacol. PMID: 31899216



## Implementation at CCTE/EPA

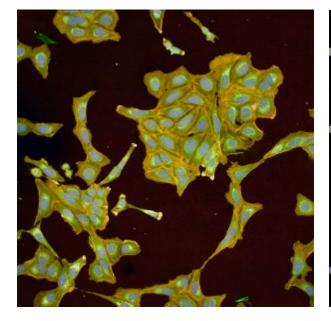


## **Laboratory Workflow**

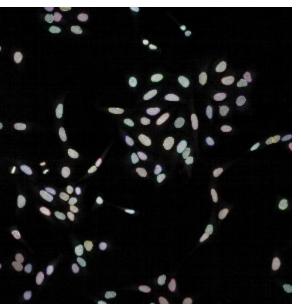




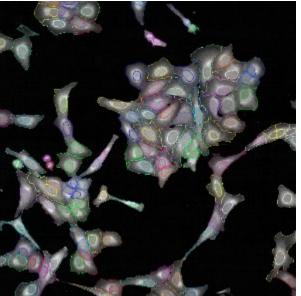
## Image Analysis Workflow → Image Segmentation



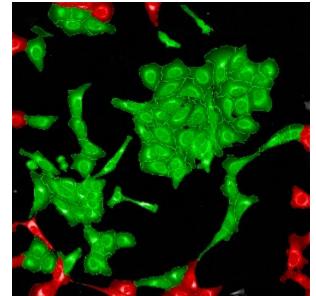
1. find nuclei

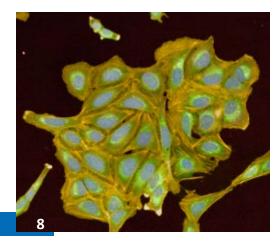


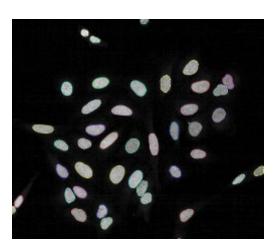
2. find cell outline

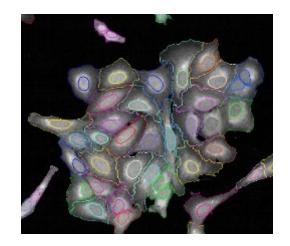


3. reject border objects





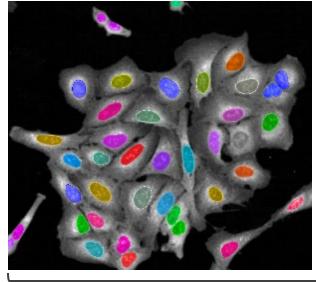


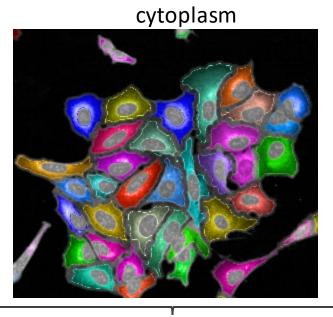




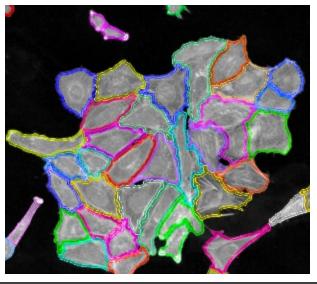
## **Define Cellular Compartments**

nuclei

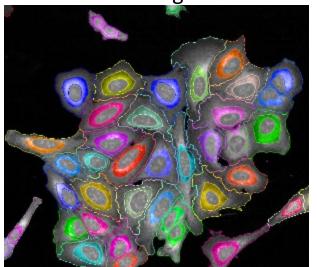


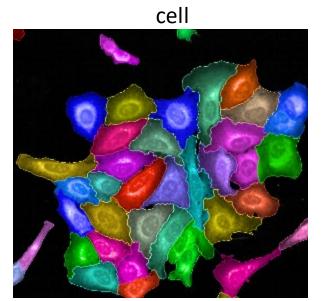


membrane



