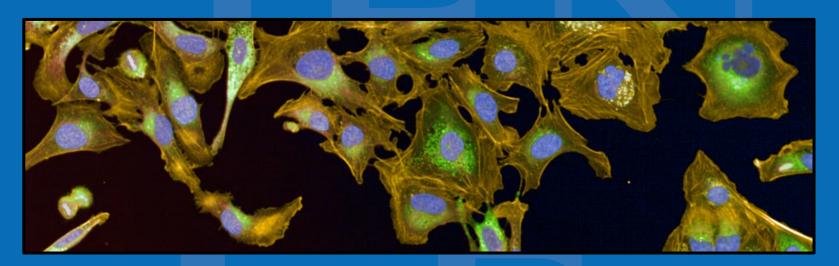


Application of Cell Painting, an Imaging-Based High Throughput Phenotypic Profiling Assay for Bioactivity Screening of Environmental Chemicals

Johanna Nyffeler

ORISE postdoctoral grantee at USEPA Center for Computational Toxicology and Exposure (CCTE)





Conflict of Interest Statement

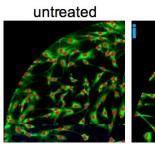
The author declares no conflict of interest.

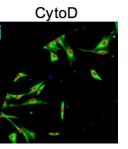
The views expressed in this presentation are those of the author and do not necessarily reflect the views or policies of the U.S. EPA.

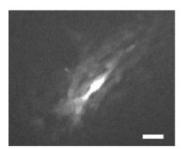


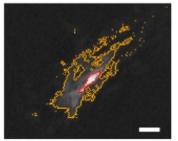
Introduction: Dr. Johanna Nyffeler

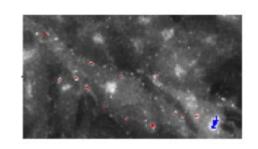
- BSc in Biochemistry, MSc in Genetics
- PhD at University of Konstanz, Germany
 - group of Dr. Marcel Leist
 - development of high-content assays for in vitro developmental neurotoxicology



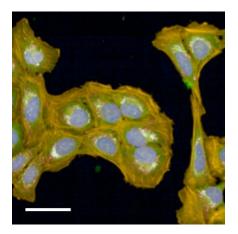








- PostDoc at Center for Computational Toxicology & Exposure (CCTE), US EPA
 - group of Dr. Joshua Harrill
 - high-throughput image-based profiling ('Cell Painting'), computational toxicology







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

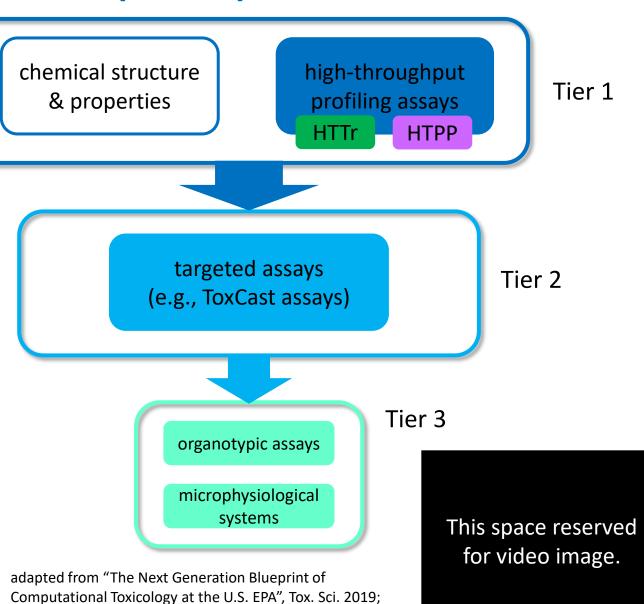
169(2):317-322. PMID: 30835285

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)

Goals

- potency estimation
- mechanistic prediction

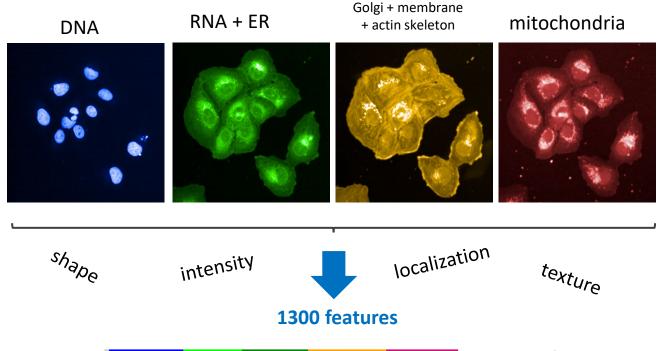




What is Imaging-Based Phenotypic Profiling?

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

'Cell Painting' assay Gustafsdottir et al. 2013 Bray et al. 2016



Flourescent labels
DNA: H-33342
RNA: SYTO14
ER: Concanavalin A-488
Actin: Phalloidin-568
Golgi + Membrane: wheat germ agglutinin (WGA) -555
Mitochondria: MitoTracker

