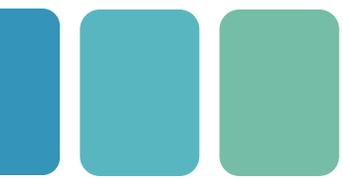


# Linking DNT In Vitro Battery Endpoints to Adverse Outcome Pathways using Omics Approaches



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Rapid Assay Development Branch  
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## Disclosure:

This work has been funded by the US. Environmental Protection Agency. I have no conflicts to declare.

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# Alignment of DNT IVB assays with AOPs



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## Neurotoxicology

journal homepage: [www.elsevier.com/locate/neuro](http://www.elsevier.com/locate/neuro)

Archives of Toxicology (2019) 93:2759–2772  
<https://doi.org/10.1007/s00204-019-02551-1>

REGULATORY TOXICOLOGY



### Development and analysis of an adverse outcome pathway network for human neurotoxicity

Nicoleta Spinu<sup>1</sup> · Anna Bal-Price<sup>2</sup> · Mark T. D. Cronin<sup>1</sup> · Steven J. Enoch<sup>1</sup> · Judith C. Madden<sup>1</sup> · Andrew P. Worth<sup>2</sup>

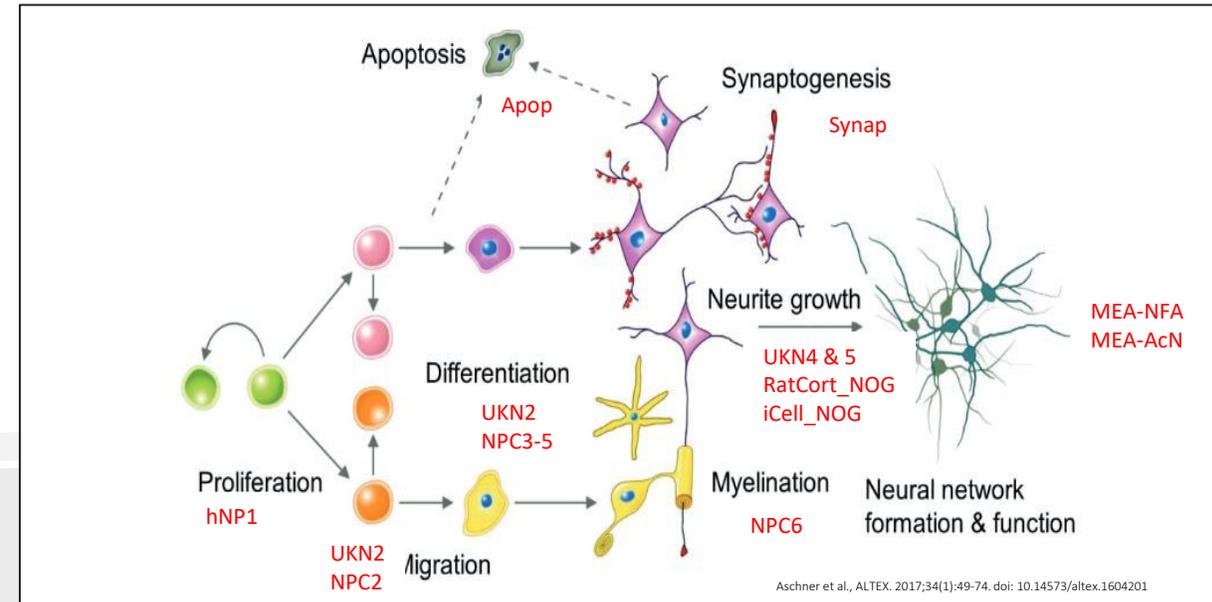
Received: 15 June 2019 / Accepted: 14 August 2019 / Published online: 23 August 2019  
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## Identification of neurotoxicology (NT)/developmental neurotoxicology (DNT) adverse outcome pathways and key event linkages with *in vitro* DNT screening assays

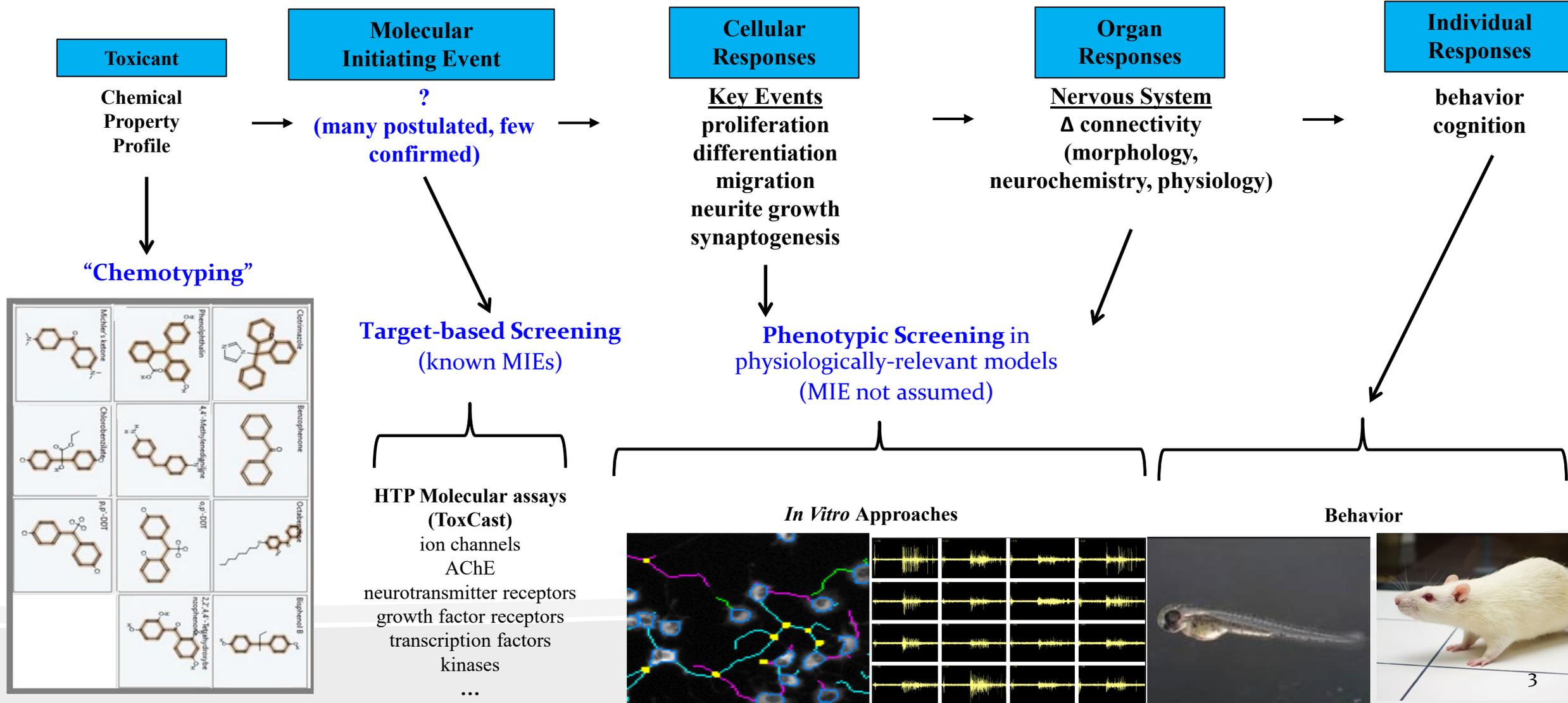
Emily M. Pitzer<sup>a,\*</sup>, Timothy J. Shafer<sup>b</sup>, David W. Herr<sup>a</sup>

<sup>a</sup> Center for Public Health and Environmental Assessment, US Environmental Protection Agency, Research Triangle Park, NC 27711, USA

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# Aligning different data streams to the AOP concept





## Linking DNT-NAMs data to AOPs



- Few endorsed AOPs exist for DNT
  - Especially if thyroid related AOPs are excluded
  - These AOPs are not necessarily driven by data from the DNT NAMs
- Is there another approach that would use data to establish linkages between data from DNT NAMs and AOPs for neurodevelopment?
  - Could this also help to establish human relevance?