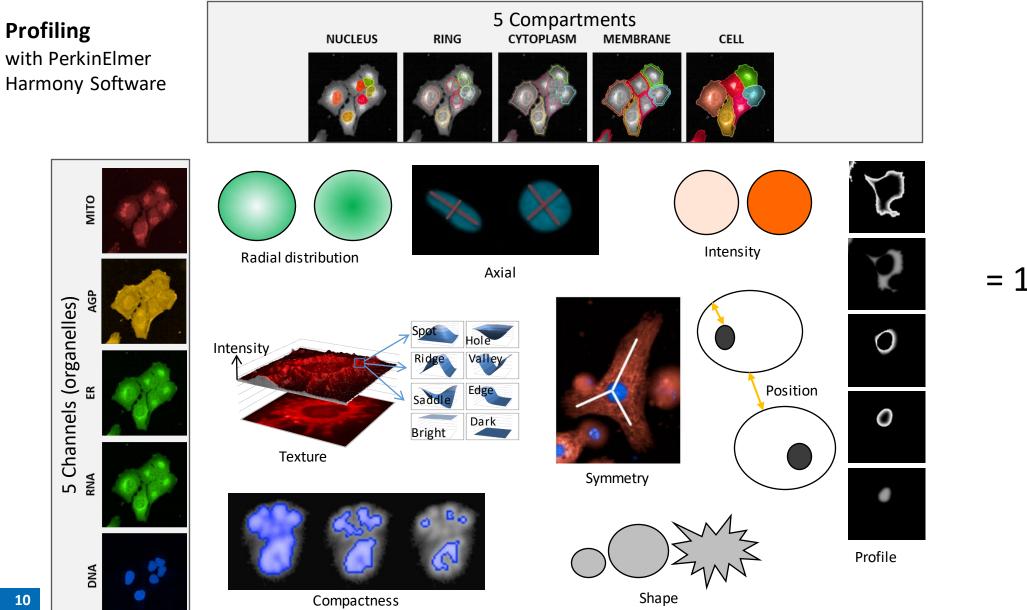
ring







## **Image Processing**

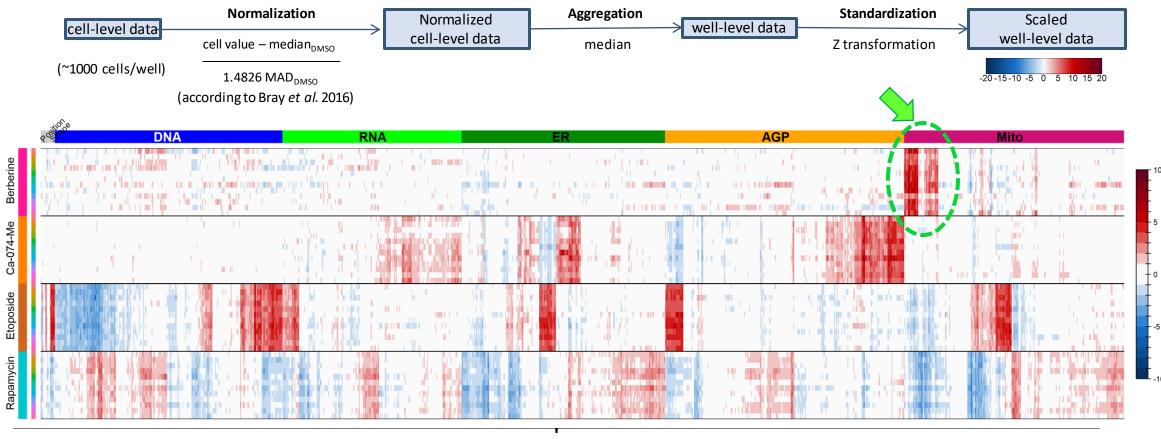


= 1300 features per cell

With illustrations from Perkin Elmer



## **Example Chemicals: Quantitative Observation**



1300 features

### Qualitative observations can be quantified

adapted from Nyffeler *et al.* (2020) Toxicol Appl Pharmacol. PMID: 31899216



# Aim for CCTE/EPA



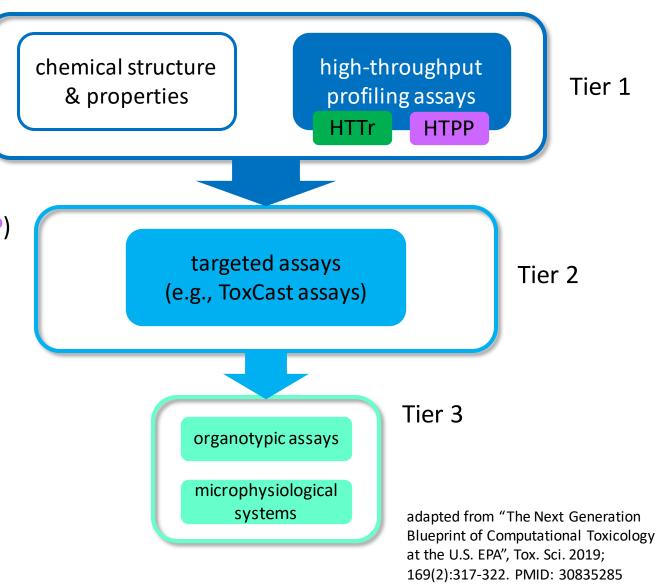
### Tiered Hazard Evaluation Strategy based on New Approach Methods (NAMs)

### **Profiling Assays**

- untargeted
- measure large number of endpoints (e.g., transcripts, phenotypic features)
- high-throughput transcriptomics (HTTr) (Harrill et al. 2021, PMID: 33538836)
- high-throughput phenotypic profiling (HTPP) (Nyffeler et al. 2020, PMID: 31899216)

