



Computational approaches to integrate DNT NAMs for fit-for-purpose identification of DNT hazard

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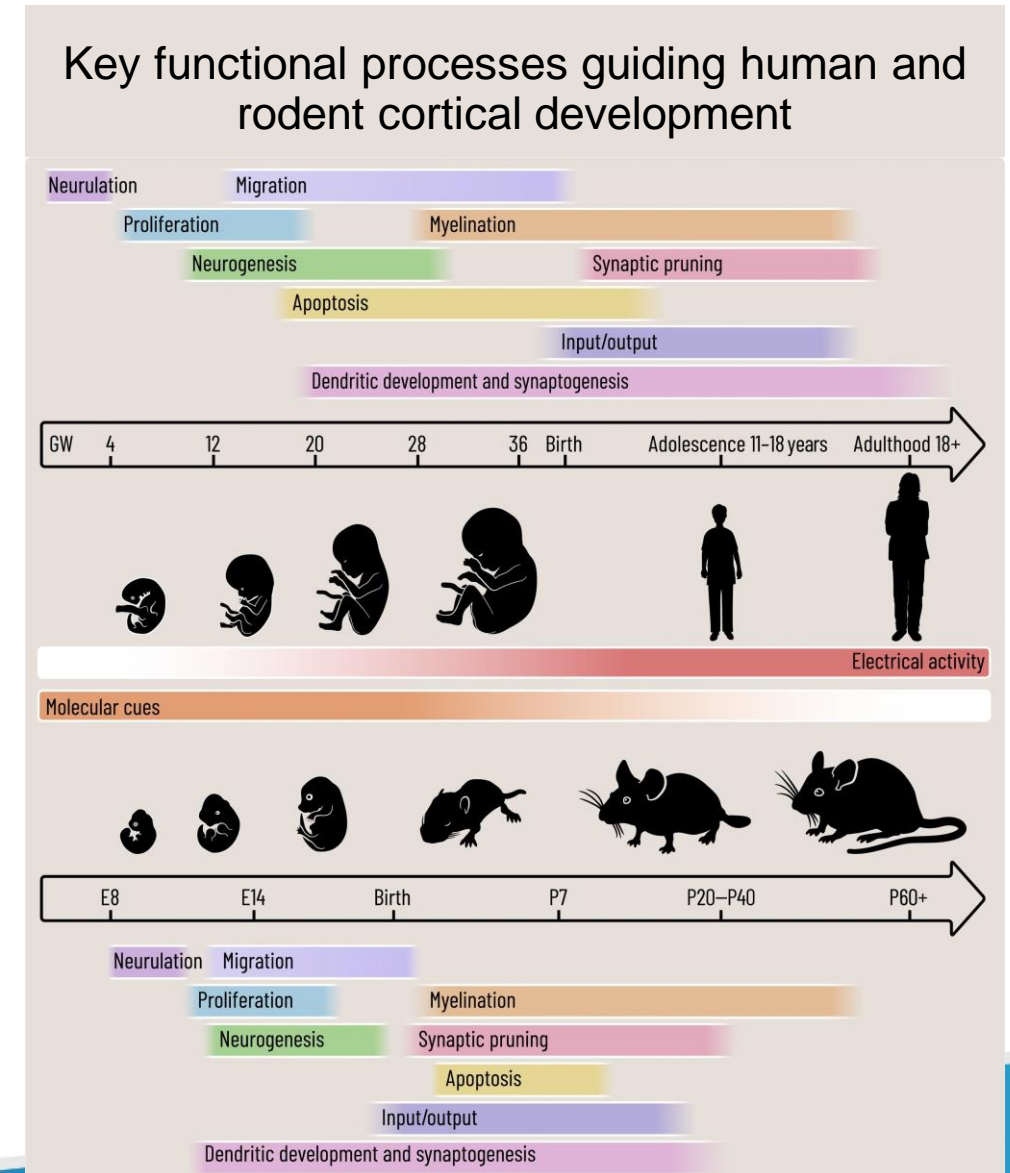
Office of Research and Development
Center for Computational Toxicology and Exposure
Biomolecular and Computational Toxicology Division
Computational Toxicology and Bioinformatics Branch

Conflict of Interest Statement

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

Background on developmental neurotoxicity (DNT) new approach methods (NAMs)

- ❖ Neurodevelopmental disability is the most prevalent chronic medical condition encountered in pediatrics (Zablotsky et al. 2019).
- ❖ Both **genetic and environmental risk factors** have been identified as underlying causes driving this prevalence.
- ❖ DNT NAMs battery: multi-dimensional DNT screening assays that cover complex neurobiological space: temporal, different 'key events' in neurodevelopment, cell-types, and species.
- ❖ Challenges in evaluating DNT NAMs:
 - No single *in vitro* screening assay can recapitulate all critical cellular events of neurodevelopment.
 - Some compounds may disrupt specific cellular events at different stages of development.
 - Some neural cell-types may be differentially sensitive to perturbation.



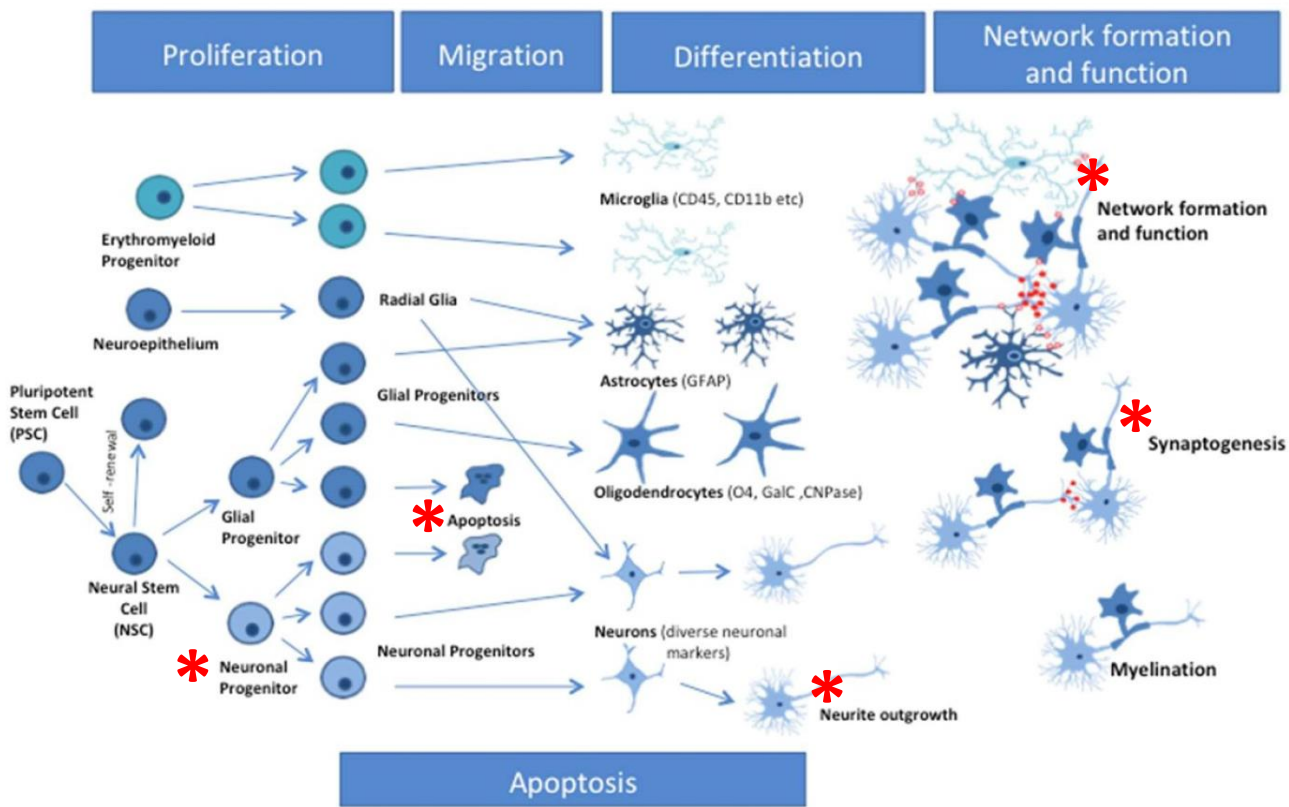
Chini and Hanganu-Opatz. 2021. Trends in Neuro.

Overview

DNT NAMs: multi-dimensional DNT screening assays that cover complex neurobiological space: multiple 'key events' in neurodevelopment, different cell-types, developmental timepoints, and species.

- ❖ How does a broad screening battery collectively inform DNT-relevant bioactivity?
- ❖ Can we build a model to classify compounds that demonstrate *in vivo* DNT bioactivity?
- ❖ Can we identify biological gaps in the current EPA DNT NAM battery and/or broader ToxCast/ Tox21 database?