# Could Omics data help link data from in vitro DNT assays to AOPs?



Toxicology and Applied Pharmacology 354 (2018) 81–93



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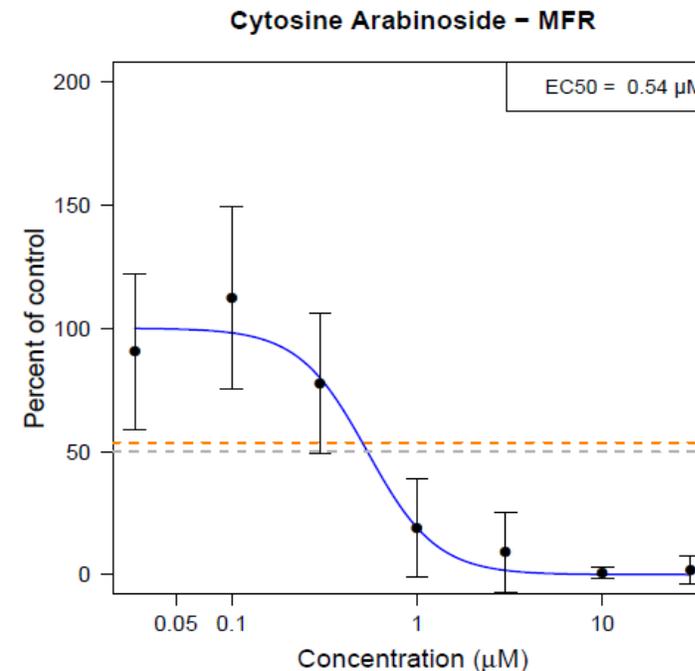
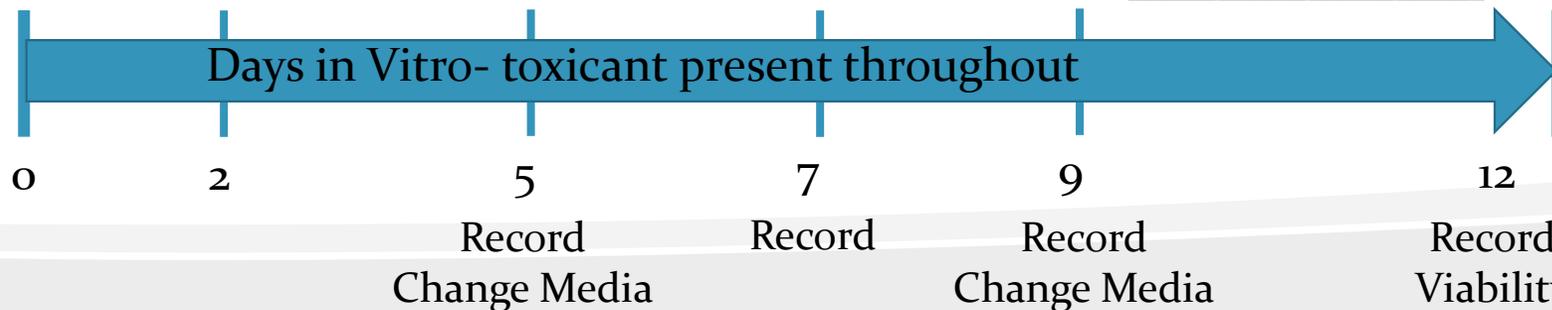
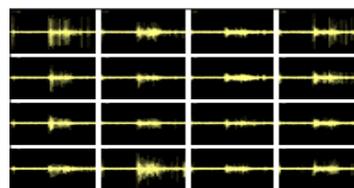
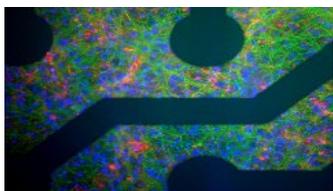
Toxicology and Applied Pharmacology

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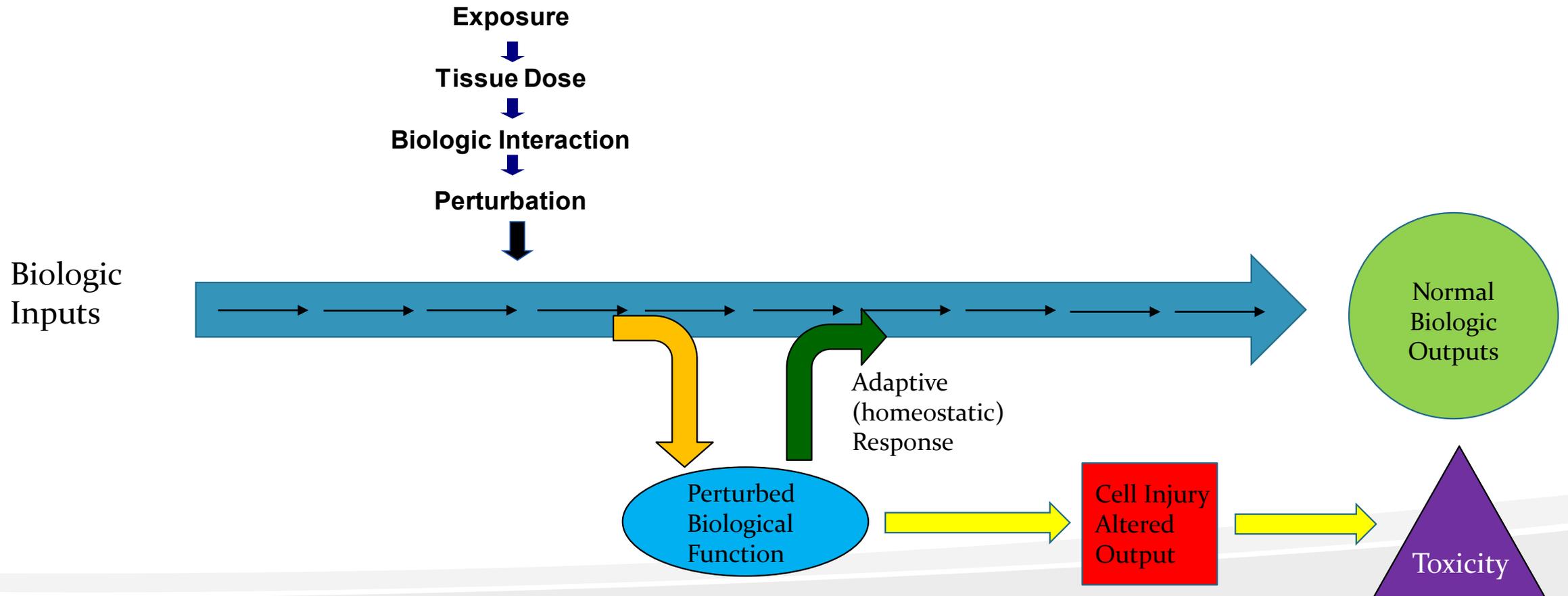
## Defining toxicological tipping points in neuronal network development<sup>☆</sup>