### Focus

• Prioritization: False positives are preferred over false negatives



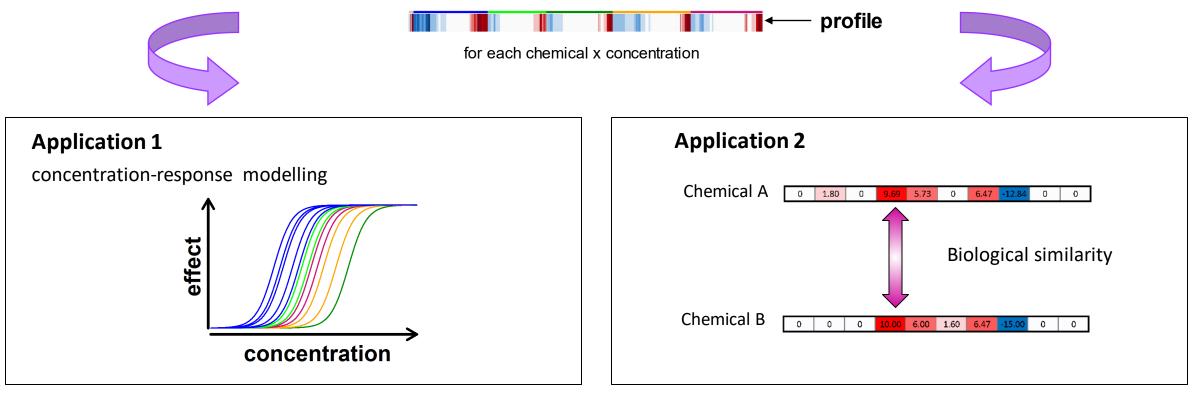


## **Challenges of Environmental Chemicals**

- Often low expected bioactivity
- Often lack a specific molecular target in human-based cell models
- 'Poly-pharmacology'
- Responses can be associated with general cell stress
- ⇒ more challenging for hit identification than drug-like chemicals



## **Two Applications**



### Potency estimation:

### in vitro point-of-departure (POD)

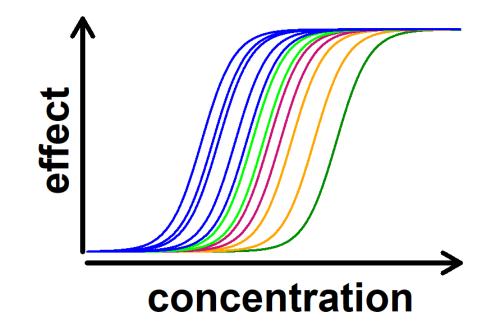
- Nyffeler *et al.* (2020) Toxicol Appl Pharmacol. PMID: 31899216
- Willis *et al.* (2020). SLAS Discov. PMID: 32546035
- Nyffeler *et al.* (2021). SLAS Discov. PMID: 32862757

Compare profiles with annotated reference chemicals → putative mechanisms

work in progress



# Application 1: Potency Estimation





## **U-2 OS ToxCast Screen Experimental Design**

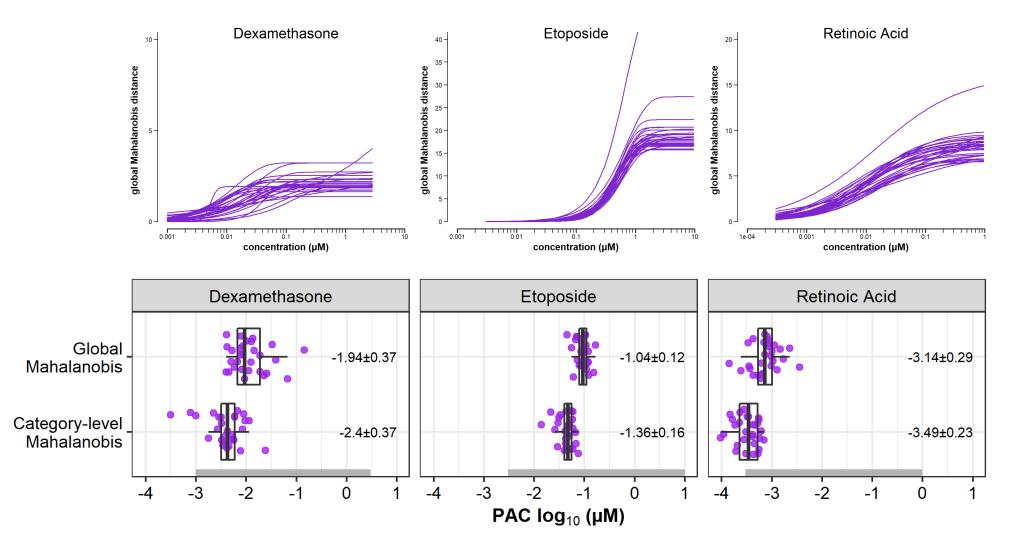
Parameter	Multiplier	Notes	
Cell Type(s)	1	U-2 OS	
Time Points:	1	24 hours	
Chemicals	1,202	<ul> <li>TSCA Chemicals of interest to US EPA</li> <li>Includes 462 APCRA case study chemicals</li> <li>Includes 179 chemicals with annotated molecular targets</li> </ul>	
Concentrations:	8	3.5 log <sub>10</sub> units; ~half-log <sub>10</sub> spacing	
Biological Replicates:	4		