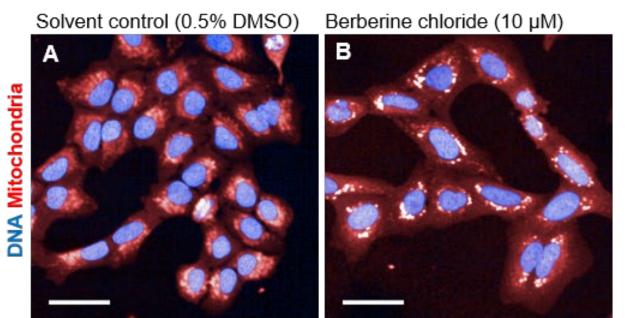
Nyffeler et al. 2020



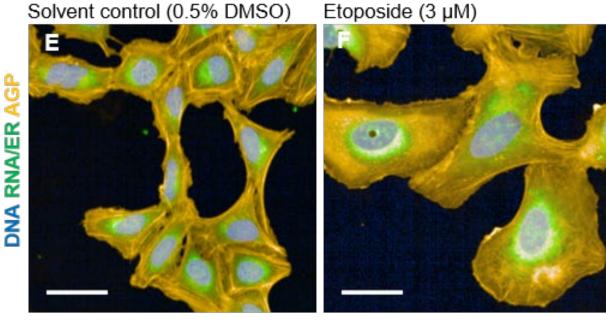
Cell Painting = Phenotypic Profiling
High-Throughput Phenotypic Profiling = HTPP



Example Chemicals: Qualitative Observation



→ Mitochondrial compactness/texture

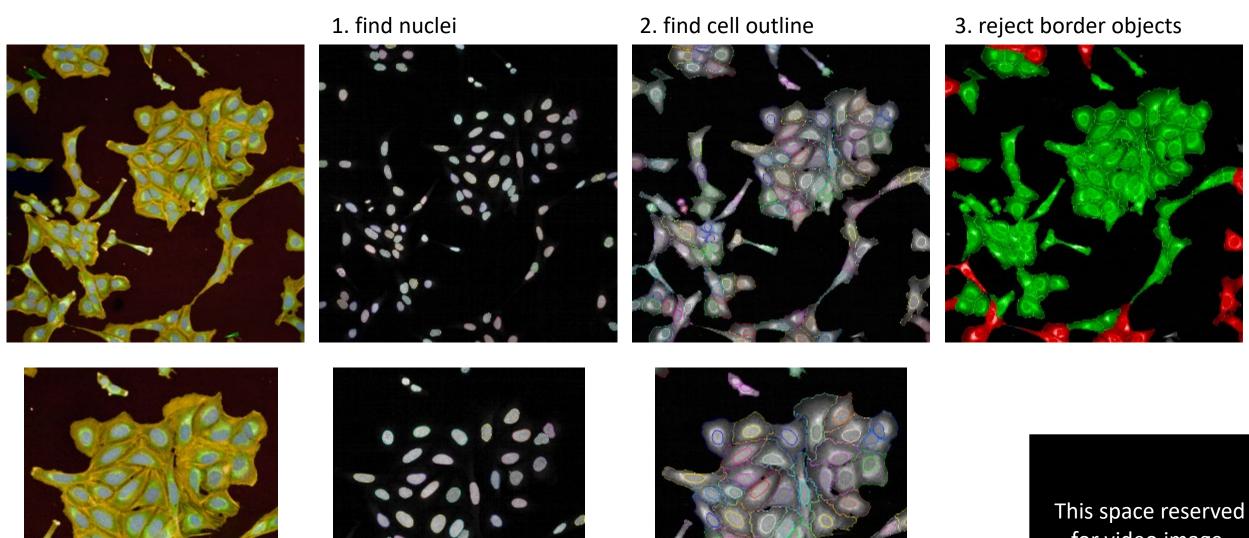


→ Cells are larger

⇒ Strong phenotypes are observable qualitatively



Image Analysis Workflow → Image Segmentation



for video image.