Christopher L. Frank<sup>a,1</sup>, Jasmine P. Brown<sup>a,2</sup>, Kathleen Wallace<sup>a</sup>, John F. Wambaugh<sup>b</sup>, Imran Shah<sup>b</sup>, Timothy J. Shafer<sup>a,\*</sup>



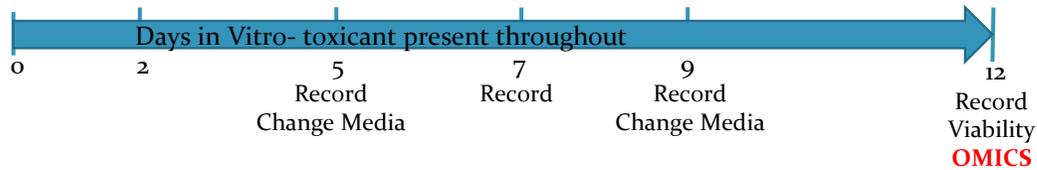
**Critical concentration ("tipping point") determined = 0.046 µM for Cytosine Arabinoside**

# The Concept of Toxicological “Tipping Points”

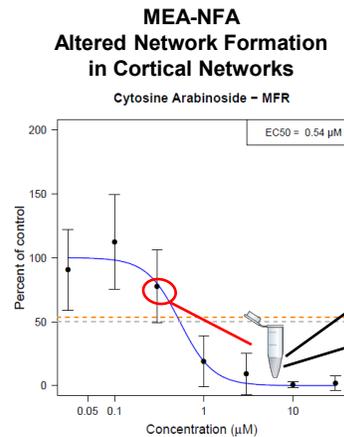
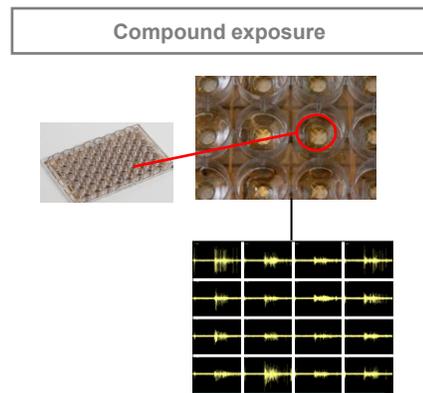


The “Tipping Point” is a mathematical determination of the point at which the system transitions from homeostasis to toxicity

# Proof-of-Concept for collecting -omics information from the rat network formation assay



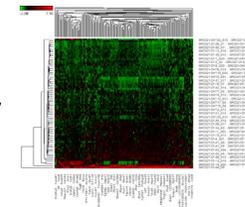
## Step 1: Chemical Dose Identification with Functional Assay