### Reference chemicals run on each plate

	Chemical	Molecular Target	Tested Range
Weak	Dexamethasone	Glucocorticoid receptor agonist	0.001-3 μM
Medium	all-trans-Retinoic Acid	Retinoic acid receptor agonist	$0.0003 - 1  \mu M$
Strong	Etoposide	DNA topoisomerase inhibitor	0.03 - 10 μM
Extra strong	Trichostatin A	Histone deacetylase inhibitor	1 μΜ



## **Reproducibility: Potencies**

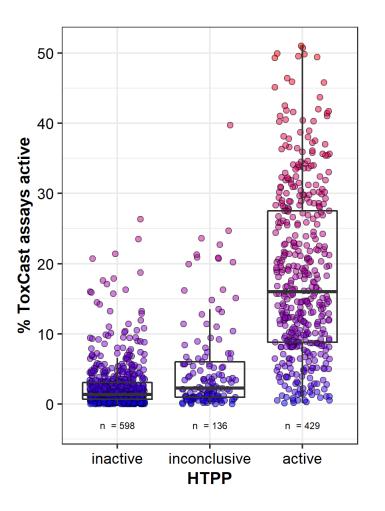


⇒ Potency estimates vary less than ½ an order of magnitude



## **HTPP Screening Results (1)**

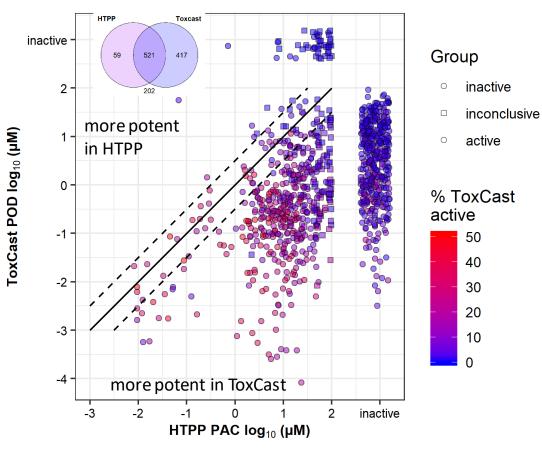
Active chemicals:



- → ~ 40% of chemicals were active
- $\Rightarrow$  Most activity is > 10  $\mu$ M
- ⇒ Chemicals active in HTPP are more often active in many ToxCast assays

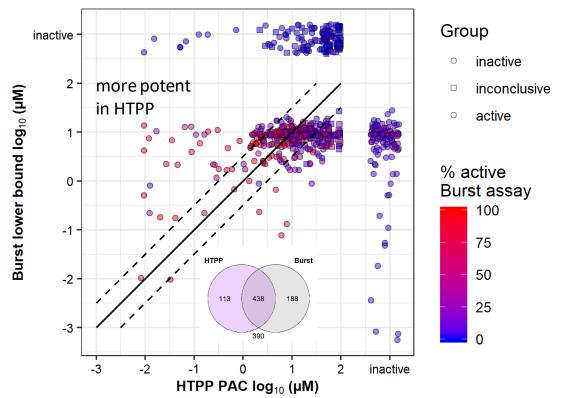


## **HTPP Screening Results (2)**



### Comparison with ToxCast screening results:

⇒ HTPP with a single cell line less sensitive than all ToxCast assays combined....