Define Cellular Compartments

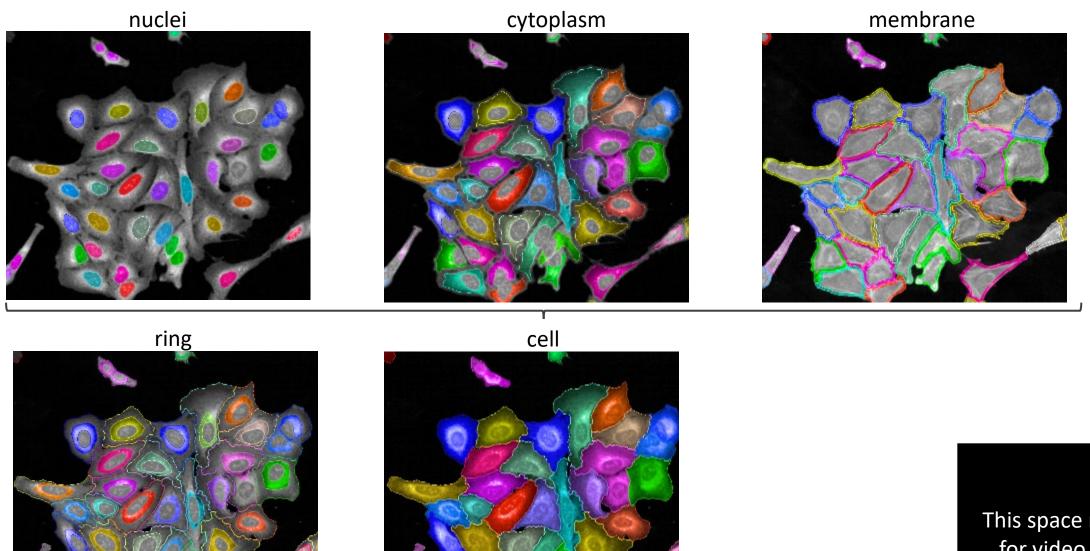


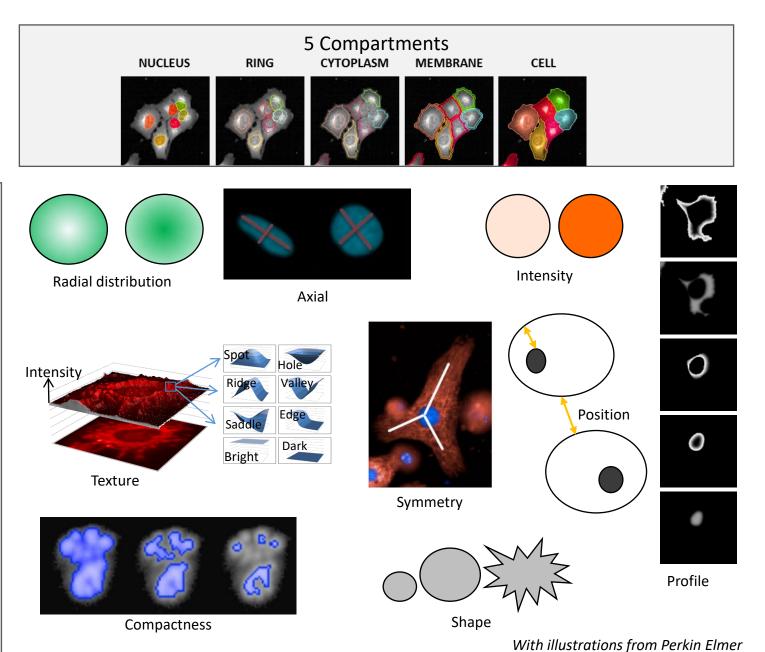


Image Processing

Profilingwith Perkin Elmer
Harmony Software

AGP

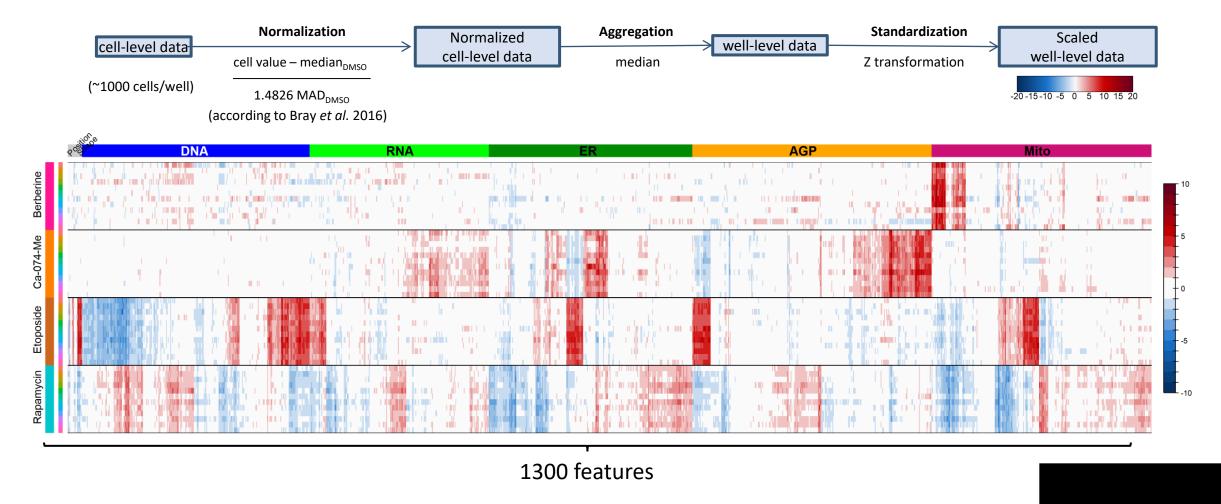
5 Channels (organelles)



= 1300 features



Example Chemicals: Quantitative Observation



Qualitative observations can be quantified



Two Applications

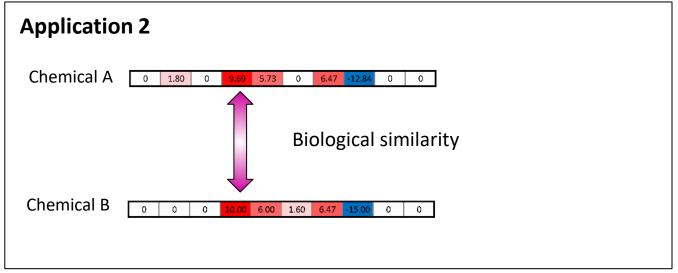




for each chemical x concentration



Application 1 concentration-response modelling concentration concentration



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



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	 TSCA Chemicals of interest to US EPA Includes 462 APCRA case study chemicals Includes 179 chemicals with annotated molecular targets 	
Concentrations:	8	3.5 log ₁₀ units; ~half-log ₁₀ spacing	
Biological Replicates:	4		



Kavlock et al. (2018) Chem. Res. Tox; 31(5): 287-290 International collaboration of regulatory scientists focused on next generation chemical risk assessment including deriving quantitative estimates of risk based on NAM-derived potency information and computational exposure estimates.