Selected Treatments

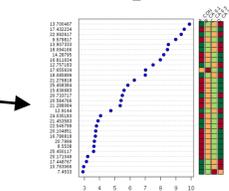
Cell Pellet  
media

## Step 2: Identifying Key Molecular Events Involved in Neurodevelopment



Transcriptomics

+



Metabolomics

## 3: Integrated Core Analysis

**Transcriptomic level**  
Disease Analysis (NDD)  
Molecular/Cellular  
Function Categories

**Metabolomic level**  
Disease Analysis (NDD)  
Molecular/Cellular  
Function Categories

**Combined Analysis**

Disease Analysis (NDD)  
Molecular/Cellular  
Function Categories

# Proof-of-Concept for collecting -omics information from the rat network formation assay



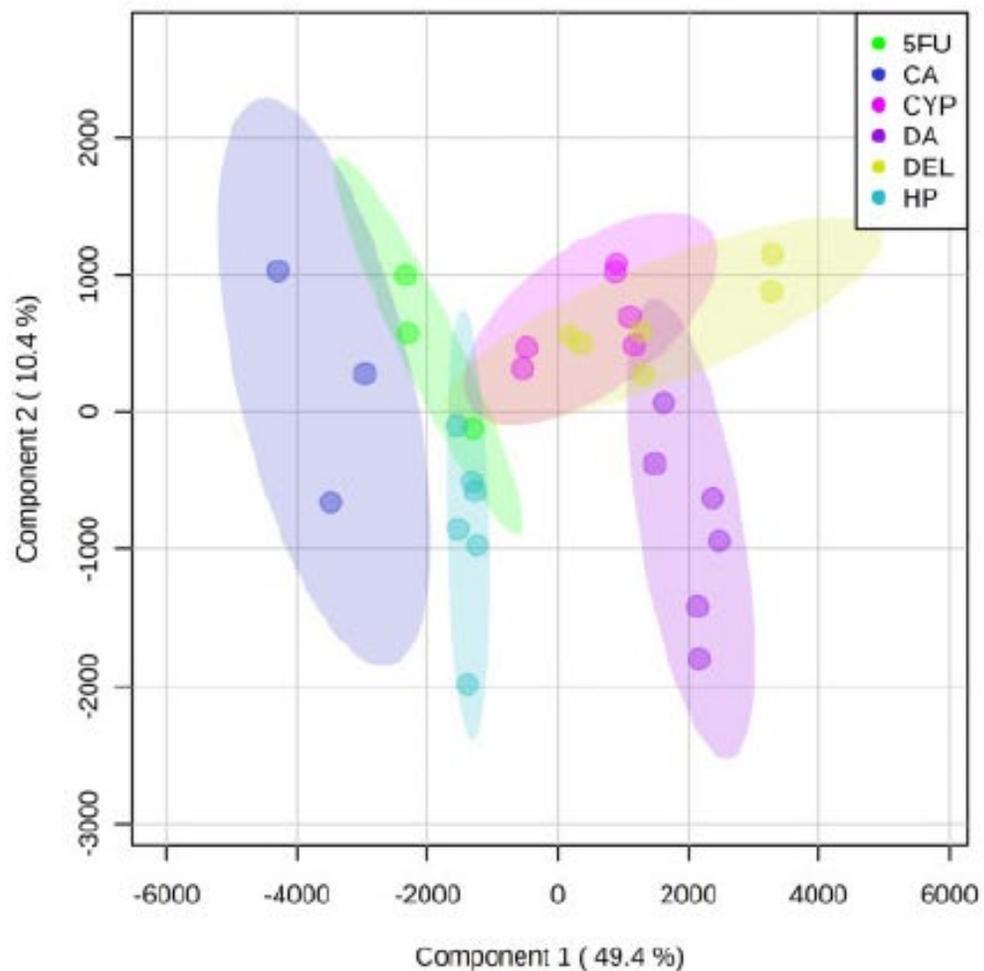
**Table 1.** Compounds Tested

Compound Name (Abbreviation)	Proposed Target	CAS No.	DTXSID	Tipping Point <sup>a</sup> ( $\mu\text{M}$ [ $\pm 95\%$ CI])	Concentrations Tested ( $\mu\text{M}$ )	Min Viability <sup>b</sup> EC <sub>50</sub> ( $\mu\text{M}$ )	Solvent Used	Source
Cytosine arabinoside (CA)	DNA synthesis inhibitor	147-94-4	DTXSID3022877	0.046 [0.04–0.06]	1	0.84	DMSO	Sigma-Aldrich
5-Fluorouracil (5FU)	DNA synthesis inhibitor	51-21-8	DTXSID2020634	0.14 [0.10–0.17]	1	2.28	DMSO	Sigma-Aldrich
Domoic acid (DA)	Glutamate Receptor Agonist	14277-97-5	DTXSID20274180	0.21 [0.9–0.25]	0.3	NA	Water	Sigma-Aldrich
Cypermethrin (CM)	Voltage-gated Sodium Channel Modulator	52315-07-8	DTXSID1023998	0.28 [0.19–0.38]	3.10	NA	DMSO	Chem Service, Inc.
Deltamethrin (DM)	Voltage-gated Sodium Channel Modulator	52918-63-5	DTXSID8020381	0.05 [0.04–0.15]	3.10	NA	DMSO	Chem Service, Inc.
Haloperidol (HP)	Dopamine Receptor Agonist	52-86-8	DTXSID4034150	0.31 [0.15–0.49]	3	13.9	DMSO	Sigma-Aldrich

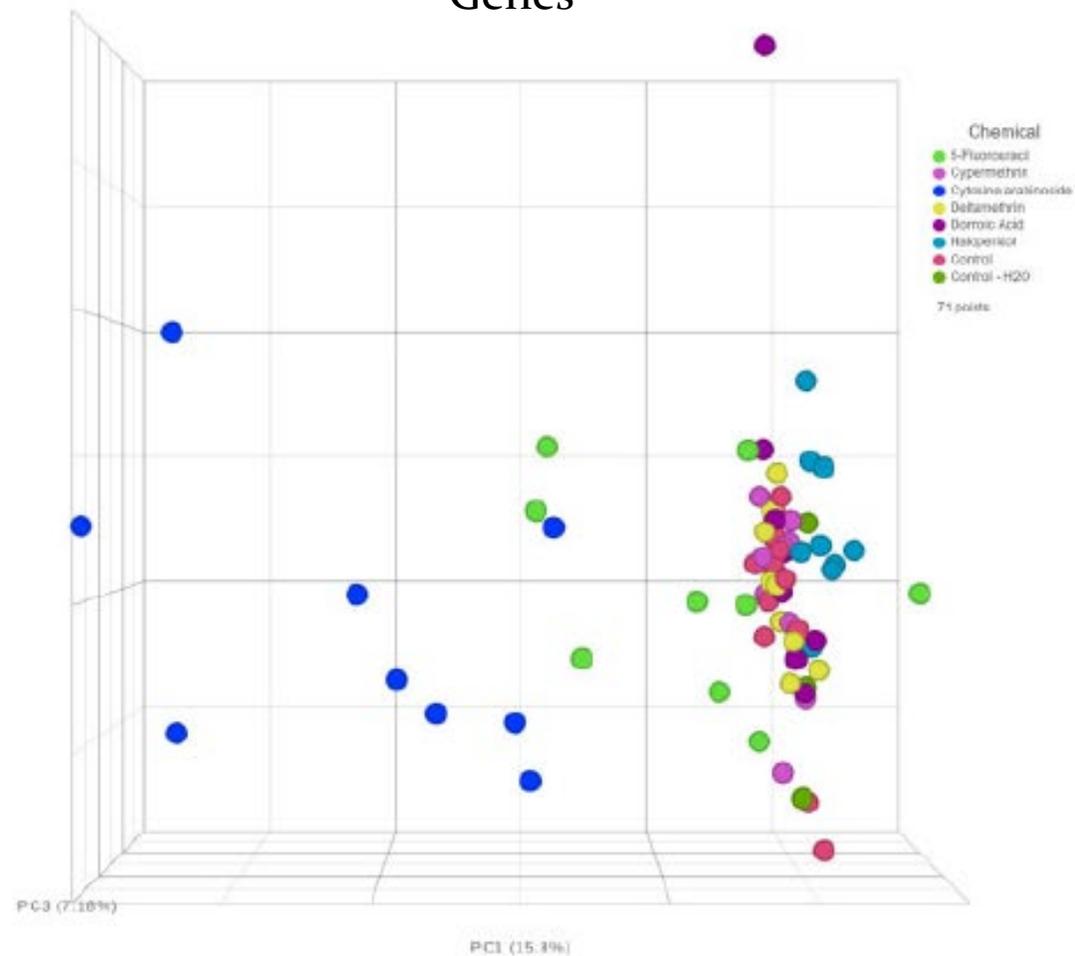
# Principal Components Analysis for Differentially Expressed Genes and Metabolites indicate treatment effects



### Metabolites



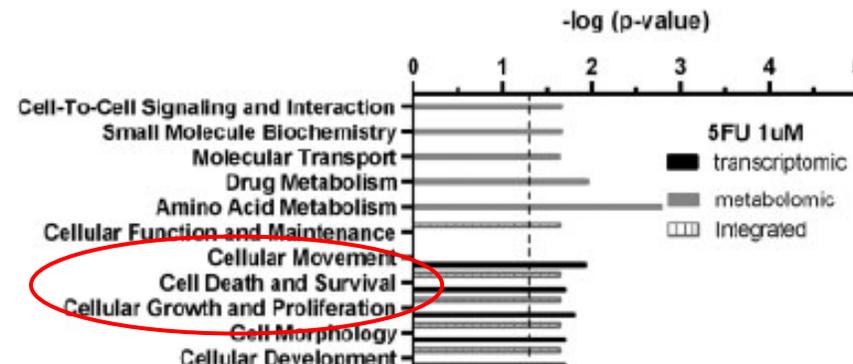
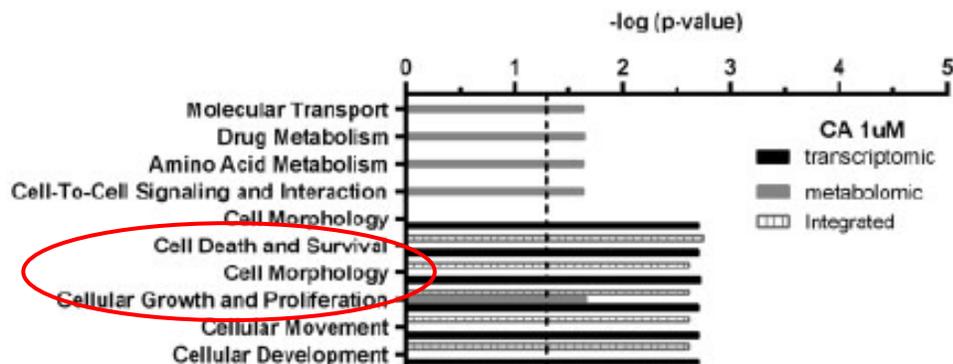
### Genes