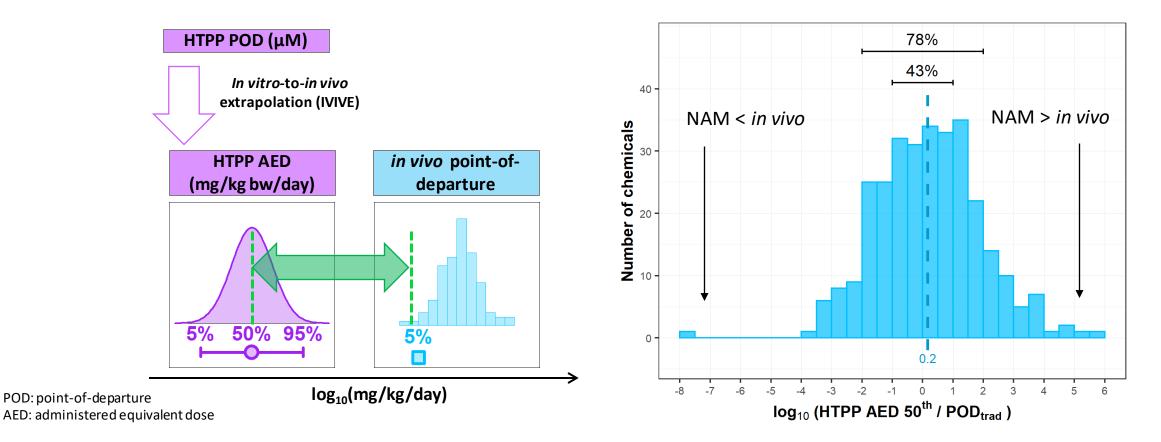


... but more sensitive than the ToxCast cytotoxicity burst estimate



## **Comparison to in vivo Effect Values**

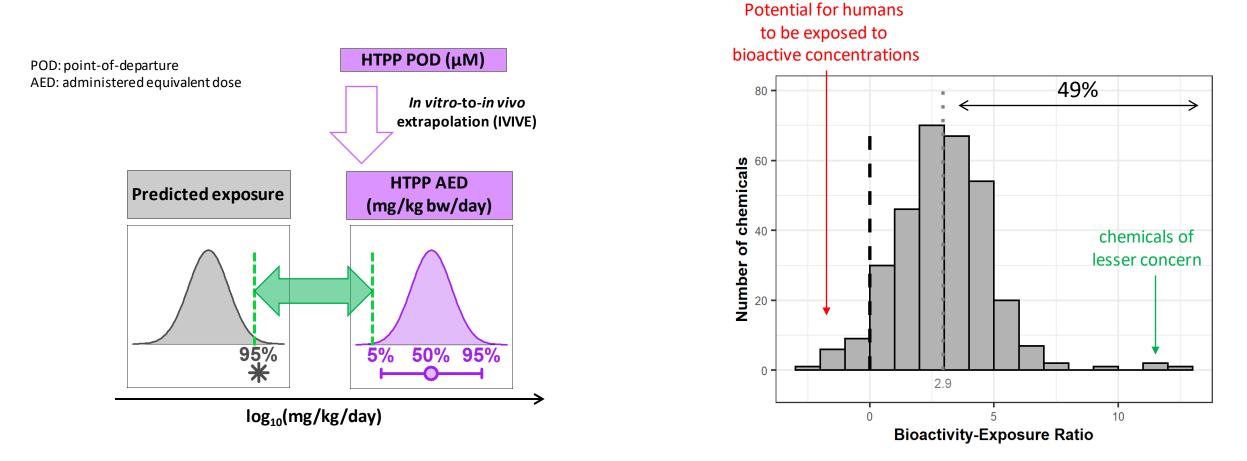
• 303 chemicals were active and had pharmacokinetic (PK) information



> 78% of HTPP AED are within 2 orders of magnitude of the *in vivo* POD



## **Comparison to Exposure Estimates**



- ⇒ for 49% of chemicals, predicted exposure is > 1000x lower than estimated bioactivity
- for a small set of chemicals, the BER was negative, indicating a potential for humans to be exposed to bioactive concentrations of these chemicals





# Mechanistic Prediction



## **Feature Selection & Profile Comparison**

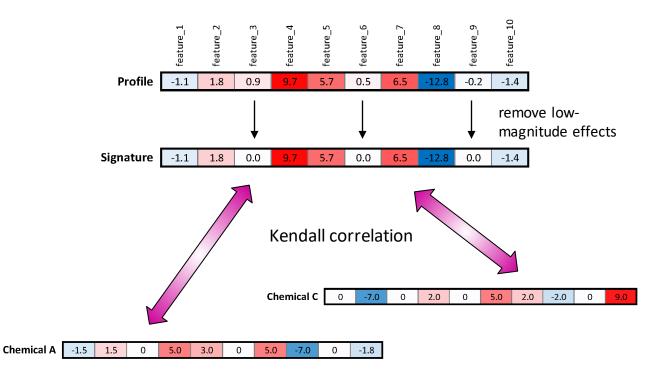
### **Feature Selection**

### 1300 features

- **1.** remove features that do not provide any information (i.e. have 0 variance)
- 2. remove features that are not reproducible (high variation between treatments of different biological replicates)
- **3.** remove features that are highly correlated (using recursive feature elimination)