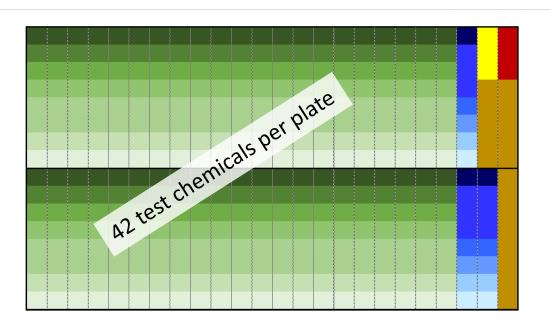
APCRA Chemicals



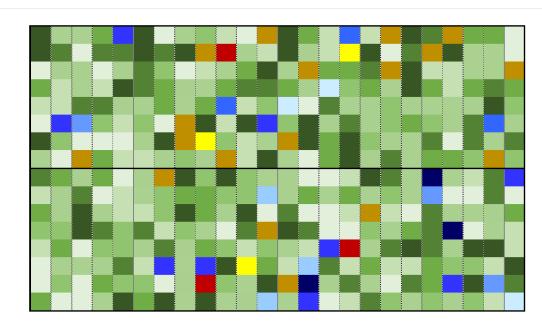
PK parameters necessary for *in vitro* to *in vivo* extrapolation (IVIVE) *in vivo* toxicity data



U-2 OS ToxCast Screen Dose Plate Design





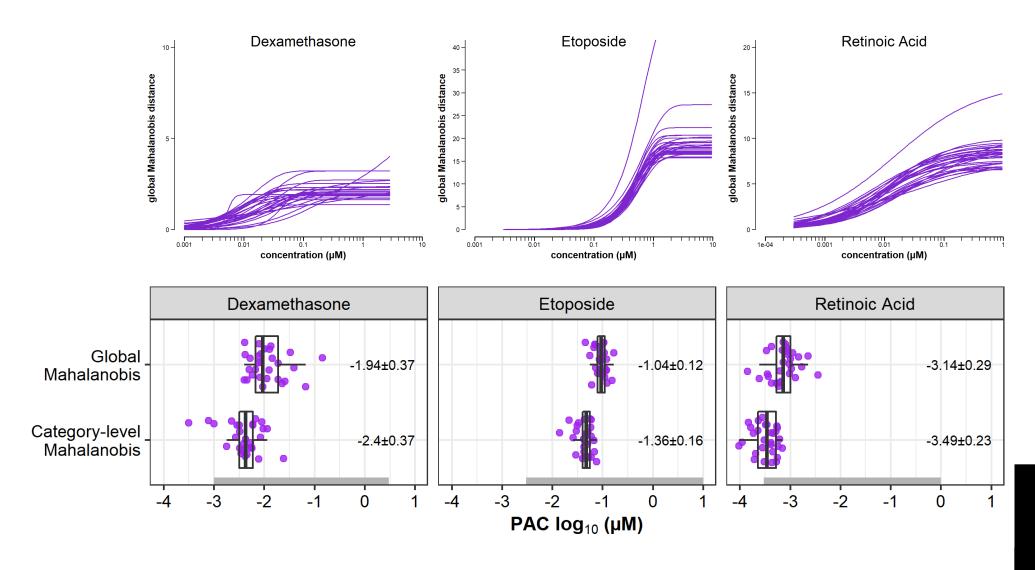


Label	Reference Chemicals:	Molecular Mechanism-of-Action	Test Concentrations
Α	Etoposide	DNA topoisomerase inhibitor	0.03 - 10 μΜ
В	all-trans-Retinoic Acid	Retinoic acid receptor agonist	0.0003 – 1 μM
С	Dexamethasone	Glucocorticoid receptor agonist	0.001 – 3 μM
D	Trichostatin A	Histone deacetylase inhibitor	1 μΜ
Е	Staurosporine	Cytotoxicity control	1 μΜ
F	DMSO	Vehicle control	0.5 %

each test plate is uniquely randomized→ no systematic edge effects



Reproducibility: Potencies

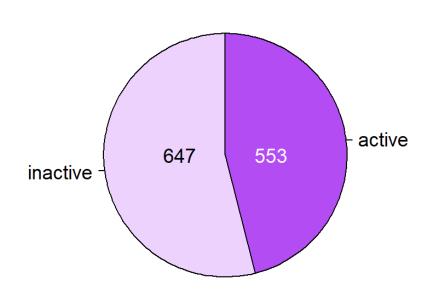


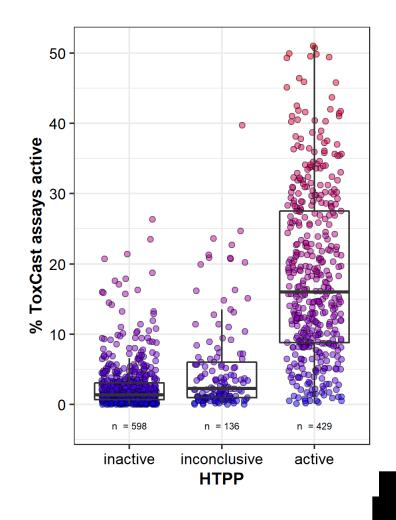
⇒ Potency estimates vary less than ½ an order of magnitude



HTPP Screening Results (1)

Active chemicals:



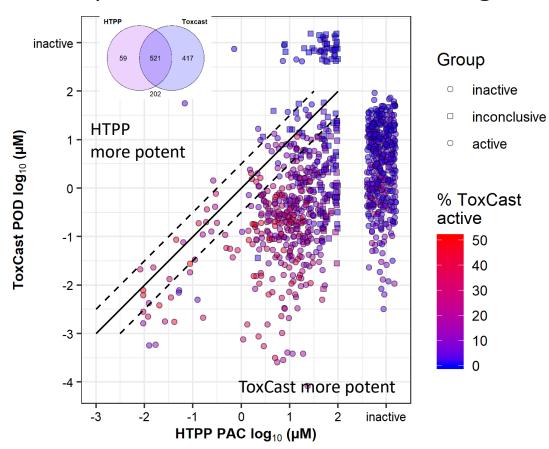


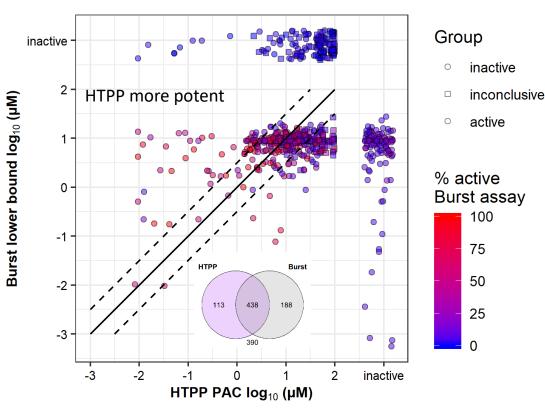
- ⇒ ~ 40% of chemicals were active
- \Rightarrow Most activity is > 10 μM
- **⇒** Chemicals active in HTPP are more often 'promiscuous' in ToxCast