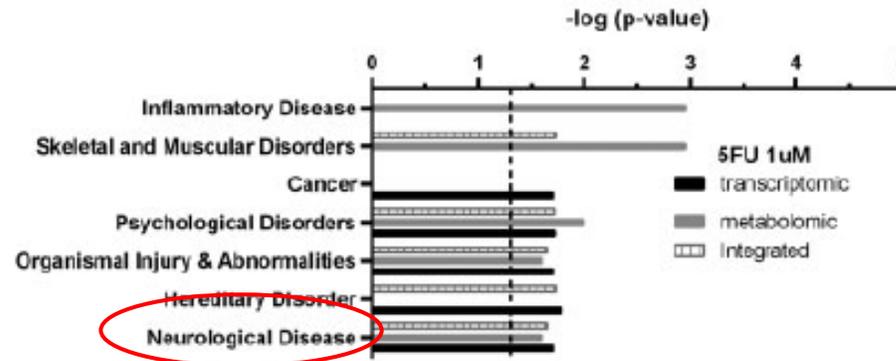
# Categories Associated With Functional Alterations in Network Development



## Cellular and Molecular Function



## Disease and Disorders



# Canonical Pathways Associated With Functional Alterations in Network Development



**Table 3.** Canonical Pathways Associated With Altered Network Formation Using Integrated Pathway Analysis

Cytosine Arabinoside (CA)	5-Fluorouracil (5FU)	Domoic Acid (DA)	Cypermethrin (CM)	Deltamethrin (DM)	Haloperidol (HP)
<ul style="list-style-type: none"> <li>• CREB signaling in neurons</li> <li>• Axonal guidance signaling</li> <li>• Osteoarthritis pathway</li> <li>• Human embryonic stem cell pluripotency</li> <li>• Hepatic fibrosis/hepatic stellate cell activation</li> </ul>	<ul style="list-style-type: none"> <li>• WNT/<math>\beta</math>-catenin signaling</li> <li>• Axonal guidance signaling</li> <li>• tRNA charging</li> <li>• Hepatocyte Growth Factor (HGF) signaling</li> <li>• Nicotinamide adenine dinucleotide (NAD) signaling pathway</li> <li>• Hepatic fibrosis/hepatic stellate cell activation</li> </ul>	<ul style="list-style-type: none"> <li>• tRNA charging</li> <li>• Phenylalanine degradation IV (mammalian, via side chain)</li> <li>• Glycine biosynthesis III</li> <li>• Purine nucleotides de novo biosynthesis II</li> <li>• Tyrosine biosynthesis IV</li> </ul>	<ul style="list-style-type: none"> <li>• Alanine biosynthesis III</li> <li>• Glycine biosynthesis III</li> <li>• Catecholamine biosynthesis</li> <li>• tRNA charging</li> <li>• Glycine Biosynthesis III</li> <li>• Thio-molybdenum cofactor biosynthesis</li> </ul>	<ul style="list-style-type: none"> <li>• (S)-reticuline biosynthesis II</li> <li>• Tyrosine degradation I</li> <li>• 4-hydroxybenzoate biosynthesis</li> <li>• 4-hydroxyphenylpyruvate biosynthesis</li> <li>• tRNA charging</li> </ul>	<ul style="list-style-type: none"> <li>• Phenylalanine degradation I</li> <li>• Threonine degradation II</li> <li>• Tyrosine biosynthesis IV</li> <li>• Glycine biosynthesis III</li> <li>• tRNA charging</li> </ul>

# Upstream Regulators Associated with Functional Alterations in Network Development

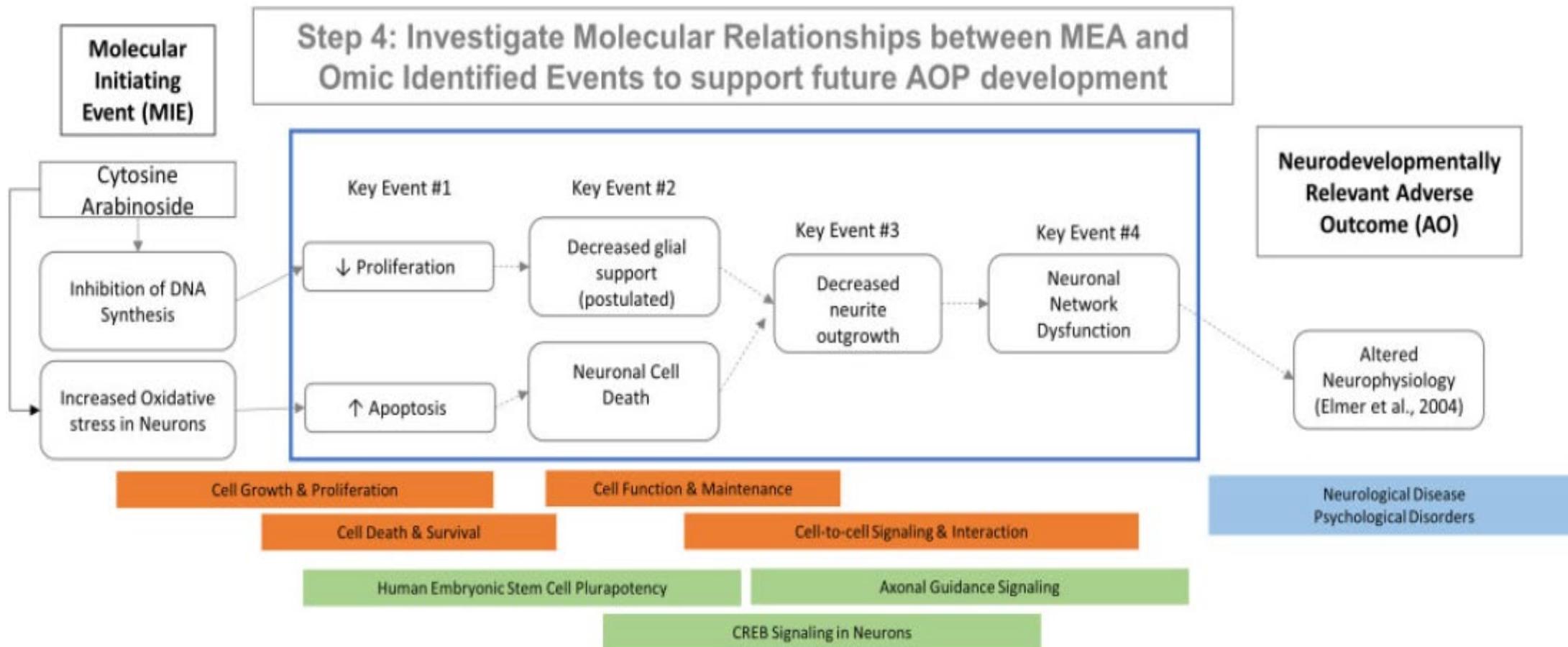


**Table 4.** Top Upstream Regulators Associated with Chemical Exposures That Alter Network Formation

	Cytosine Arabinoside (CA)	5-Fluorouracil (5FU)	Domoic Acid (DA)	Cypermethrin (CM)	Deltamethrin (DM)	Haloperidol (HP)
Transcriptomic regulators	CREB1↑	SOX2↓	LOC102724788/ PRODH	ARX	ATXN1	CREB1
	SOX2↓	TCF7L2↓	SLC1A2	LOC102724788/ PRODH	GLS	CA9
	ZBTB17	PRMT1↓	ELAVL4	SLC1A2	HPRT1	CPT1B
	S100A8↑	PPP3CA	GLS	KLF9	ELAVL4	
Metabolomic regulators	<i>β</i> -estradiol	NOTCH1	Kynurenic acid	SLC23A2	K+	Afatinb GSKJ4

Predicted activation (↑) and inhibition (↓). Lack of a directional arrow indicates that activation or inhibition could not be predicted.