Neurodevelopmental processes in the EPA DNT NAM battery



Bal-price et al. 2018

Table 2. Proposed Assays for Evaluation As an In Vitro DNT Battery

Process	Assays	References
* Proliferation	→ hNP1 NPC1	Harrill et al. (2018) Baumann et al. (2016) and Barenys et al. (2017)
* Apoptosis	UKN1	Balmer et al. (2012)
Migration	→ hNP1 NPC2	Harrill et al. (2018) Baumann et al. (2016) and Barenys et al. (2017)
Neuron differentiation	UKN2 NPC3	Nyffeler et al. (2017) Baumann et al. (2016) and Barenys et al. (2017)
Oligodendrocyte differentiation & maturation	NPC5/6	Baumann et al. (2016) and Barenys et al. (2017)
* Neurite outgrowth	→ iCell gluta (hN2) UKN 4 & 5 (rat) NPC4	Harrill et al. (2018) Krug et al. (2013) Baumann et al. (2016) and Barenys et al. (2017)
* Synaptogenesis	→ Rat primary synaptogenesis	Harrill et al. (2018)
* Network formation	→ MEA-NFA (rat)	Brown et al. (2016) and Frank et al. (2018)

Sachana, M., et.al. 2019, Toxicological Sciences

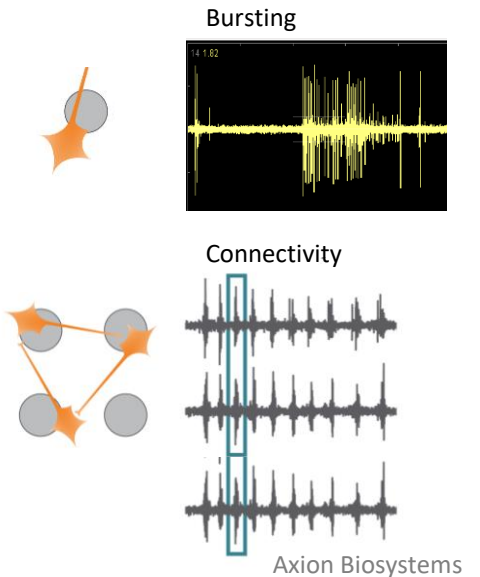
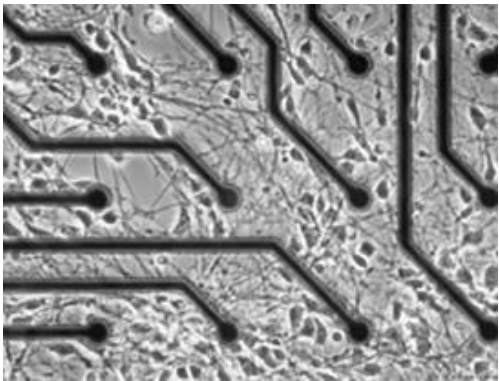
Experimental models in the EPA DNT NAM battery

Microelectrode Array (MEA) Network Formation Assay (NFA)

← 92 chemicals →

High Content Imaging

48-well culture plate
16 electrodes per well



Cell culture	Activity type	# endpoints
Primary rat cortical neurons (DIV 5, 7, 9, 12)	↓↑ General activity	4
	↓↑ Network connectivity	8
	↓↑ Bursting	5
	Cytotoxicity	2

96-well culture plate
Immunohistochemistry

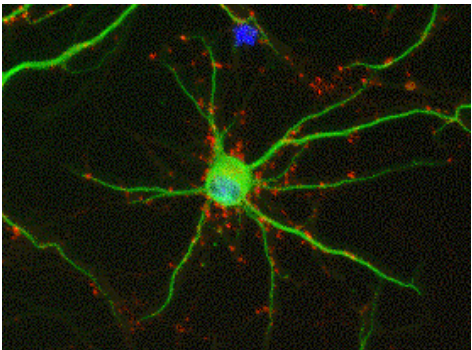
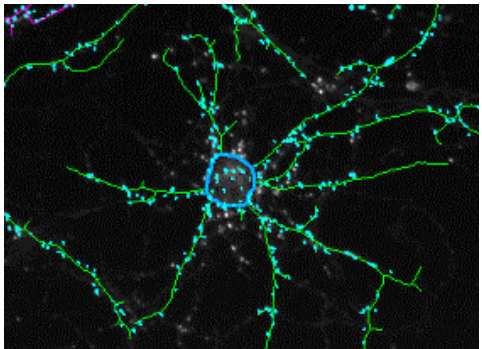


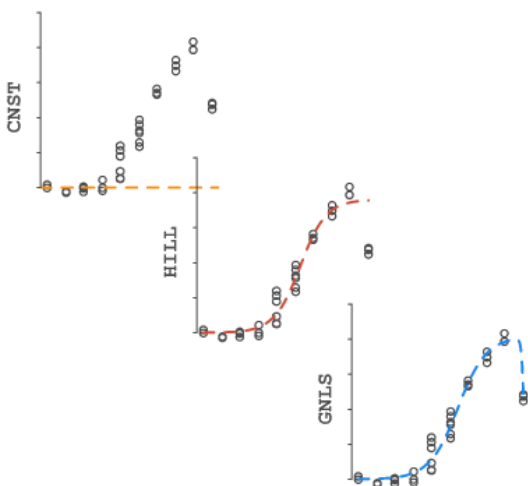
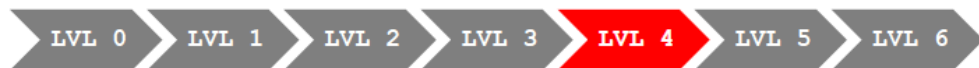
Image Analysis



Cell culture	Assays/ Key events	# endpoints
Primary rat cortical neurons	Neurite Outgrowth (NOG)	4
	Synaptogenesis and Neurite maturation	8
Human hN2 neural cells	NOG	4
Human hNP1 neuroprogenitors	Proliferation	3
	Apoptosis	2

Defining bioactivity using the ToxCast pipeline

Model fitting (constant, hill, gain-loss)



$$\mu_i = 0$$

$$\mu_i = \frac{1}{1 + 10^{(ga - x_i)gw}}$$

$$g_i = \frac{1}{1 + 10^{(ga - x_i)gw}}$$

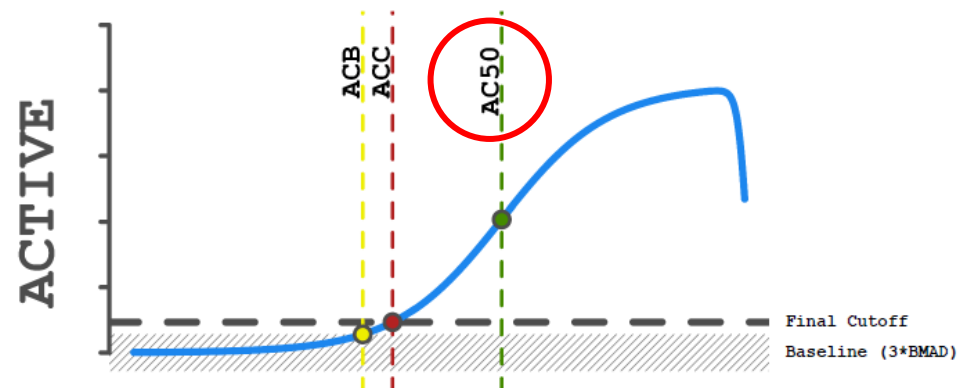
$$l_i = \frac{1}{1 + 10^{(x_i - la)lw}}$$

$$\mu_i = tp * g_i * l_i$$

Select winning model and hit-calling



Point of departure estimates:

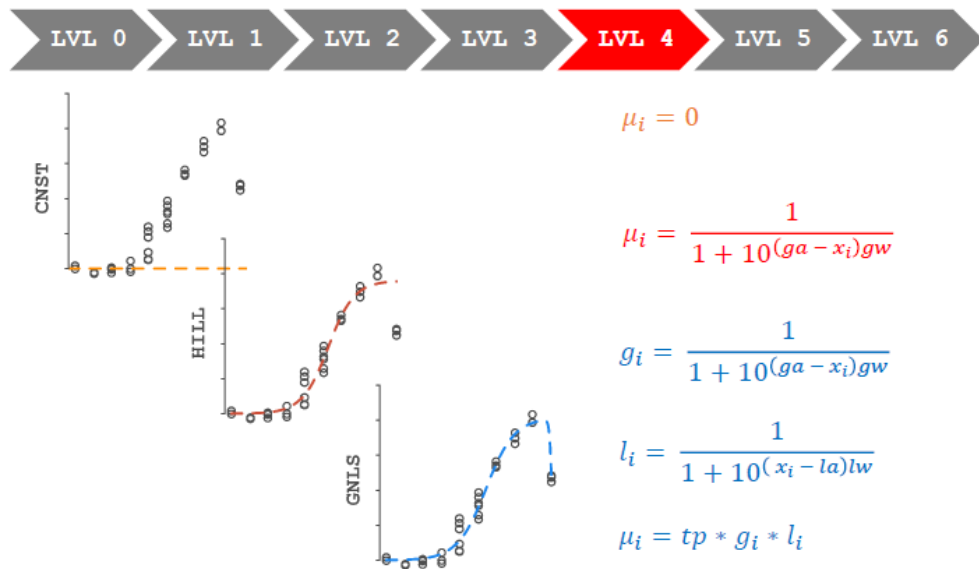


ToxCast pipeline (tcpl) R package (version 2.0.3 [publicly available](#))
(Filer et al. 2017)

https://cran.r-project.org/web/packages/tcpl/vignettes/Data_processing.html#level-4