HTPP Screening Results (2)

Comparison with ToxCast screening results:





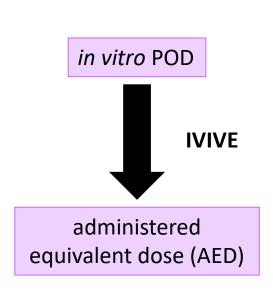
Less potent than ToxCast POD

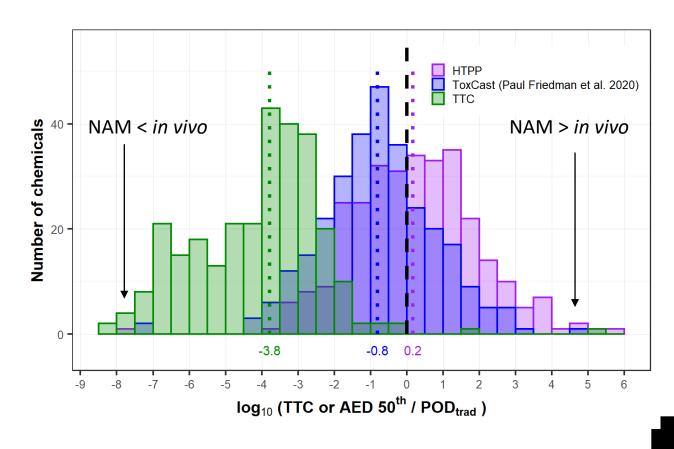
More potent than the ToxCast cytotoxicity burst estimate



Comparison to in vivo Effect Values & other NAMs

303 chemicals were active and had pharmacokinetic (PK) information

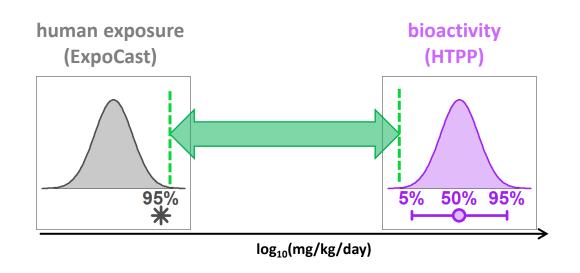




- → HTPP AEDs are higher than ToxCast-derived AEDs and TTC values
- ⇒ 78% of HTPP AED are within 2 orders of magnitude of the *in vivo* POD

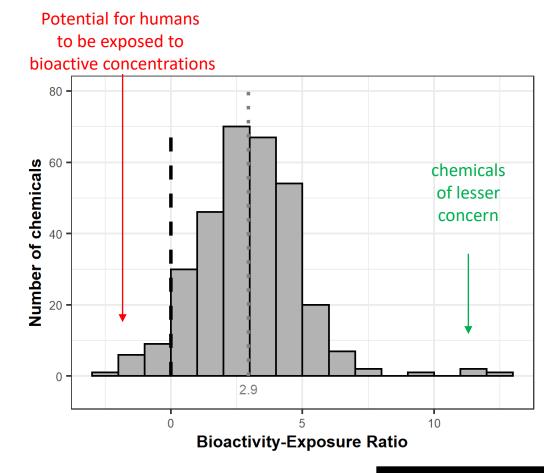


Comparison to Exposure Estimates





for a small set of chemicals, the BER was negative, indicating a potential for humans to be exposed to bioactive concentrations of these chemicals