# A Putative AOP based on NAMs and Omics Data



## Next Step: Confirm with additional data



Chemical	Functional Category	CAS No.	DTXSID	Tipping Point* ( $\mu\text{M}$ [ $\pm 95\%$ CI])	Concentrations Tested ( $\mu\text{M}$ ) LOW/MID/HIGH	Min AC <sub>50</sub> Endpoint*	Min AC <sub>50</sub> Viability ( $\mu\text{M}$ )*
Tebuconazole	Fungicides	107534-96-3	DTXSID9032113	1.341 [0.122-4.382]	0.3, 1, 3	0.99	20.4
Flusilazole	Fungicides	85509-19-9	DTXSID3024235	0.085 [0.043-0.291]	0.03, 0.1, 0.3	1.6	13.2
Dieldrin	GABA Modulators	60-57-1	DTXSID9020453	1.035 [0.347-2.177]	0.3, 1, 3	0.7	11.6
Heptachlor Epoxide	GABA Modulators	1024-57-3	DTXSID1024126	1.737 [1.288-1.995]	0.3, 1, 3	1.5	19.5
Lindane	GABA Modulators	58-89-9	DTXSID2020686	0.799 [0.267-4.782]	0.3, 1, 3	0.5	ND
Fipronil	GABA Modulators	120068-37-3	DTXSID4034609	0.799 [0.158-3.089]	0.3, 1, 3	15.6	25.3
Cadmium	Metals	7790-78-5	DTXSID4040183	0.036 [0.036-0.043]	0.01, 0.03, 0.1	0.04	0.48
Lead Acetate	Metals	6080-56-4	DTXSID3031521	2.916 [1.828-4.382]	1, 3, 10	0.5	16.1
Tributyltin oxide	Metals	56-35-9	DTXSID9020166	0.005 [0.004-0.006]	0.001, 0.003, 0.01	0.02	0.03
Triethyltin bromide	Metals	2767-54-6	DTXSID9040712	0.022 [0.017-0.024]	0.003, 0.01, 0.03	0.07	0.06
Sodium Arsenite	Metals	7784-46-5	DTXSID5020104	0.092 [0.055-0.122]	0.03, 0.1, 0.3	1.0	1.1
Chlorpyrifos	Organophosphates	2921-88-2	DTXSID4020458	1.23 [0.072-1.828]	0.3, 1, 3	13.1	13.7
Chlorpyrifos oxon	Organophosphates	5598-15-2	DTXSID1038666	0.071 [0.039-0.133]	0.03, 0.1, 0.3	0.4	0.3
PBDE-47	Other	5436-43-1	DTXSID3030056	1.155 [0.302-2.262]	0.3, 1, 3	0.01	11.6
Permethrin	Other	52645-53-1	DTXSID8022292	0.871 [0.64-1.288]	0.3, 1, 3	5.7	9.2
Bis(2-ethylhexyl) phthalate	Other	117-81-7	DTXSID5020607	0.11 [0.055-0.224]	0.03, 0.1, 0.3	2.0	12.3
Methylchloroisothiazolinone	Other	26172-55-4	DTXSID9034286	0.436 [0.245-0.64]	0.1, 0.3, 1	2.0	4.61
Paraquat dichloride <sup>‡</sup>	Other	1910-42-5	DTXSID7024243	0.142 [0.102-0.205]	0.03, 0.1, 0.3	0.8	1.7

\*Frank et al. (2018).

\*From ToxCast; may differ from values reported in Frank et al. (2017).

‡CAS No. and DTXSID were incorrectly reported in Frank et al 2017 and 2018.

# Treatments caused more metabolomic than transcriptomic changes



## Differentially Expressed Genes (DEGs) and Significant Metabolites at Tested Doses

Chemical	No. DEGs			No. Metabolites		
	LOW	MID	HIGH	LOW	MID	HIGH
Tebuconazole	1	51	923	101	82	350
Flusilazole	-	-	-	133	72	49
Dieldrin	-	-	-	107	137	65
Heptachlor Epoxide	-	-	-	149	54	139
Lindane	-	-	-	557	202	86
Fipronil	96	125	208	139	75	135
Cadmium	-	-	8	114	432	62
Lead Acetate	96	221	895	148	295	392
Tributyltin oxide	-	13	103	148	157	117
Triethyltin bromide	-	-	-	585	399	244
Sodium Arsenate	-	-	9	48	81	415
Chlorpyrifos	-	1	-	65	75	48
Chlorpyrifos oxon	-	-	-	44	79	68
PBDE-47	-	-	-	41	96	125
Permethrin	-	-	32	78	285	125
Bis(2-ethylhexyl) phthalate	-	-	-	243	137	113
Methylchloroisothiazolinone	-	-	249	215	384	194
Paraquat	-	-	1	57	93	114

Many fewer DEGs observed here. Therefore, focus on compounds with concentration-dependent changes in DEGs.



# Top Diseases and Functions Associated With Functional Alterations in Network Development



## Tebuconazole

- Cell Death and Survival
- Neurological Disease
- Organismal Injury and Abnormalities
- Cellular Development
- Nervous System Development and Function

## Lead Acetate

- Cell Morphology,
- Cellular Assembly and Organization
- Molecular Transport
- Cell-To-Cell Signaling and Interaction
- Nervous System Development and Function
- Neurological Disease
- Psychological Disorders

## Tributyltin

- Amino Acid Metabolism
- Cell-To-Cell Signaling and Interaction
- Cell Death and Survival
- Cellular Growth and Proliferation
- Neurological Disease
- Nervous System Development and Function

## Fipronil

- Cell Morphology
- Cell Development
- Tissue Morphology
- Neurological Disease
- Nervous System Development and Function,