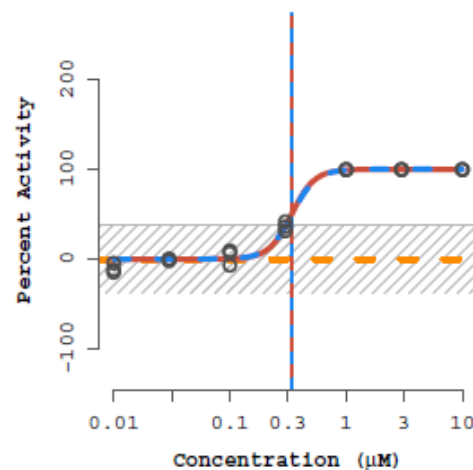
Defining bioactivity using the ToxCast pipeline

Model fitting (constant, hill, gain-loss)



Select winning model and hit-calling

CCTE_Shafer_MEA_dev_bursting_electrodes_number_dn



```
ASSAY:  ABID2500 (CCTE_Shafer_MEA_dev_bursting_electrodes_number_dn)
NAME:    Methylmercuric(II) chloride
CHID:    20813  CASRN: 115-09-3
SPID(S): EX000383
M4ID:    42619839

HILL MODEL (in red):
tp      ga      gw
val:    100    -0.469  4.43
sd:     1.28    0.0287  2.18

GAIN-LOSS MODEL (in blue):
tp      ga      gw      la      lw
val:    101    -0.465  4.21    1.22    10.2
sd:     NaN    NaN    NaN    NaN    NaN

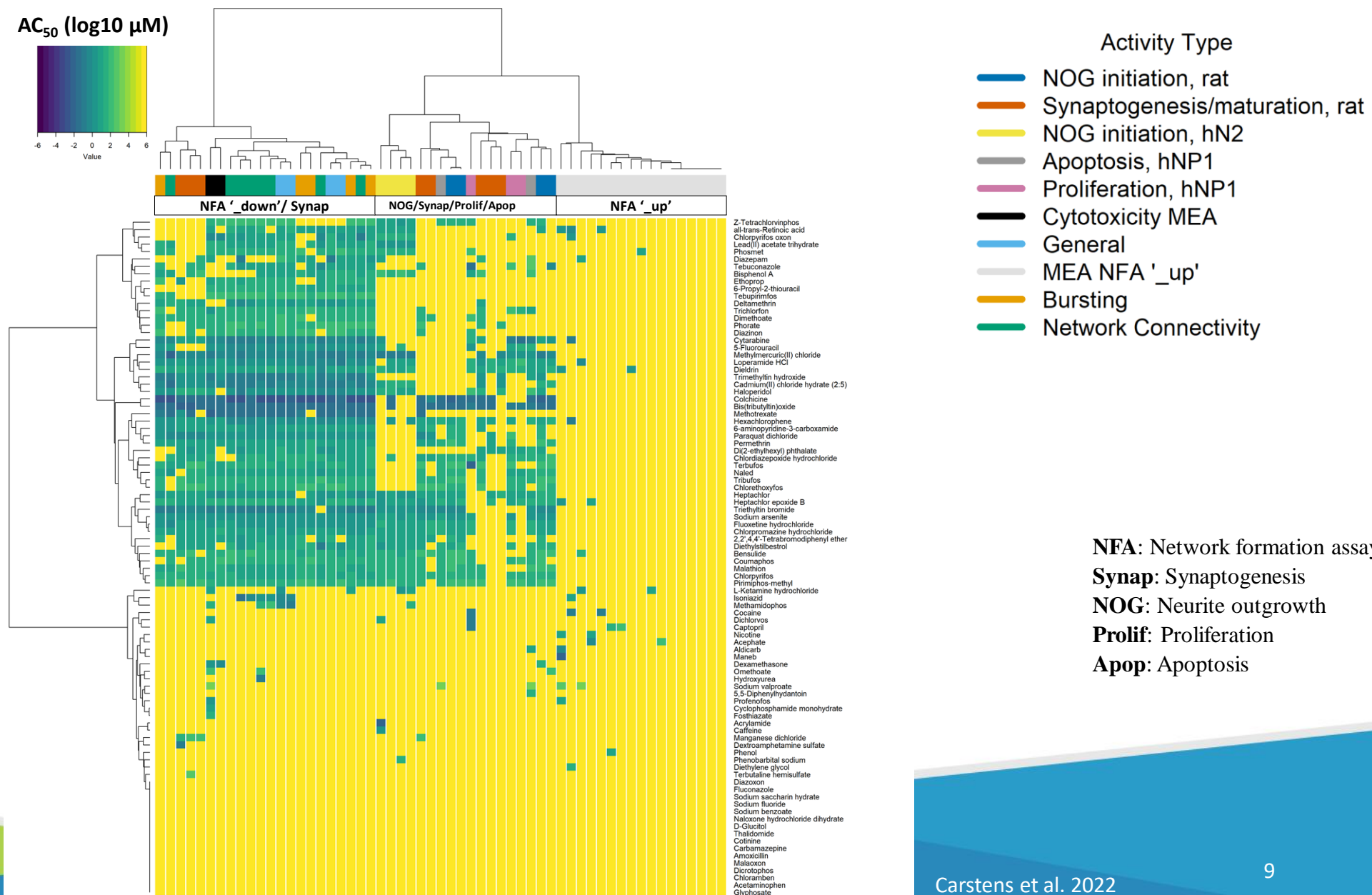
CNST      HILL      GNLS
AIC:  240.39  135.21  139.17
PROB:  0      0.88   0.12
RMSE:  67.13  5.4    5.39

MAX_MEAN: 100      MAX_MED: 100      BMAD: 12.8
COFF:  38.3  HIT-CALL: 1  FITC: 41  ACTP: 1
FLAGS:
```

ToxCast pipeline (tcpl) R package (version 2.0.3 [publicly available](https://cran.r-project.org/web/packages/tcpl/vignettes/Data_processing.html#level-4))
(Filer et al. 2017)

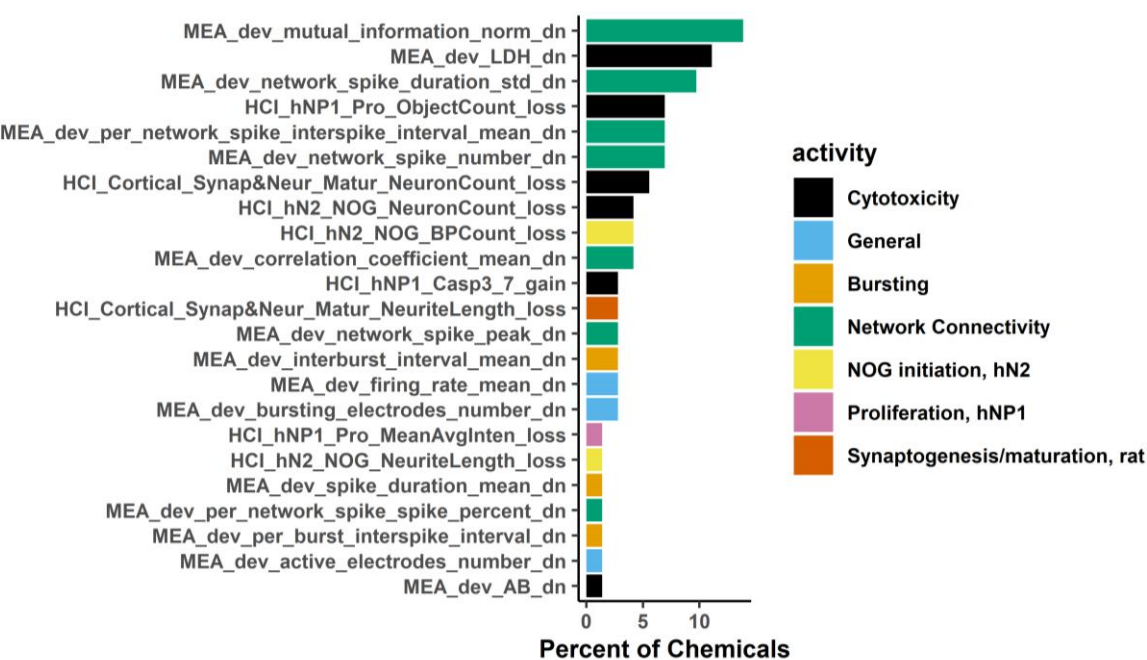
https://cran.r-project.org/web/packages/tcpl/vignettes/Data_processing.html#level-4

How does a broad screening battery collectively inform DNT-relevant bioactivity?