Feature Selection & Profile Comparison

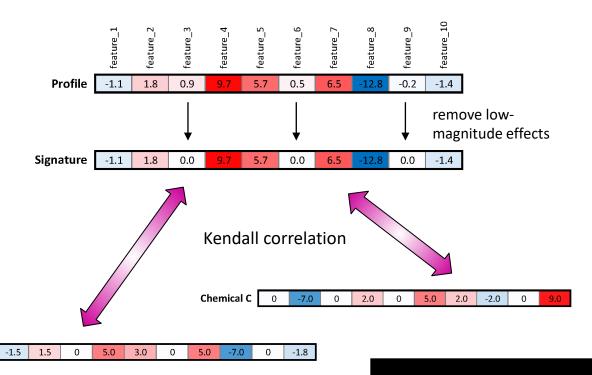
Feature Selection

1300 features

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

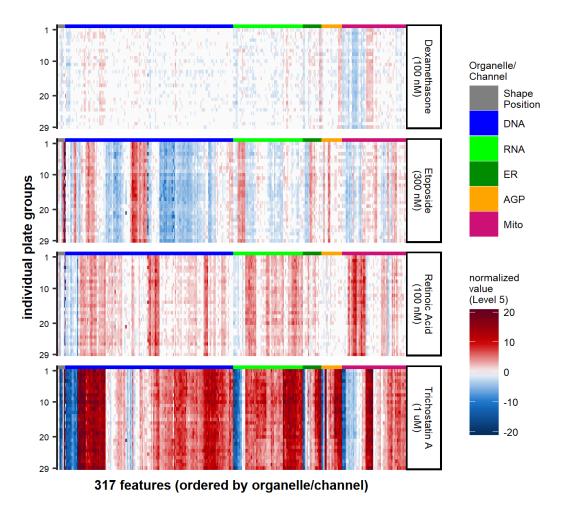
317 features

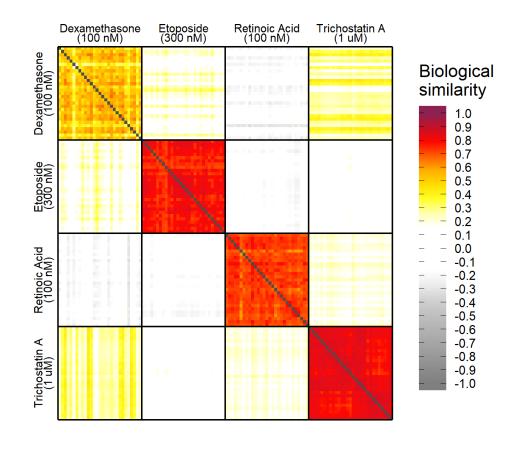
Profile Comparison





Reproducibility: Phenotypic Profiles







Phenotypic profiles are highly reproducible across different plates

Hypothesis: Chemicals with similar mechanisms will display similar profiles.



Example: Nuclear Receptor Modulators (I)

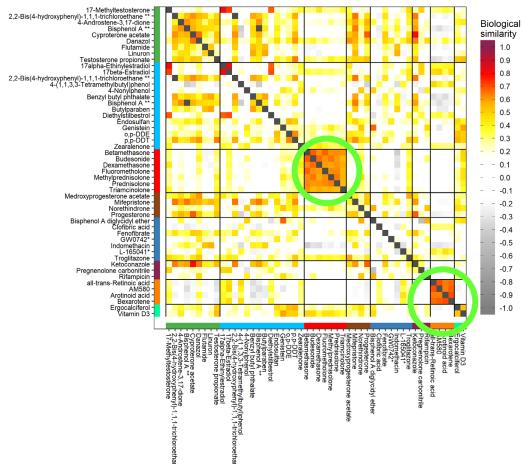
52 chemicals were annotated as targeting a nuclear receptor

Preliminary results. Do not cite or quote.

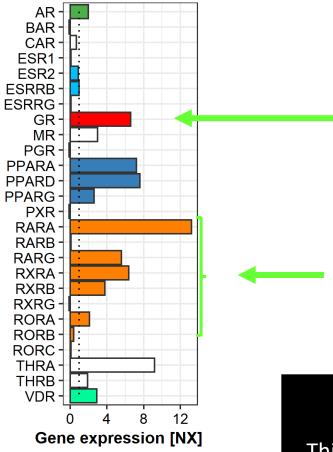
target

AR
ESR
GR
PGR
PPAR
PXR
RAR
VDR





Gene expression in U-2 OS



- ⇒ 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)



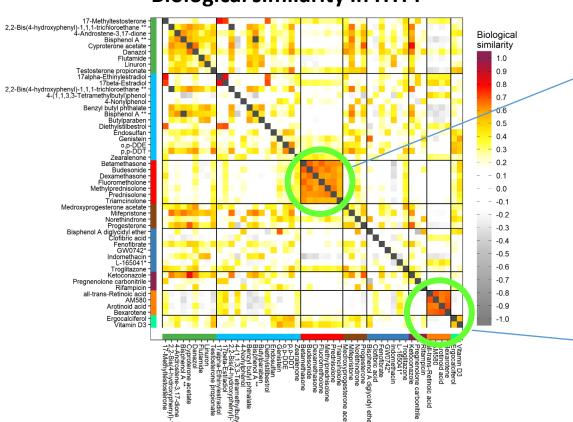
Example: Nuclear Receptor Modulators (II)

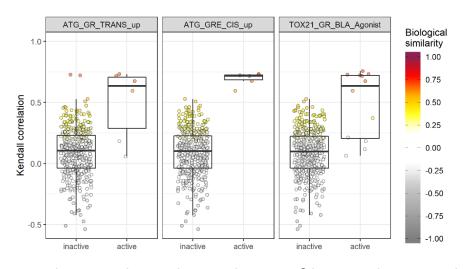




AR
ESR
GR
PGR
PPAR
PXR
RAR

VDR





Chemicals with similar profiles to dexamethasone tend to be active in **ToxCast GR assays**

Chemicals with similar profiles to all-trans retinoic acid tend to be active in **ToxCast RAR / RXR assays**

Certain molecular mechanisms result in characteristic phenotypic profiles



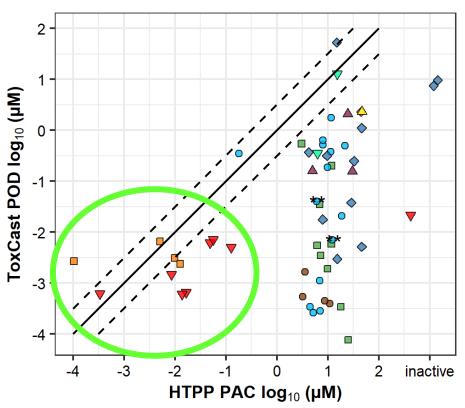
Example: Nuclear Receptor Modulators (III)

Preliminary results. Do not cite or quote.