# Top Upstream Regulators



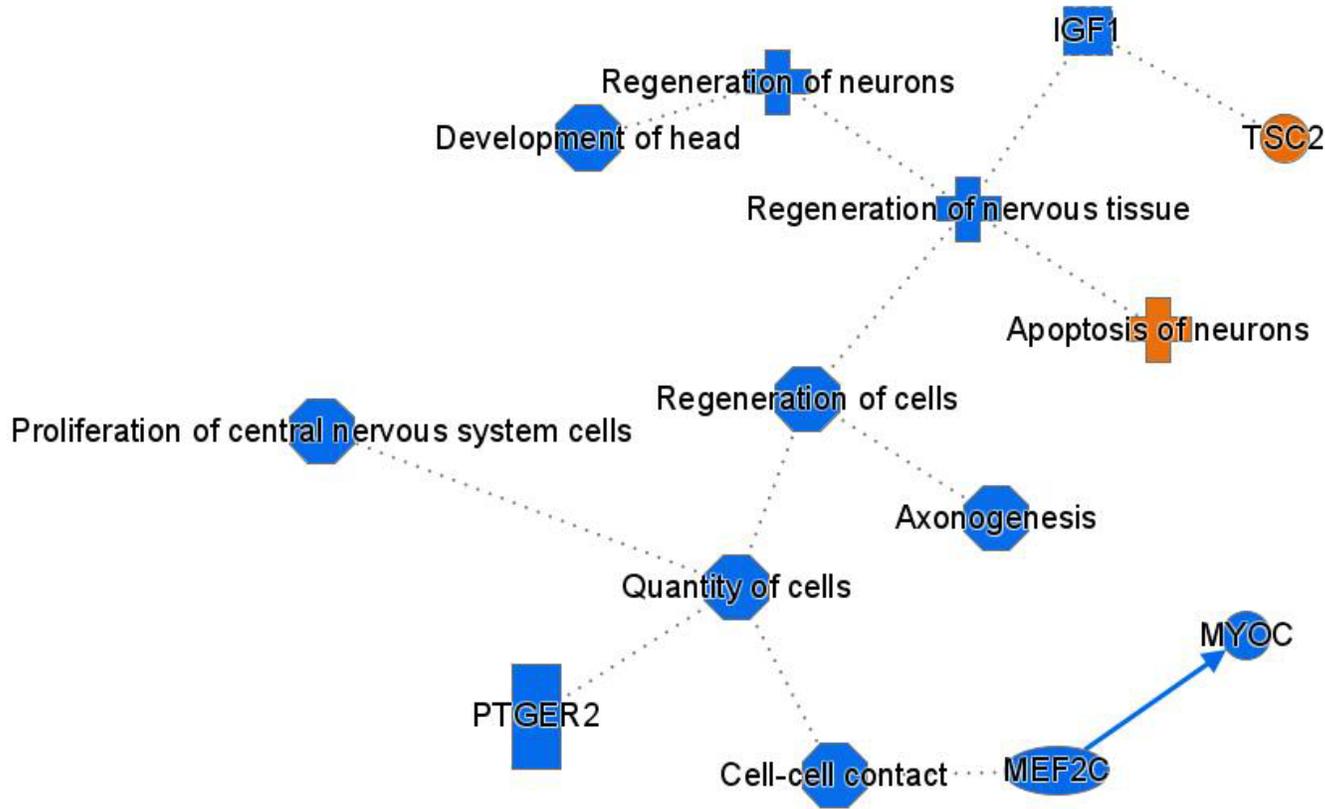
Cytosine Arabinoside (CA)	5-Fluorouracil (5FU)	Domoic acid (DA)	Cypermethrin (CM)	Deltamethrin (DM)	Haloperidol (HP)	Lead Acetate	Tributyltin Chloride	Fipronil	Tebuconazole
<p>CREB1↑</p> <p>SOX2↓</p> <p>ZBTB17</p> <p>S100A8↑</p>	<p>SOX2↓</p> <p>TCF7L2↓</p> <p>PRMT1↓</p> <p>PPP3CA</p> <p>NOTCH1</p>	<p>LOC102724788/</p> <p>PRODH</p> <p>SLC1A2</p> <p>ELAVL4</p> <p>GLS</p>	<p>ARX</p> <p>LOC102724788/</p> <p>PRODH</p> <p>SLC1A2</p> <p>KLF9</p> <p>SLC23A2</p>	<p>ATXN1</p> <p>GLS</p> <p>HPRT1</p> <p>ELAVL4</p>	<p>CREB1</p> <p>CA9</p> <p>CPT1B</p>	<p>SOX2↓</p> <p>KDM1A</p> <p>PPP3CA</p> <p>inosine↓</p> <p>MYOC↓</p> <p>MAPT</p> <p>CX3CL1</p> <p>HTT</p> <p>APP↓</p>	<p>MAPT</p> <p>PSEN1</p> <p>APP</p> <p>HTT</p> <p>BDNF↑</p> <p>MKNK1↑</p> <p>ATN1</p> <p>FMR1↓</p> <p>GRN</p> <p>CYP27A1</p>	<p>SOX2↑</p> <p>BDNF</p> <p>CHMP2B</p> <p>CREB1</p> <p>HTT</p> <p>GRN</p> <p>SERPINF1</p> <p>CSF1</p>	<p>TARDBP↑</p> <p>PROM1</p> <p>SH3TC2↑</p> <p>LMNB1</p> <p>SIRT2↑</p> <p>SOX2↑</p> <p>NR1H3</p> <p>NR1H2</p> <p>CYP27A1↑</p>

Our analysis indicates that there are a number of common upstream regulators across the active chemicals. However, there is also considerable heterogeneity in the identified upstream regulators.

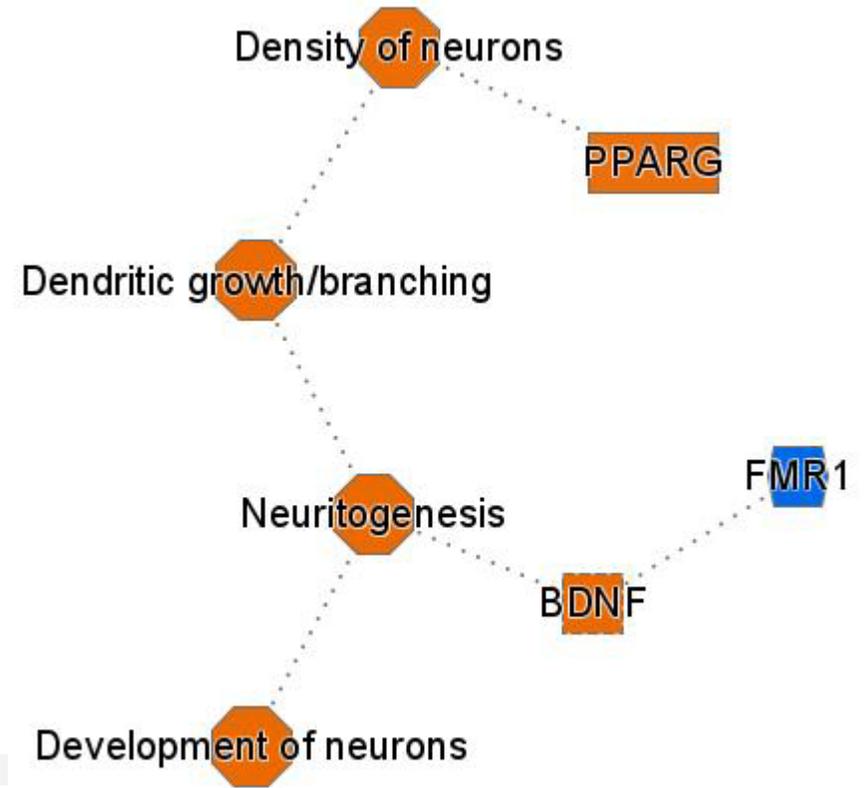
# Analysis could inform AOPs



Lead- Mid dose



Tributyltin- Mid dose





## Conclusions



- To date, there are few AOPs focused on DNT Adverse Outcomes
  - Especially if thyroid-related AOPs are considered (excluded)
- The Omics approaches identify common diseases, molecular functions, upstream regulators and canonical pathways associated with positive responses in the DNT NAMs
  - Data for more chemicals are needed to confirm this
  - If common regulators and pathways are identified, this might result in novel DNT NAMs
- Omics approaches also identify unique responses to each chemical at multiple levels of organization.
- Using Omics approaches and NAMs data, putative AOPs can be identified
  - These putative AOPs need development, but provide a link between DNT NAMs data, key events and adverse outcomes related to human disease



Thank you!  
Questions?