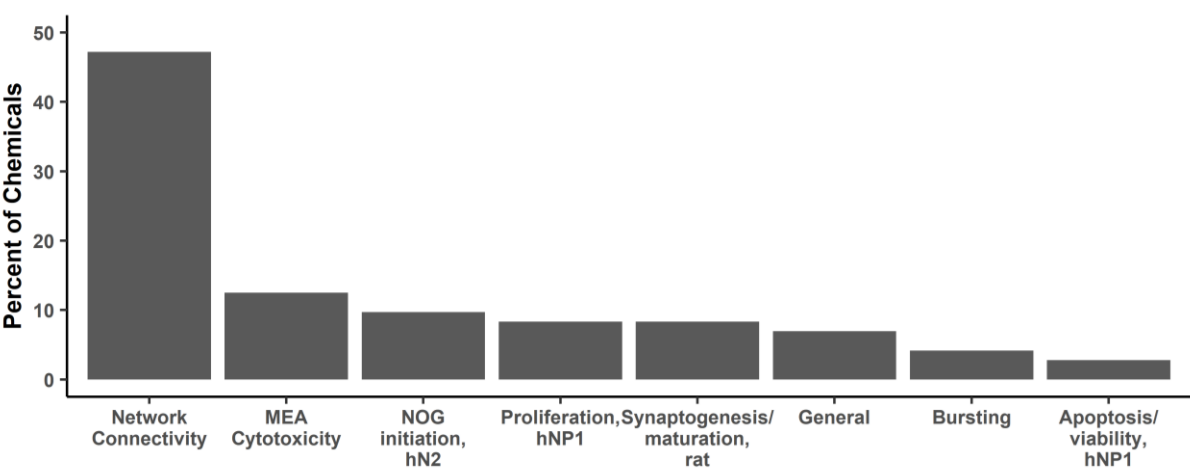


Evaluating sensitivity in the DNT NAM battery

Minimum potency by endpoint



Minimum potency by activity type



Network connectivity and cytotoxicity are the most sensitive ‘activity types’ in the battery.

Evaluating sensitivity in the DNT NAM battery

Feature importance

- MUNDY_HCI_hNP1_Pro_ObjectCount_loss
- MUNDY_HCI_hN2_NOG_NeuronCount_loss
- CCTE_Shafer_MEA_dev_network_spike_peak_dn
- CCTE_Shafer_MEA_dev_LDH_dn
- CCTE_Shafer_MEA_dev_active_electrodes_number_dn
- CCTE_Shafer_MEA_dev_mutual_information_norm_dn
- CCTE_Shafer_MEA_dev_bursting_electrodes_number_dn
- CCTE_Shafer_MEA_dev_network_spike_number_dn
- MUNDY_HCI_Cortical_NOG_NeuriteLength_loss
- MUNDY_HCI_hNP1_Casp3_7_gain
- CCTE_Shafer_MEA_dev_firing_rate_mean_dn
- CCTE_Shafer_MEA_dev_per_network_spike_spike_number_mean_dn
- CCTE_Shafer_MEA_dev_AB_dn
- CCTE_Shafer_MEA_dev_correlation_coefficient_mean_dn
- CCTE_Shafer_MEA_dev_network_spike_duration_std_dn
- CCTE_Shafer_MEA_dev_spike_duration_mean_dn
- CCTE_Shafer_MEA_dev_burst_duration_mean_dn
- CCTE_Shafer_MEA_dev_burst_rate_dn
- CCTE_Shafer_MEA_dev_per_burst_spike_percent_dn
- CCTE_Shafer_MEA_dev_per_network_spike_interspike_interval_mean_dn
- CCTE_Shafer_MEA_dev_interburst_interval_mean_dn
- CCTE_Shafer_MEA_dev_per_network_spike_spike_percent_dn
- MUNDY_HCI_hN2_NOG_BPCount_loss
- MUNDY_HCI_hN2_NOG_NeuriteLength_loss
- MUNDY_HCI_Cortical_NOG_NeuriteCount_loss
- MUNDY_HCI_Cortical_Synap.Neur_Matur_NeuriteCount_loss
- MUNDY_HCI_Cortical_NOG_BPCount_loss
- MUNDY_HCI_Cortical_Synap.Neur_Matur_NeuriteLength_loss
- MUNDY_HCI_Cortical_Synap.Neur_Matur_CellBodySpotCount_loss
- MUNDY_HCI_Cortical_Synap.Neur_Matur_NeuronCount_loss



A supervised machine learning algorithm identifies cytotoxicity (hNP1 cell line) as the most important endpoint in predicting minimum potency in the DNT NAM battery.

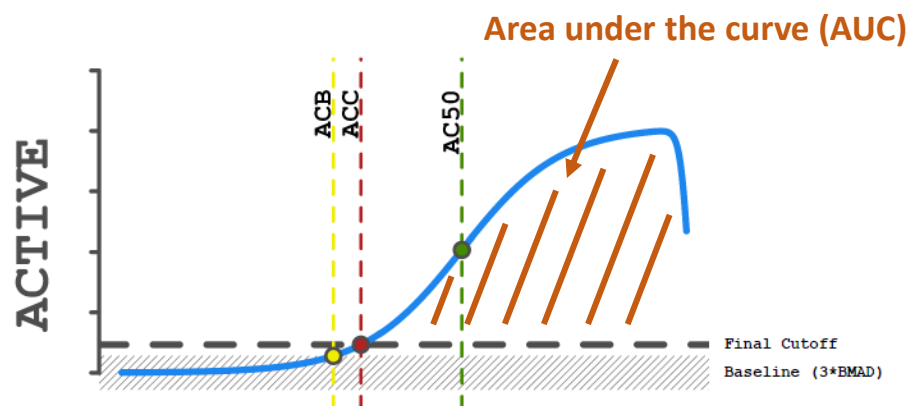
randomForest(minimum.potency ~ + potency.matrix)

%IncMSE: Percent increase in mean square error (MSE)

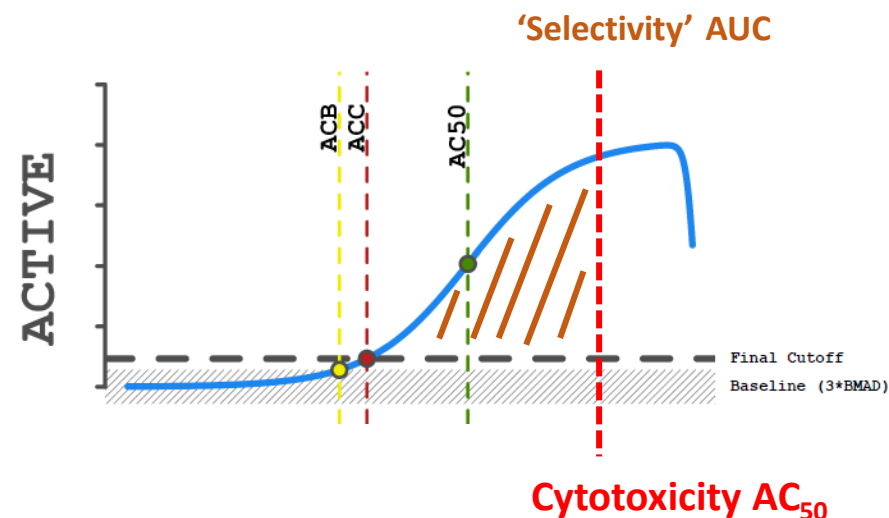
Calculating a *selectivity* metric:

Selectivity: activity at concentrations lower than cytotoxicity.

Point of departure estimates:

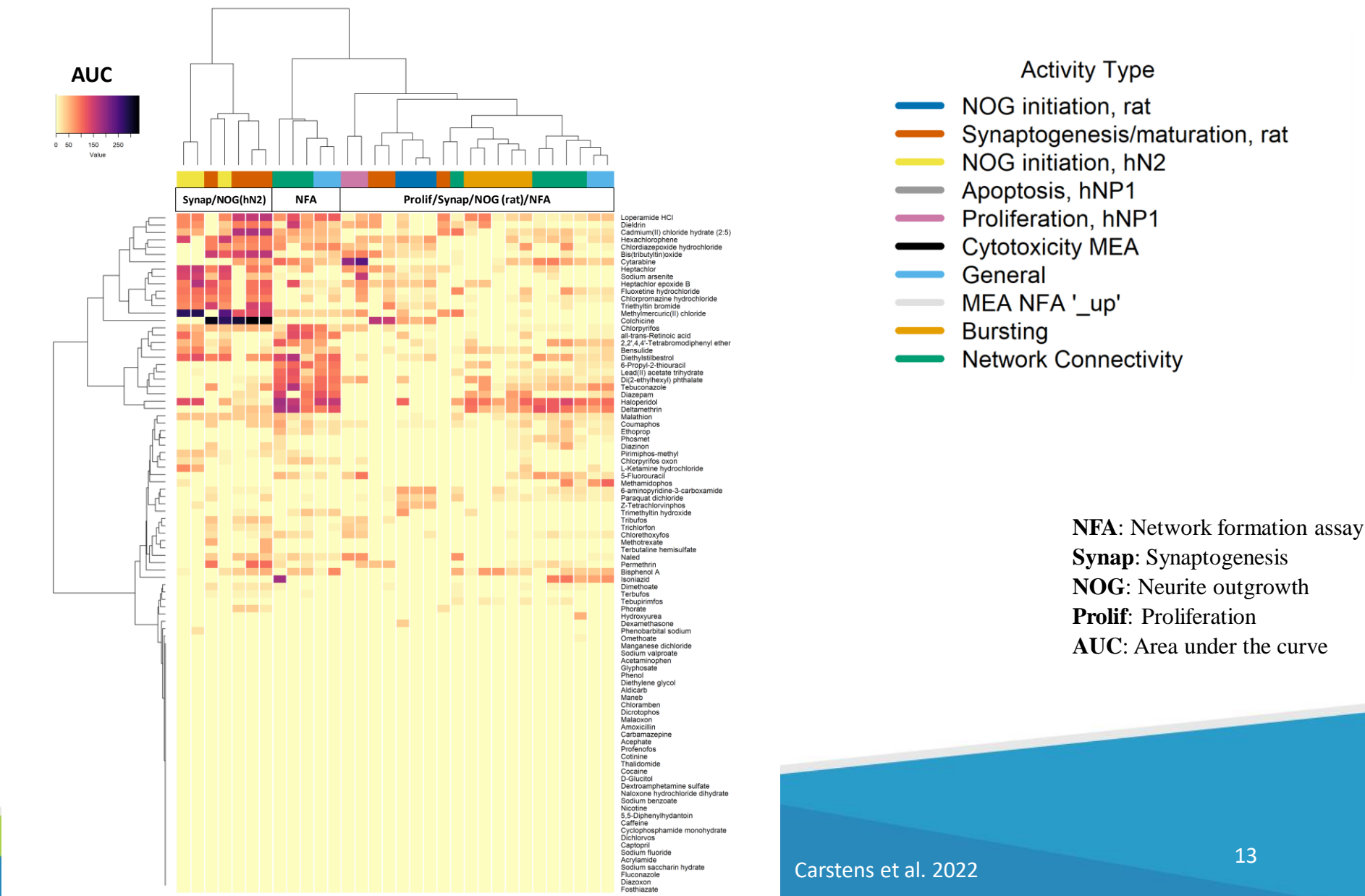


Point of departure estimates:



https://cran.r-project.org/web/packages/tcpl/vignettes/Data_processing.html#level-4

