

# In Vitro Transcriptomic Points of Departure (tPODs) for 42 diverse chemicals in a human (Caco-2) and fish (RT-gill-W1) cell line

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# Transcriptomic Points of Departure (tPOD)

- transcriptomics provide lots of information efficiently <sup>1</sup>
- in recent years growing interest in benchmark dose (BMD) modeling of transcriptomic data to identify an exposure level associated with a significant, concerted change in gene expression = tPOD<sup>2</sup>
- 2024: US EPA published Transcriptomic Assessment Product (ETAP) to yield transcriptomic-based reference values
  - tPODs from 5-day animal study associate with chronic animal toxicity data
- so, what about tPODs from in vitro data? Might they inform of chronic toxicity estimates?

1: Head, J.A., et al. *Environmental Toxicology and Chemistry*, 5954.

2: Johnson, K.J., et al *Toxicological Sciences*, 190(2), 127-132.

# Objectives

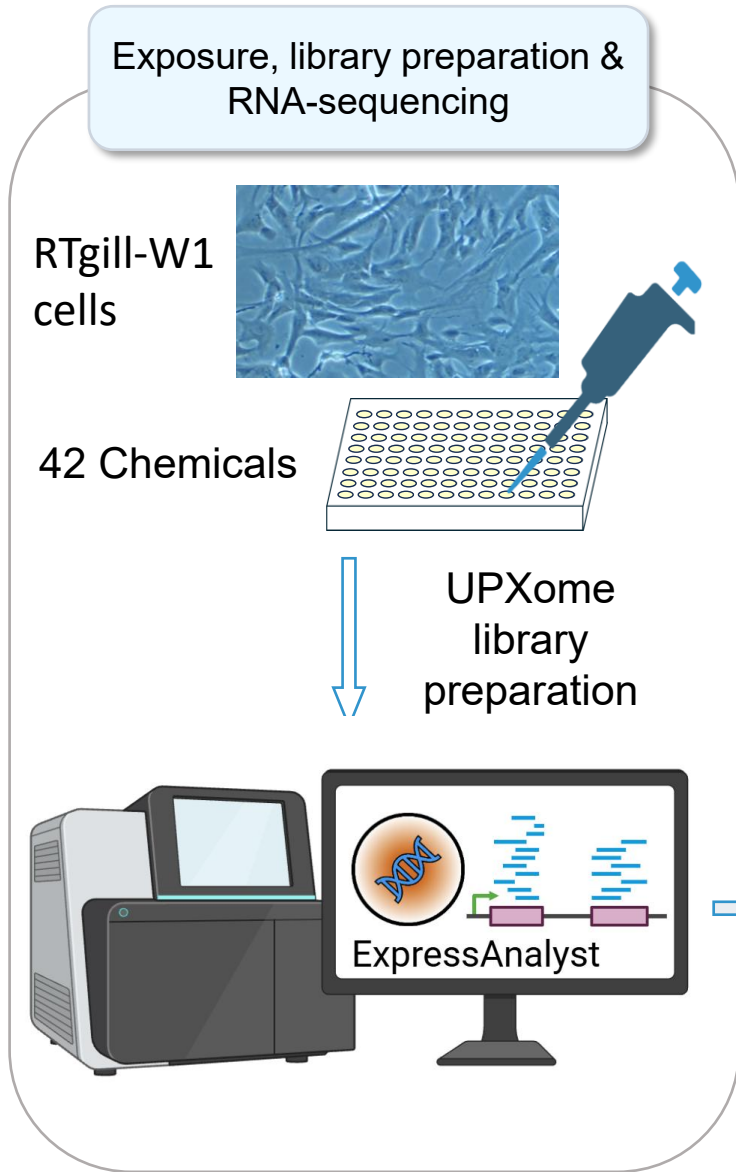
- to determine if tPOD data from in vitro studies of rainbow trout (*Oncorhynchus mykiss*) gill (RTgill-W1) cells and human intestinal epithelial cells (Caco-2) exposed to chemicals of varying properties associate with chronic toxicity data for each respective compound

## Secondary objectives:

- test and refine the TPD-Seq workflow<sup>3</sup>, from in vitro exposures to sequencing to bioinformatics to IVIVE calculations to chronic data
- assess this approach's chemical domains of applicability