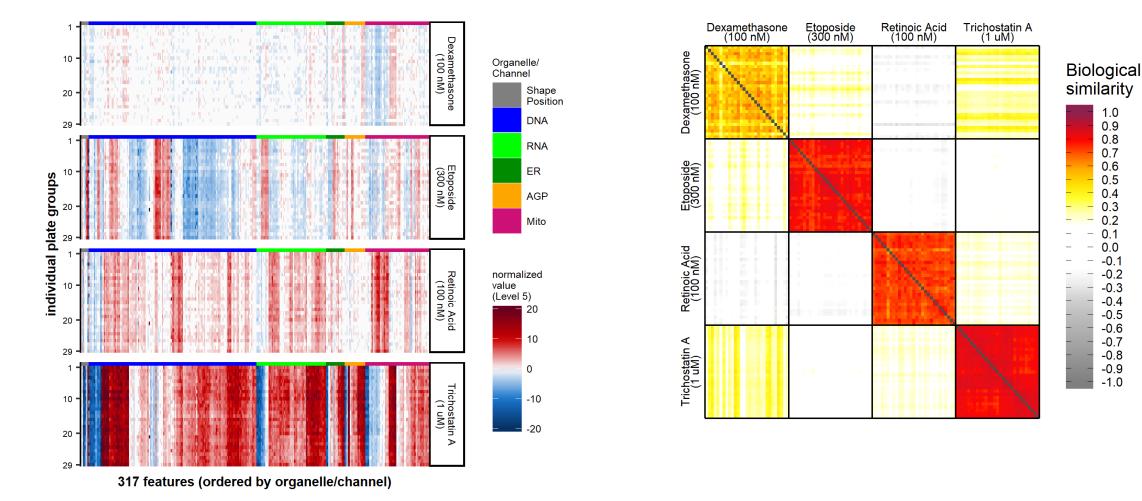
317 features

### **Profile Comparison**





## **Reproducibility: Phenotypic Profiles**



### ⇒ Phenotypic profiles are highly reproducible across different plates

Hypothesis: Chemicals with similar mechanisms will display similar profiles.

Preliminary results. Do not cite or quote.

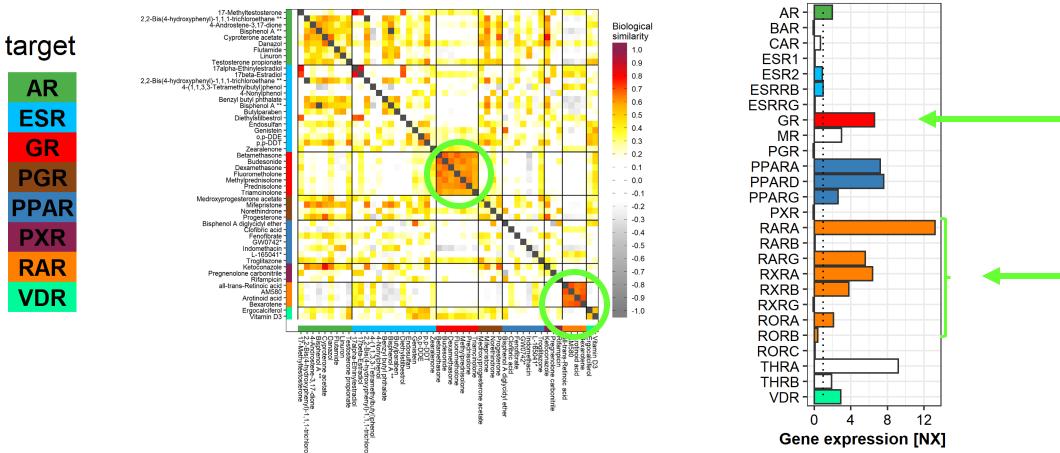


27

## **Example: Nuclear Receptor Modulators (I)**

Gene expression in U-2 OS

52 chemicals were annotated as targeting a nuclear receptor



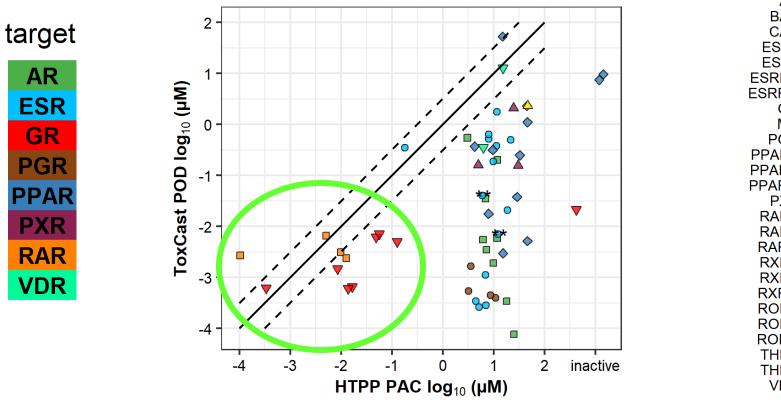
### **Biological similarity in HTPP**

- Agonists of the GR and of RAR/RXR display characteristic profiles
- Expression of a target does not guarantee that characteristic profiles are observed (e.g., PPAR)

Preliminary results. Do not cite or quote.