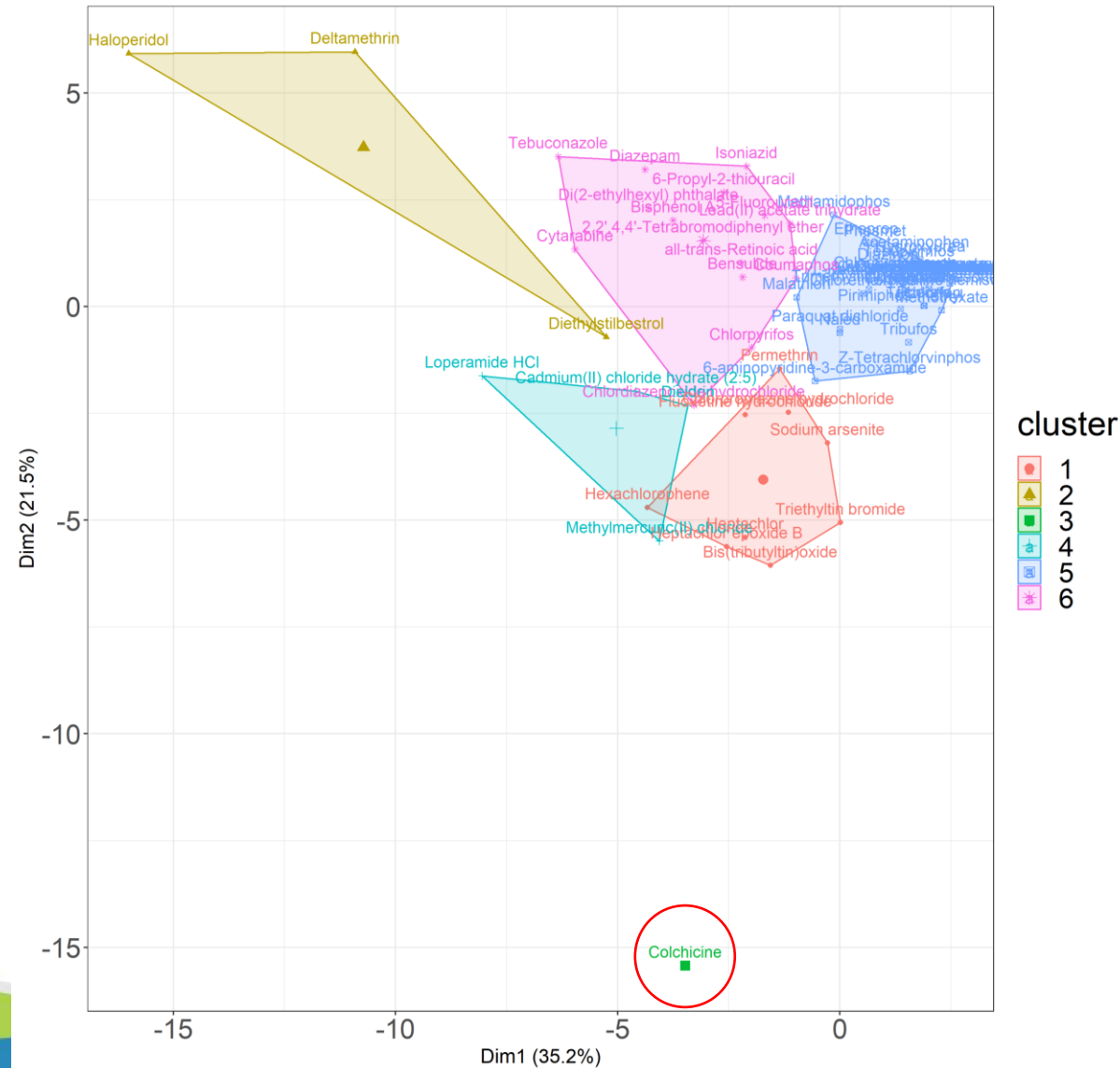
Evaluating *selectivity* is informative for identifying patterns of activity.



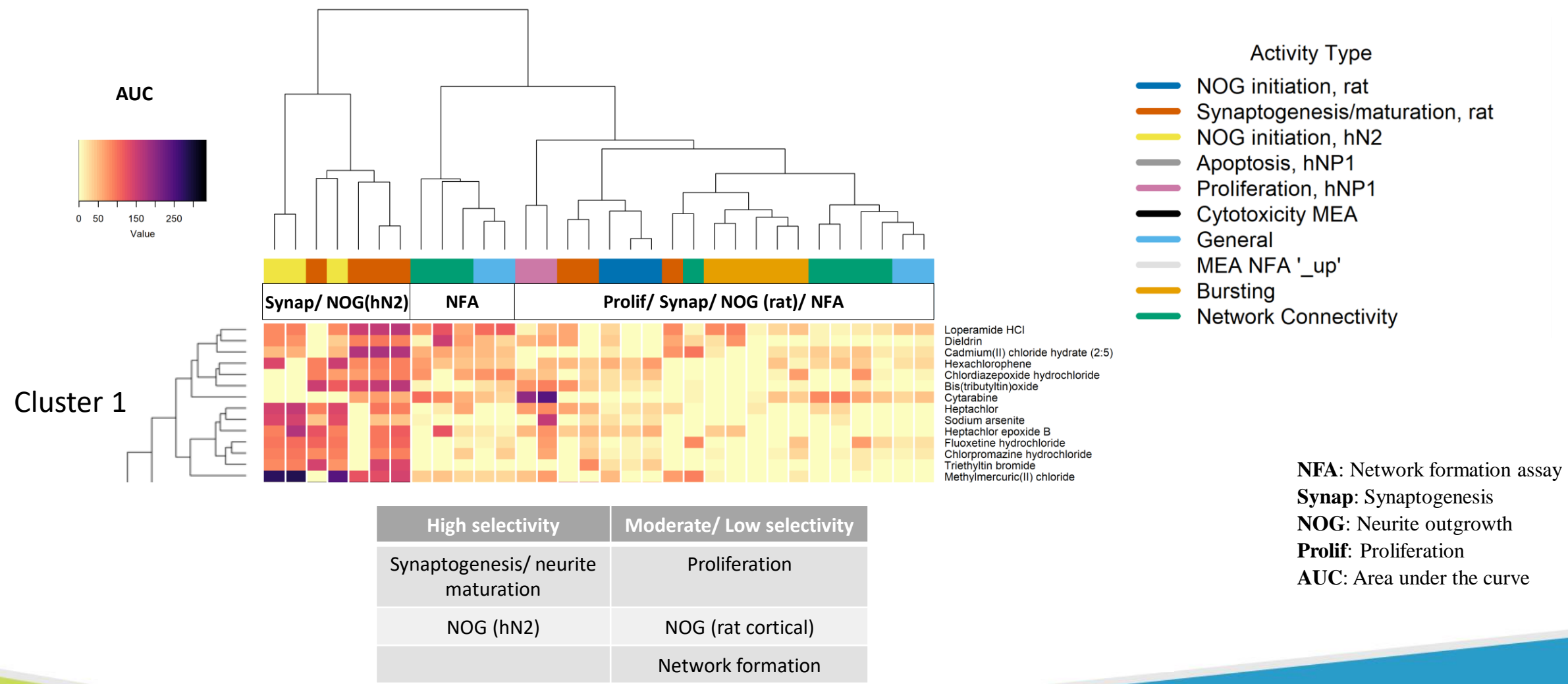
Evaluating *selectivity* is informative for identifying patterns of activity.