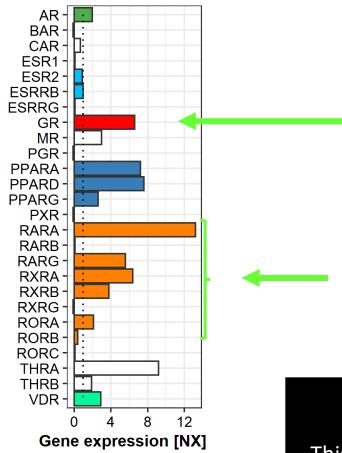
target

AR
ESR
GR
PGR
PPAR
PXR
RAR
VDR

Comparison to ToxCast potencies



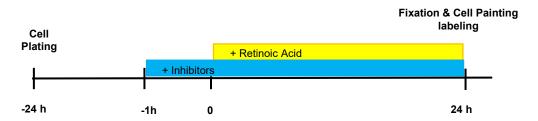
Gene expression in U-2 OS

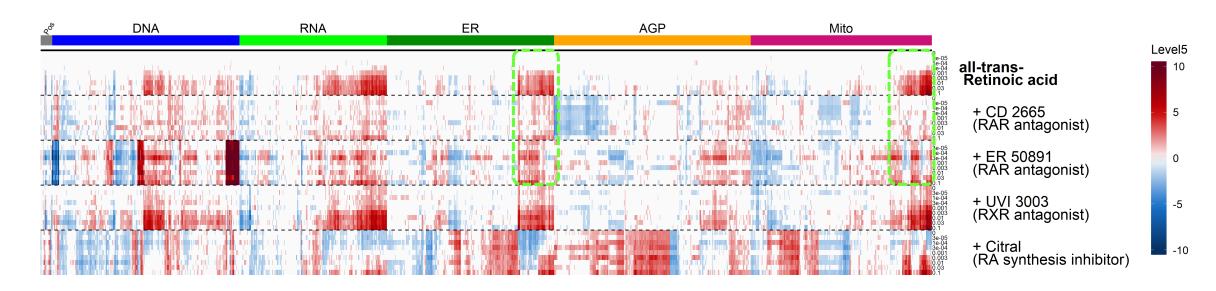


- For two receptor systems that are expressed (GR, RAR/RXR) potencies were comparable with ToxCast
- ⇒ For all other receptors, we are much less sensitive than ToxCast (off-target effects?)



Pharmacological Blockade of Phenotypic Effects

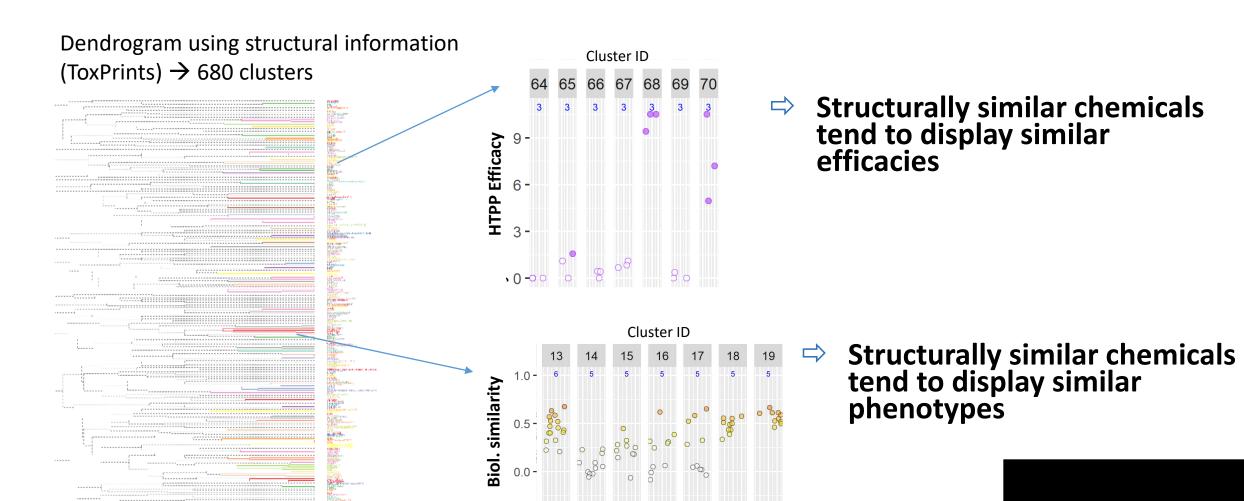




⇒ RAR but not RXR antagonists block the retinoid phenotype



Structural Similarity Translates to Biological Similarity



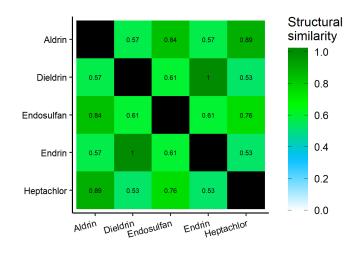
Structurally similar chemicals tend to be biologically similar

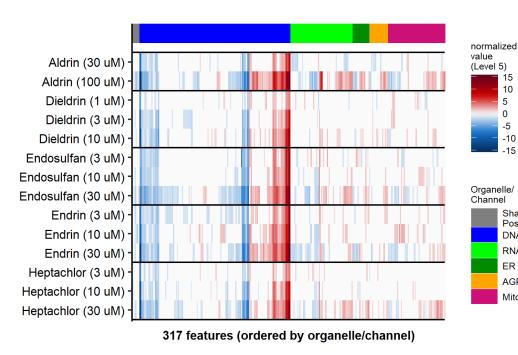


Example: Organochlorines

structural similarity

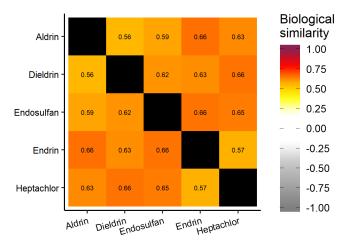
(Jaccard similarity, ToxPrints)





biological similarity

(Kendall similarity)



Organochlorines are structurally and phenotypically similar

This space reserved for video image.