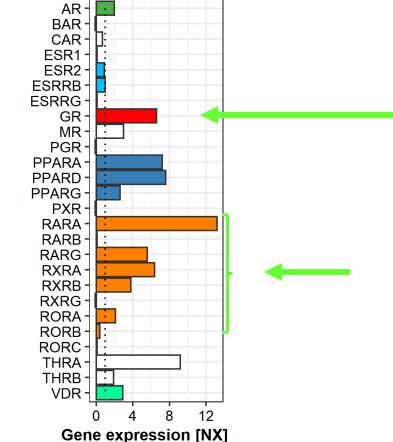


## **Example: Nuclear Receptor Modulators (II)**



**Comparison to ToxCast potencies** 

### Gene expression in U-2 OS



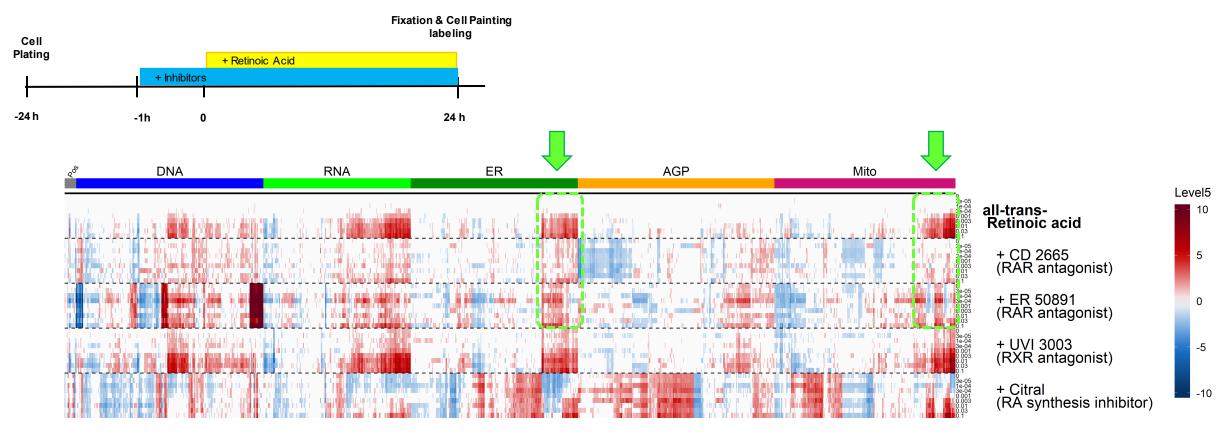
⇒ For two receptor systems that are expressed (GR, RAR/RXR) potencies were comparable with ToxCast

For receptors with no/low expression in this cell line, HTPP was less sensitive than ToxCast

Preliminary results. Do not cite or quote.



## **Pharmacological Blockade of Phenotypic Effects**



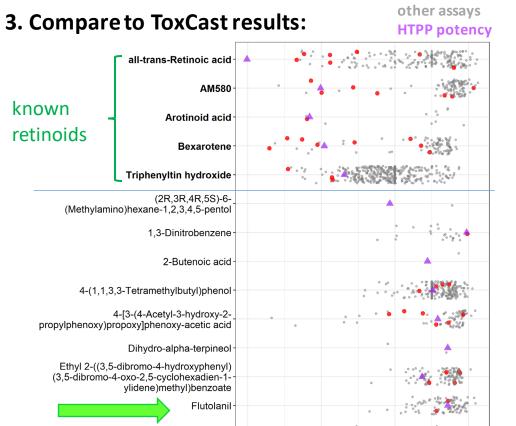
- ⇒ RAR but not RXR antagonists block the retinoid phenotype
- ⇒ Phenotypic profile is related to RAR activation



## **Application: Find Retinoid-like Chemicals**

known

retinoids



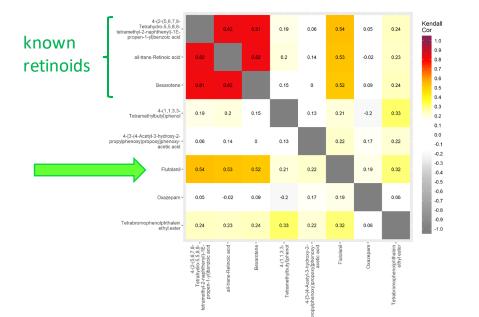
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**RAR/RXR** assays