Unsupervised machine learning algorithm

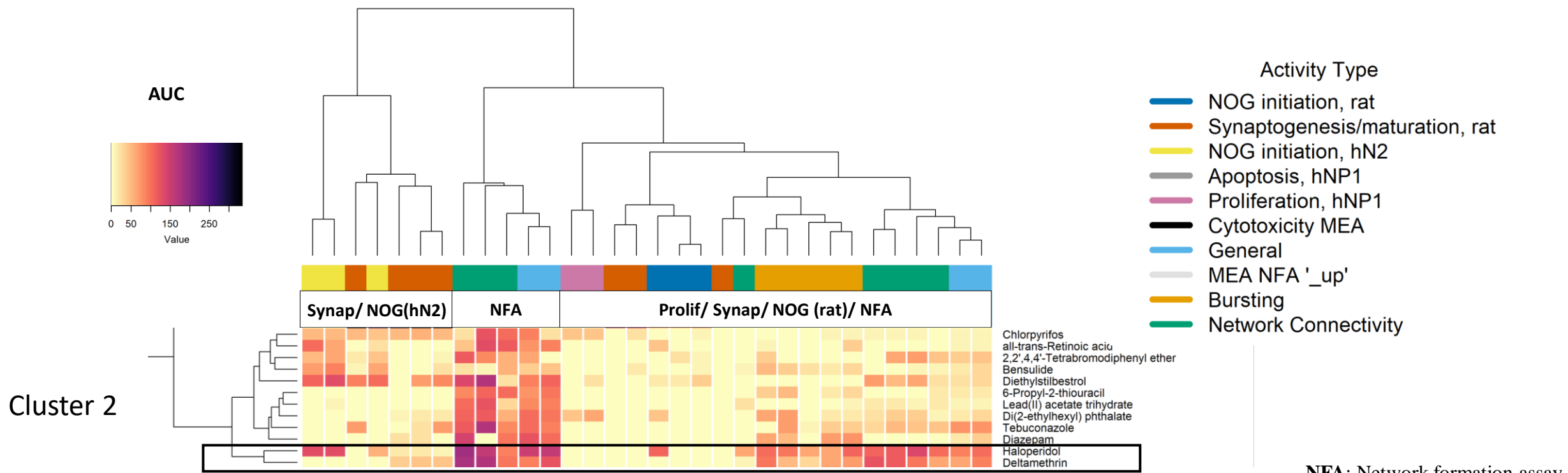
K-means clustering



Evaluating *selectivity* is informative for identifying patterns of activity.



Evaluating *selectivity* is informative for identifying patterns of activity.



Haloperidol: antipsychotic- Dopamine D₂ receptor antagonist

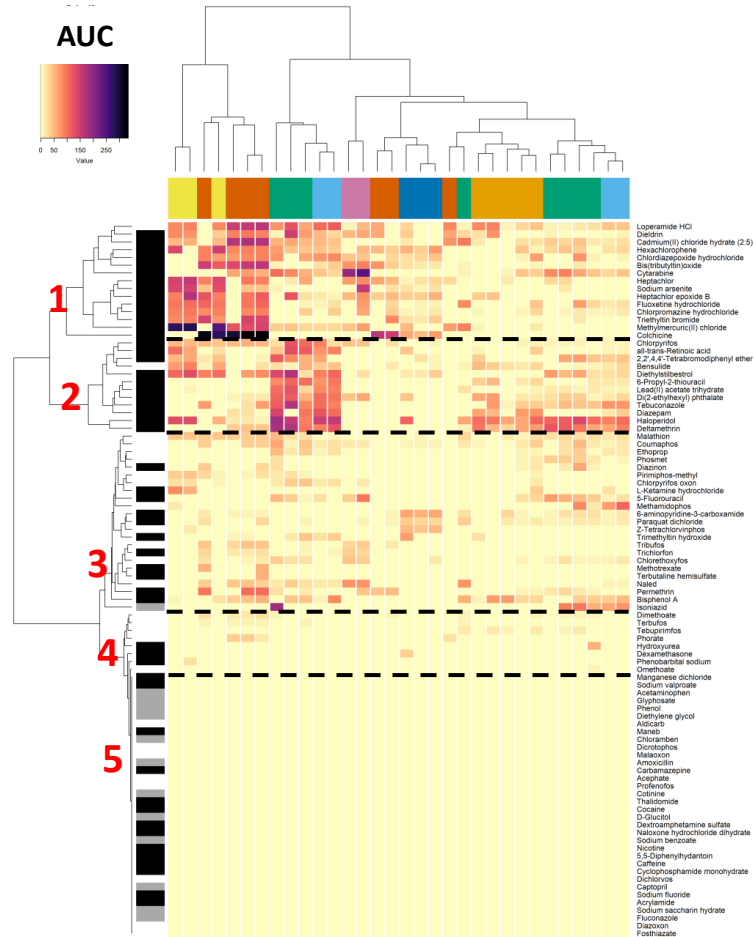
Deltamethrin: pyrethroid insecticide-voltage-gated sodium channels modulators

High selectivity	Moderate/ Low selectivity
Network connectivity	NOG (hN2)
General neuronal activity	Bursting

NFA: Network formation assay
Synap: Synaptogenesis
NOG: Neurite outgrowth
Prolif: Proliferation
AUC: Area under the curve

Carstens et al. 2022

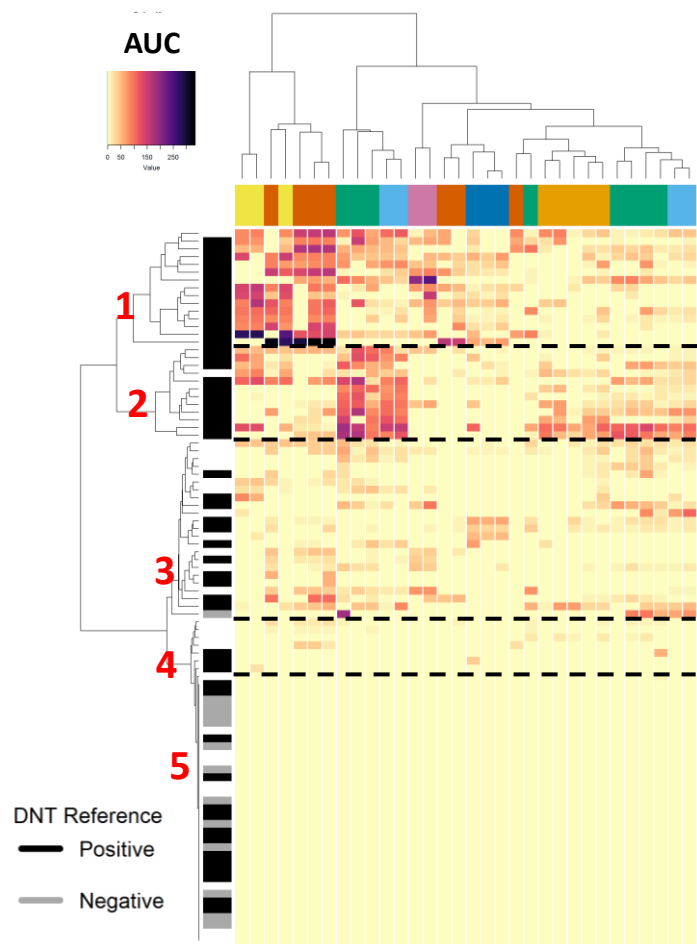
Evaluating *selectivity* is informative for identifying patterns of activity.



Key findings

- Selective data is more informative in identifying differential patterns of functional bioactivity compared to non-selective data.
- A subset of compounds demonstrate cell-type specific effects (active in the NOG assay in the hN2 cell model but not rat cortical).
- Selective activity clusters do not appear to be explained by shared mode-of-action.
- Limitations in evaluating relationships between bioactivity clusters and mode-of-action or activity-structure due to a limited dataset.

Can we build a model to classify compounds that demonstrate *in vivo* DNT bioactivity?



		<i>In vivo</i> evaluation chemicals	
		Positive (53) <small>Mundy et al. 2015 Aschner et al. 2016 Harrill et al. 2018</small>	Negative (13) <small>Martin et al. under revision</small>
Classification	Cluster 1 Synap/ prolif/ NOG/ Neurite maturation	14	0
	Cluster 2 General/ network/ bursting activity/ synap	11	0
	Cluster 3 General/ network activity/ bursting/ synap/NOG	11	1
	Cluster 4 General/ network activity/ bursting/ synap/ NOG	3	0
	Cluster 5 'Inactive/ equivocal'	14	12

	Positive	Negatives
Selective activity (Clusters 1,2,3,4)	True positive: 39	False positive:1
Inactive/ equivocal (Cluster 5)	False negative: 14	True Negative: 12

<u>Selective</u>	<u>Non-selective</u>
Sensitivity= 74%	Sensitivity= 93%
Specificity= 92%	Specificity= 69%

Carstens et al. 2022

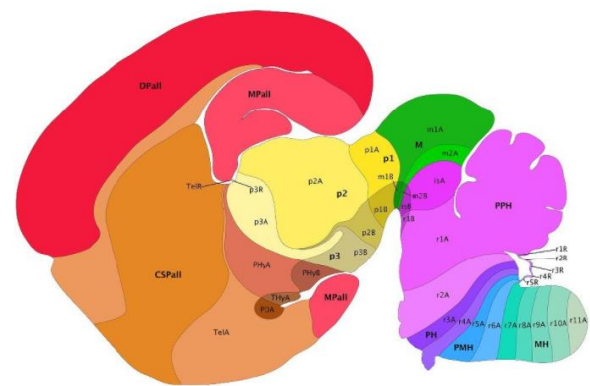
Can we identify biological gaps in the current EPA DNT NAM battery?