10

-5

-10

Position

DNA

RNA

ER

AGP

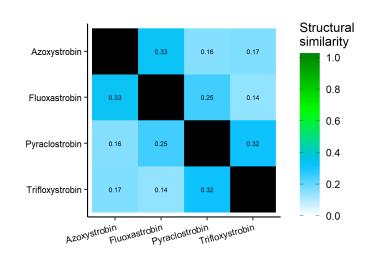
Mito

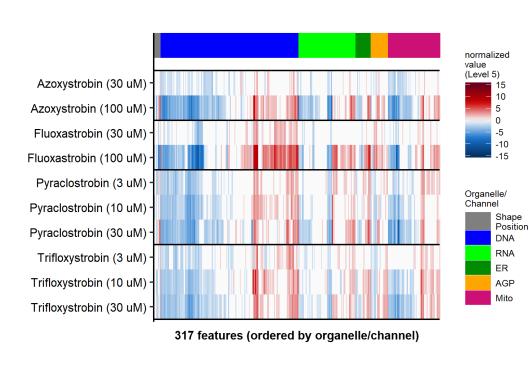


Example: Strobilurins

structural similarity

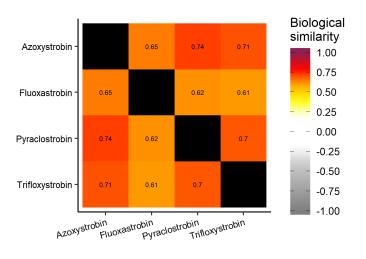
(Jaccard similarity, ToxPrints)





biological similarity

(Kendall similarity)



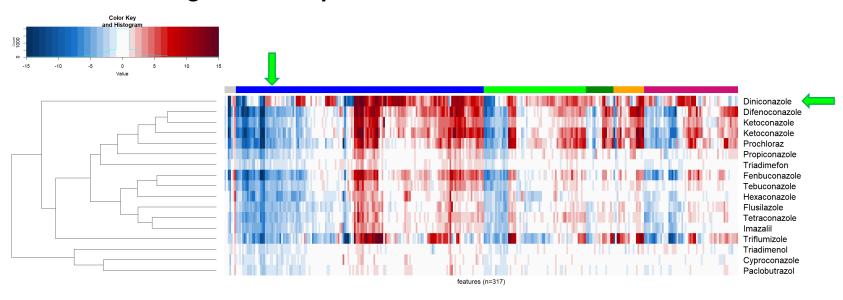
Strobilurins have less structural similarity, yet share the same molecular target and produce similar phenotypes



Example: Conazoles

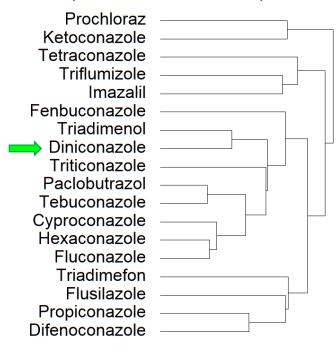
- group of fungicides
- disturb ergosterol synthesis via CYP51 and CYP61 (target absent in mammals)

biological similarity



structural similarity

(based on ToxPrints)



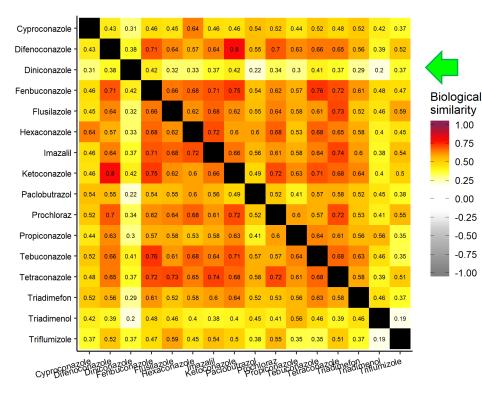
- most conazoles are phenotypically similar
- Diniconazole is phenotypically different from the other active conazoles

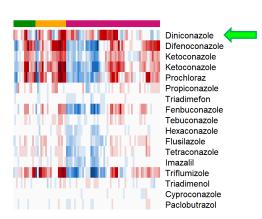


Example: Conazoles

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

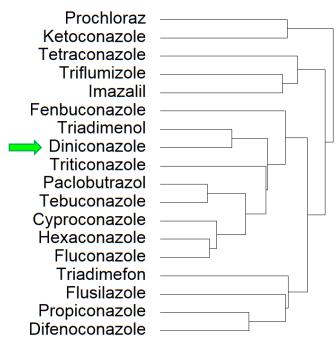
biological similarity





structural similarity

(based on ToxPrints)



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

0.75

0.50

0.25

- 0.00

-0.25

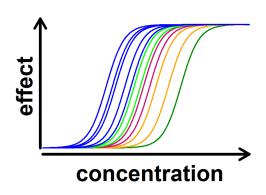
-0.50

-0.75

-1.00

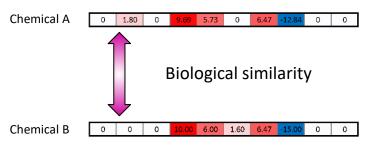


Conclusions



Application 1: Potency estimation

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



Application 2: Mechanistic prediction

- Structural similarity -> biological similarity
- Similar mechanisms → biological similarity



Outlook

Combine HTPP with HTTr

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

SOT Presentation by Dr. Joshua Harrill

"In Vitro Molecular Points-of-Departure (PODs)
from High-Throughput Profiling Assays"

Tuesday, March 23, 2021

11:15 AM-2:00 PM US Eastern Time

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



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