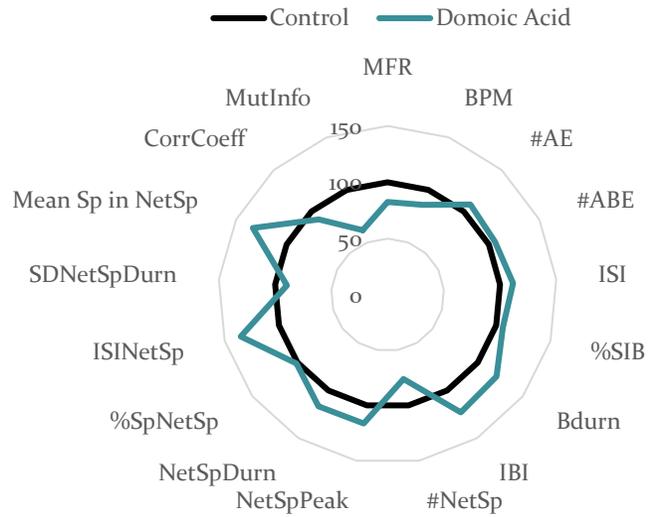


**EPA ORD Colleagues:**

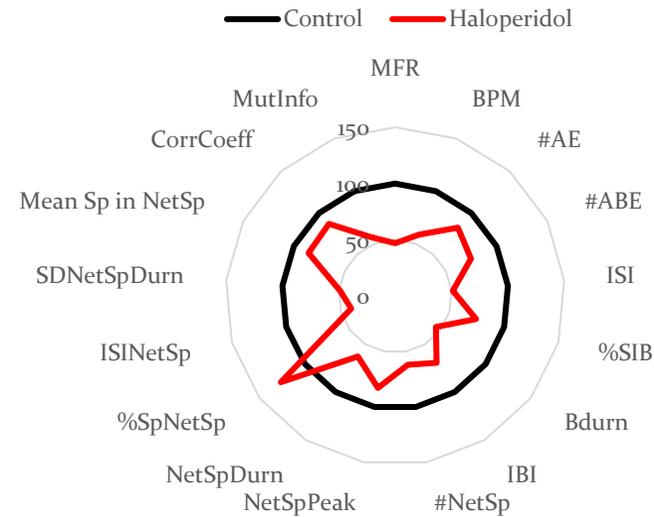
- Kathleen Wallace
- Theresa Freudenrich
- Brian Chorley
- David Gallegos
- Carmen Marable
- Chris Frank
- Matthew Henderson
- Susan Hester
- Roland Seim



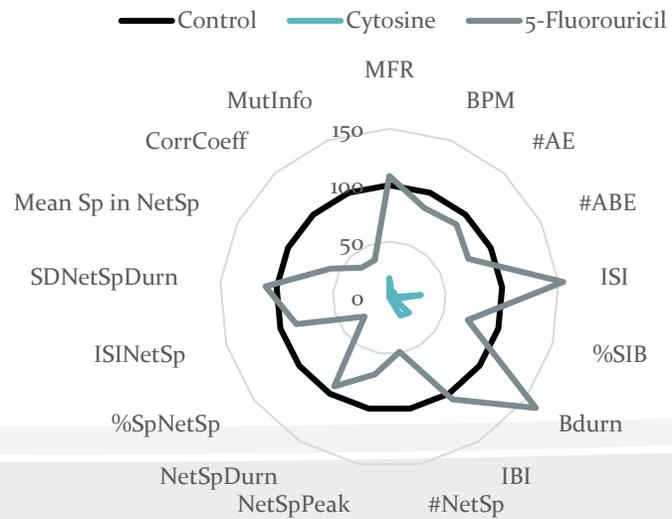
### Domoic Acid



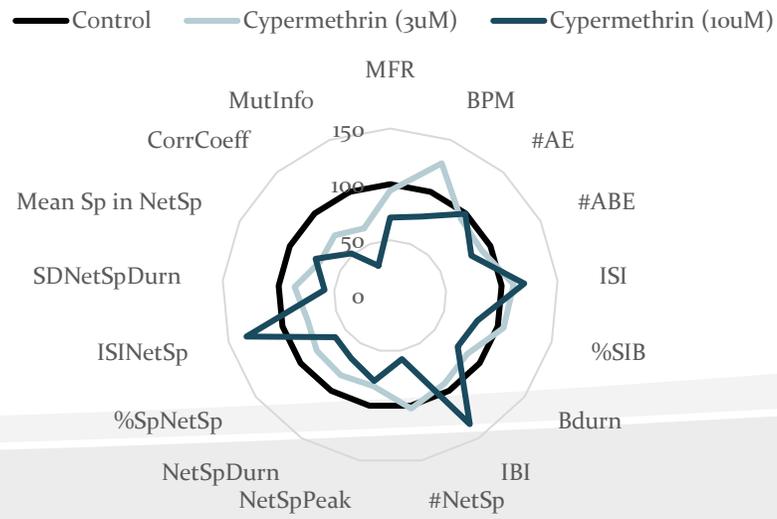
### Haloperidol



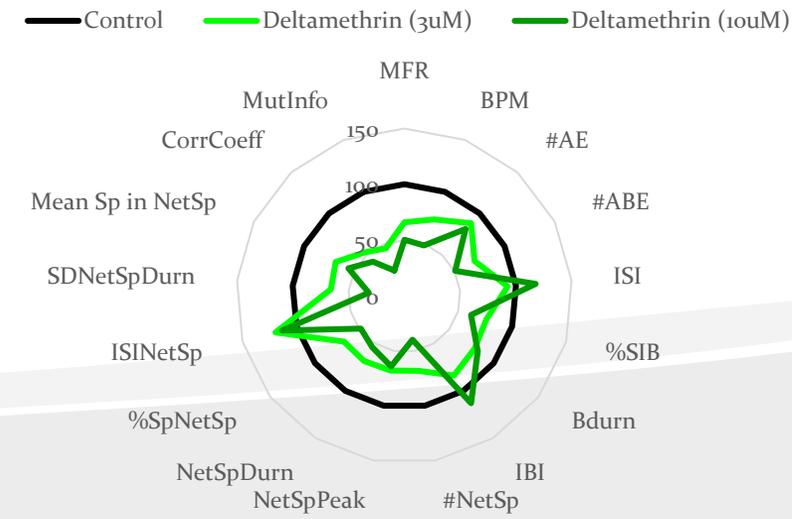
### DNA Synthesis Inhibitors



### Cypermethrin



### Deltamethrin



# Functional alterations in the MEA NFA

- Fungicides
- GABA-A Receptor
- Metals
- Parent/metabolite