Are we capturing the target mechanism in the DNT NAM battery?

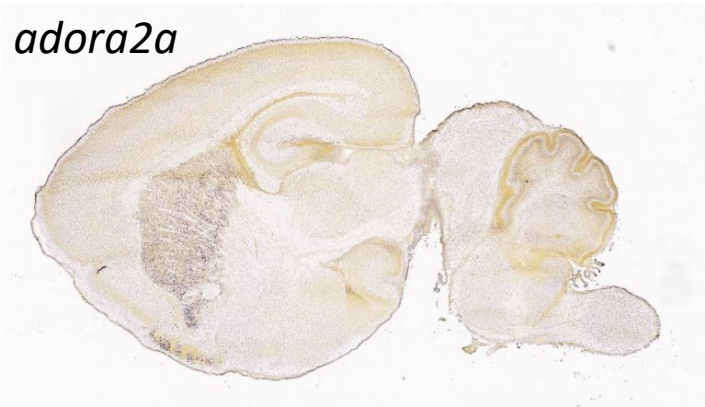
False negative: Caffeine

Caffeine targets adenosine receptor (adenosine A2a receptor)
Is adenosine expressed in cortical cells in early development?

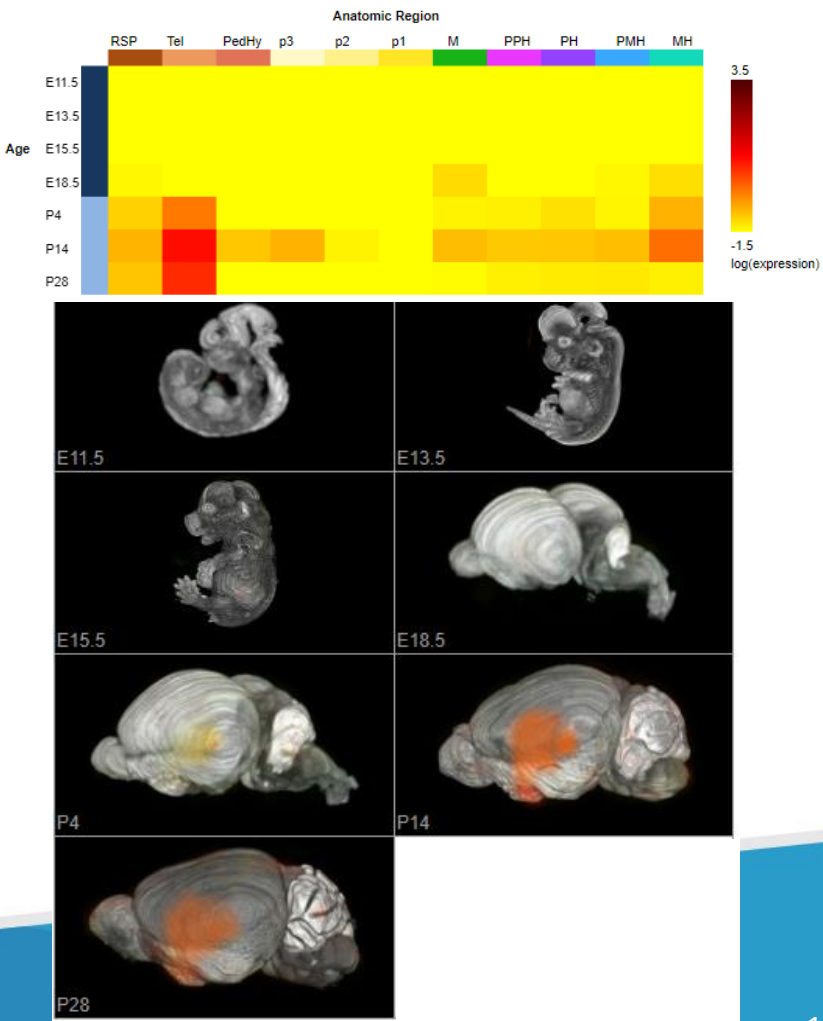
P4 mouse brain



<https://developingmouse.brain-map.org/>



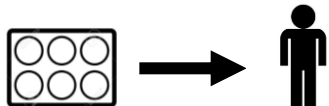
In situ hybridization



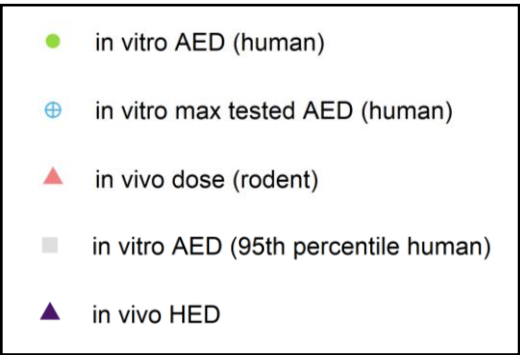
In vitro to in vivo extrapolation (IVIVE) using high-throughput toxicokinetic (HTTK) modeling

Are we testing at high enough concentrations *in vitro*?

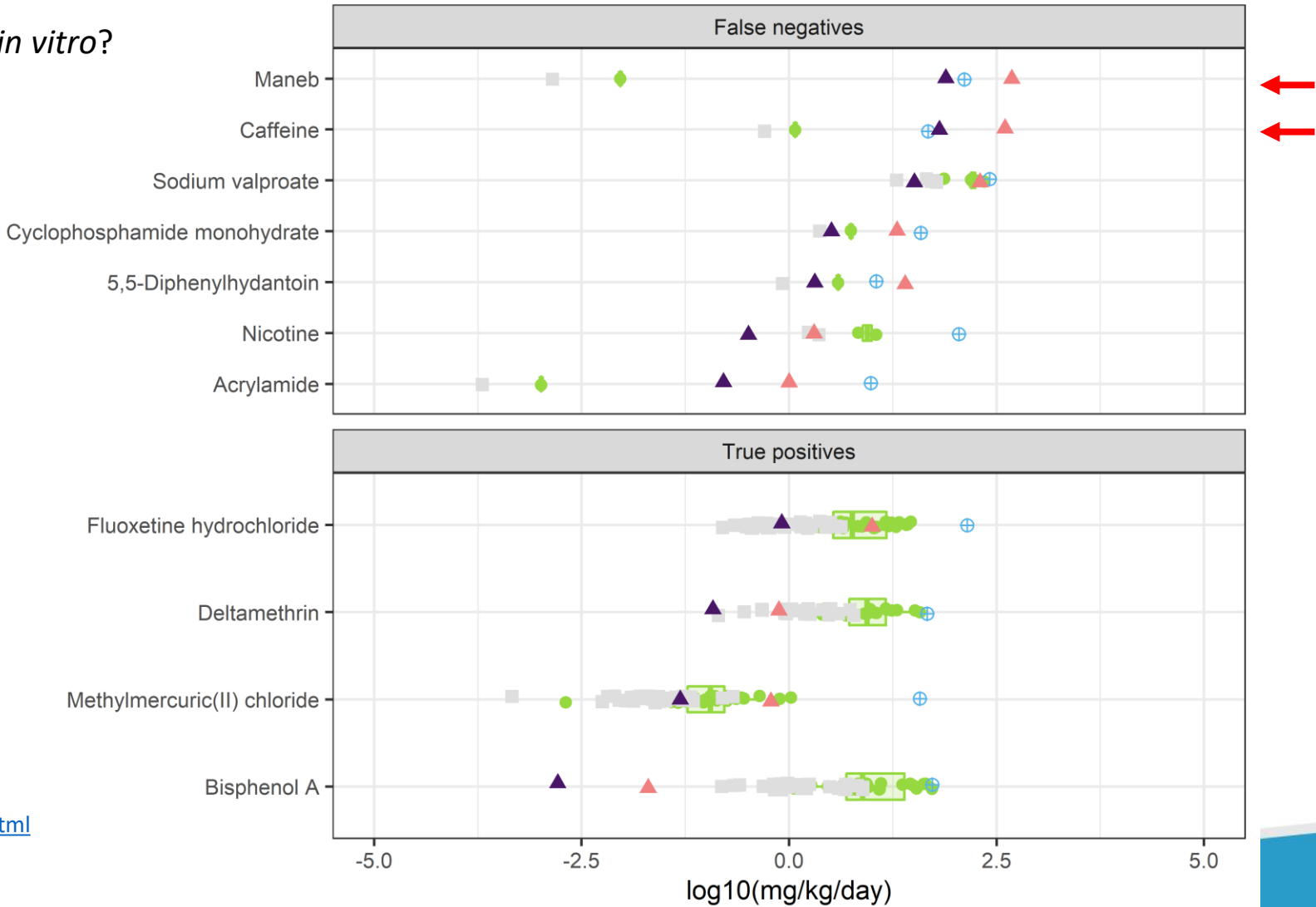
AED: administered equivalent dose



HED: human equivalent dose



'httk' R package: <https://cran.r-project.org/web/packages/httk/index.html>



Identifying gaps in the EPA DNT NAM battery

Is a false negative active in other DNT NAMs that are not included in the current EPA battery?