### 1. Compare profiles to 5 known retinoids:

 $\rightarrow$  10 candidate chemicals

### **2.** Repeat HTPP experiments:



 $\rightarrow$  Flutolanil induced a phenotype similar to retinoids

> $\rightarrow$  Flutolanil had activity in ToxCast assays targeting RAR/RXR

Methimazole

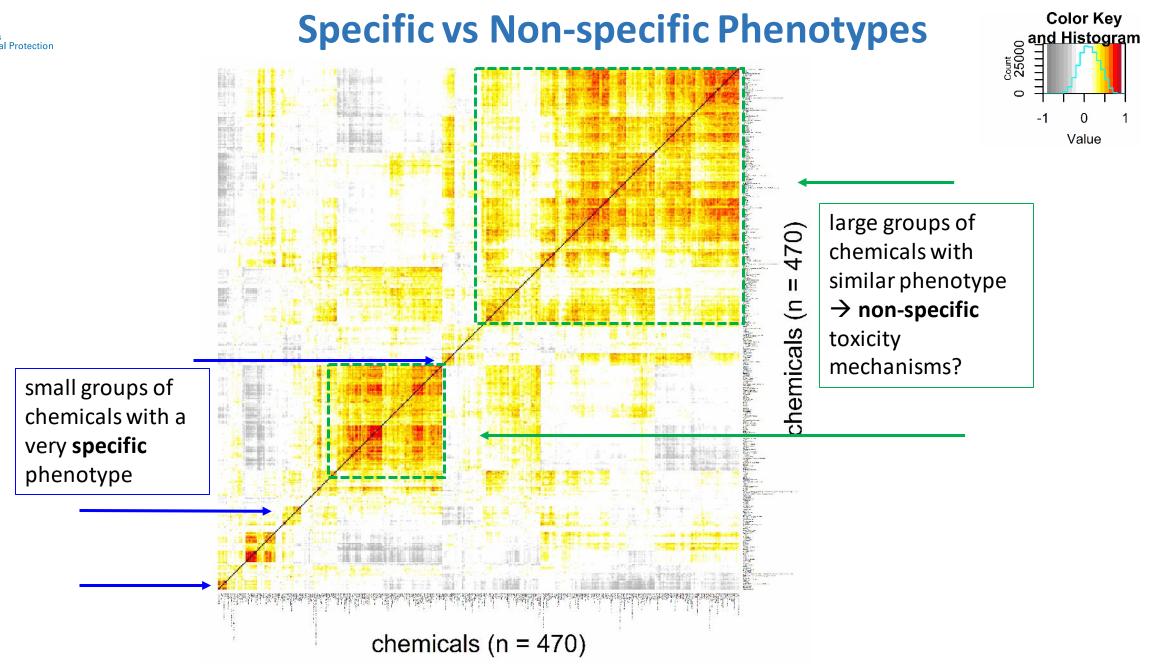
Oxazepam

+ HTPP has the potential to identify environmental chemicals with specific activities

Potency (log10 uM)



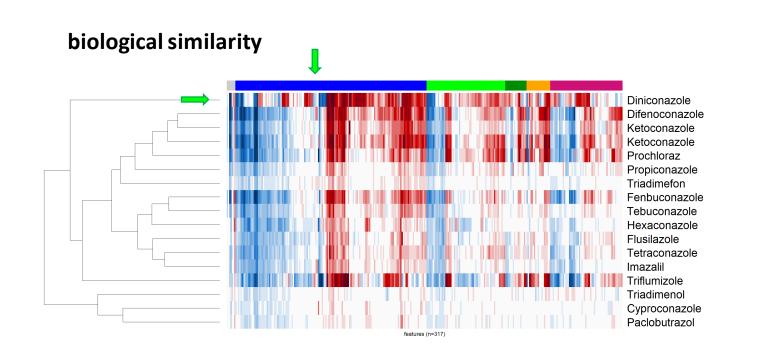
### **Specific vs Non-specific Phenotypes**





## **Application: Grouping of Conazoles**

- group of fungicides
- disturb ergosterol synthesis via CYP51 and CYP61 (target absent in mammals)



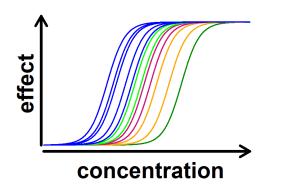
#### structural similarity (based on ToxPrints) Prochloraz Ketoconazole Tetraconazole Triflumizole Imazalil Fenbuconazole Triadimenol Diniconazole Triticonazole Paclobutrazol Tebuconazole Cyproconazole Hexaconazole Fluconazole Triadimefon Flusilazole Propiconazole

Difenoconazole

- ⇒ most conazoles are phenotypically similar
- ⇒ Diniconazole is phenotypically different from the other active conazoles

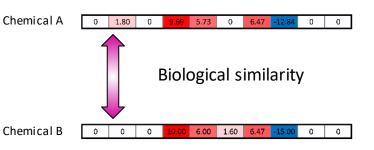


### **Conclusions**



### **Application 1: Potency estimation**

- HTPP can be used to derive *in vitro* potency estimates
- These *in vitro* potency estimates are often comparable and sometimes more conservative than *in vivo* PODs



### **Application 2: Mechanistic prediction**

• Similar mechanisms  $\rightarrow$  similar phenotypes



### Outlook

### • Combine HTPP with HTTr

- compare results, both in terms of potencies and mechanisms
- increased potential to discern molecular mechanisms

### • Expand Coverage of Biological Space

- deploy assay across diverse cell lines that express different receptors/pathways
- proof-of-concept (Gustafsdottir et al. 2013, Willis et al. 2020)
- expansion to other species



## Acknowledgements



### Office of Research and Development (ORD) Center for Computational Toxicology and Exposure (CCTE)

Harrill Lab team

- Joshua Harrill
- Clinton Willis
- Felix Harris
- Rick Brockway
- Megan Culbreth
- Dan Hallinger
- Terri Fairley

Data analysis

- Daniel Chang
- Kathy Coutros
- Logan Everett
- Derik Haggard
- Richard Judson
- Ryan Lougee
- Grace Patlewicz
- Katie Paul Friedman
- Ann Richard
- Woody Setzer
- Imran Shah
- John Wambaugh



• Scott Auerbach