False negative: Maneb

Active in 3 DNT NAM endpoints outside the US EPA battery

- UKN4: Neurite area in LUHMES cells (human dopaminergic precursor)
- UKN5: Neurite area in peripheral nervous system (human derived dorsal root ganglia)
- UKN2: Neural crest cell (NCC) migration assay (human derived)

Results from the 2020 EFSA Scientific Report on DNT NAMs (μM)

compound name	CAS #	Selection criteria	UKN4 neurite area	UKN5 peripheral neurite area	NPC4 neurite length	NPC4 neurite area	UKN2 (NCC migration)	NPC2a72h (radial glia migr.)	NPC2a120h (radial glia migr.)	NPC2b (neuron migration)	NPC2c (oligo. migr.)
Maneb	12427-38-2	in vivo positive	3.2	2	no hit	no hit	11.4	no hit	no hit	no hit	no hit

Modified from Masjothusmann et al. 2020

Table 2. Proposed Assays for Evaluation As an In Vitro DNT Battery

Process	Assays	References
*Proliferation	→ hNP1 NPC1	Harrill et al. (2018) Baumann et al. (2016) and Barenys et al. (2017)
*Apoptosis Migration	UKN1 → hNP1 NPC2	Balmer et al. (2012) Harrill et al. (2018) Baumann et al. (2016) and Barenys et al. (2017)
Neuron differentiation	UKN2 NPC3	Nyffeler et al. (2017) Baumann et al. (2016) and Barenys et al. (2017)
Oligodendrocyte differentiation & maturation	NPC5/6	Baumann et al. (2016) and Barenys et al. (2017)
*Neurite outgrowth	→ iCell gluta (hN2) UKN 4 & 5 (rat) NPC4	Harrill et al. (2018) Krug et al. (2013) Baumann et al. (2016) and Barenys et al. (2017)
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Sachana, M., et.al. 2019, Toxicological Sciences

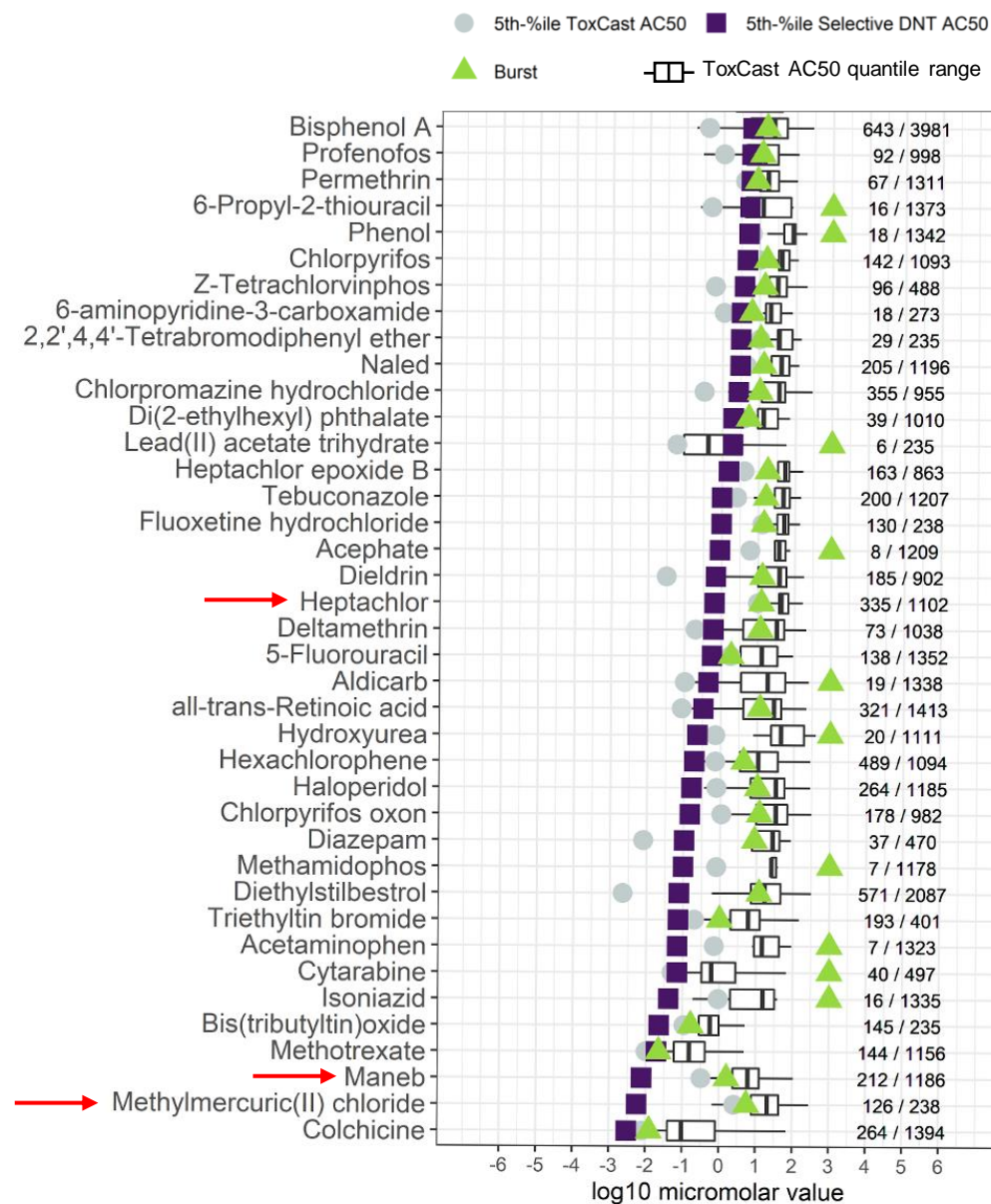
Comparison of *selective* DNT NAM activity to ToxCast/Tox21 database

ToxCast includes >1,500 assay endpoints and covers heterogeneous assay types, tissue sources, gene targets, and biological responses.

Examples of biological responses in ToxCast:

- Cell proliferation and death
- Cell differentiation
- Enzymatic activity
- Mitochondrial depolarization
- Protein stabilization
- Oxidative phosphorylation
- Reporter gene activation
- Receptor binding
- Receptor activity
- Metabolomic responses (stem cells)

<https://comptox.epa.gov/dashboard/assay-endpoints>



Conclusions

1) How does the DNT NAM battery collectively inform DNT-relevant bioactivity?

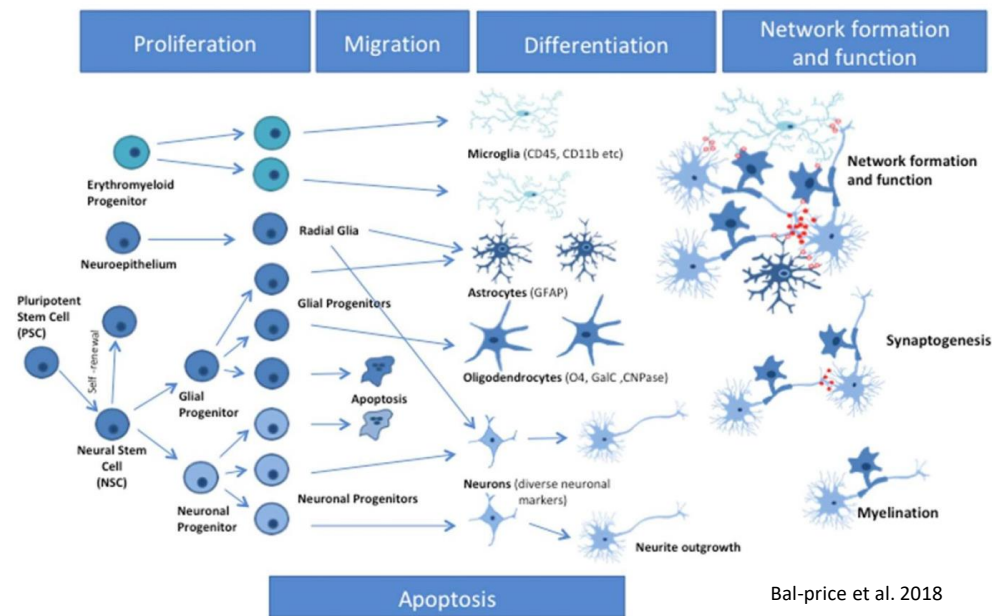
- Selective data is more informative in identifying differential patterns of functional bioactivity than non-selective data.
- Selective activity clusters do not appear to be explained by mode-of-action.

2) Can we build a model to classify compounds that demonstrate *in vivo* DNT bioactivity?

- Using the selectivity metric, DNT reference chemicals are classified with high specificity and moderate sensitivity.
- False negatives provide insight into experimental and biological limitations.

3) Can we identify gaps in the current DNT NAM battery and/or broader ToxCast/ Tox21 database?

- Identified gaps in target receptor which may be associated with cell-type, species or developmental timepoint.
- Identified gaps in the current battery of DNT NAMs that appear to be covered by assays representing additional 'key events' in neurodevelopment.
- DNT NAMs data provides added value to ToxCast/ Tox21 database from the perspective of capturing health protective potencies.





Questions?

Acknowledgements

Tim Shafer
Katie Paul Friedman
Theresa Freudenrich
Kathleen Wallace
Cina Mack
Melissa Martin
Amy Carpenter
Seline Choo
Jackson Keever
Josh Harrill
Megan Culbreth

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Assay data:

Available in ToxCast invitrodb v 3.4
<https://doi.org/10.23645/epacomptox.6062479.v6>