# Methods (TPD-Seq Workflow<sup>3</sup>)

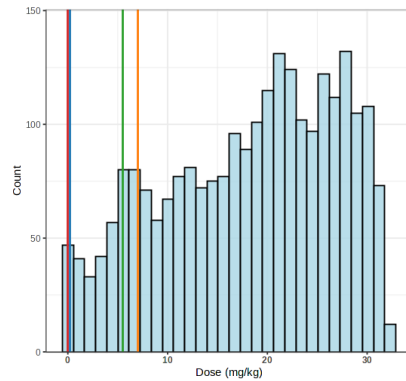
- expose cells (RTgill-W1, Caco-2) to 42 chemicals
  - blinded to chemicals tested, and so an in vitro screening strategy developed with all chemicals assessed at 100uM and then binned into 4 potency groups
- cytotoxic and transcriptomic responses measured and compared
  - 24 hour exposures; 3-fold dilution series over 8 concentrations
  - each microplate exposure had a solvent and positive control
- resulting data compared to available measures of chronic toxicity for each compound in both fish and mammalian models
- focus today on RTgill-W1 results i.e., 'omics-enhanced OECD249



tPOD calculation & modeling

*In vitro* - Mass Balance Model (IV-MBM) predicted tPODs

Nominal transcriptomic point of departure (tPODs)



Comparison with *in vitro* studies & chronic animal studies

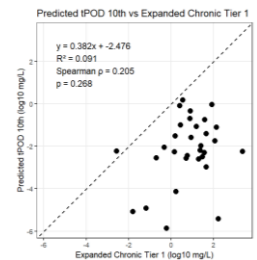
**ECOTOX** Knowledgebase

Zebrafish cell lines and high-throughput transcriptomics: advancing *in vitro* and bioinformatics methods for supporting environmental risk assessment

Peter G. Schumann<sup>1</sup>, Joseph Bundy<sup>2</sup>, Derik E. Haggard<sup>3</sup>, Logan Everett<sup>4</sup>, Joshua A. Harrill<sup>5</sup>, Felix Harris<sup>6</sup>, David Byon<sup>7</sup>, Jacob Collins<sup>8</sup>, Claudia Rivetti<sup>9</sup>, Bruno Campos<sup>10,11</sup>, Geoff Hodjoes<sup>11</sup>, Carlie A. LaLone<sup>12,13</sup>

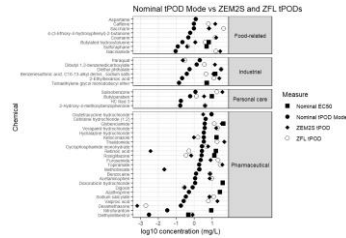
versus

*In vivo* chronic mortality and non-mortality PODs

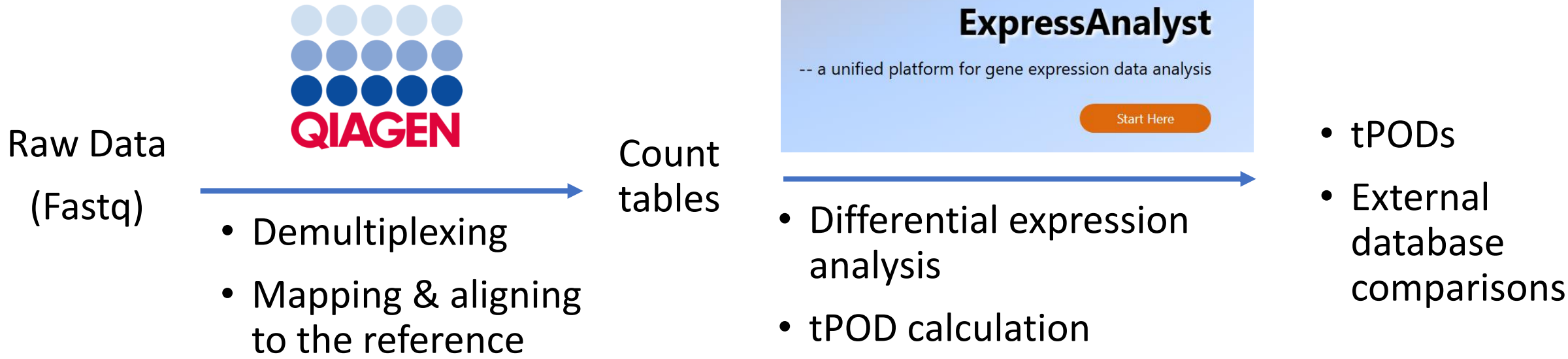


versus

*In vitro* tPODs from ZEM2S and ZFL cell lines



# Data analysis pipeline



**Demultiplexing:** separate samples by barcode/index  
**Mapping:** Locate reads on the reference genome  
**Alignment:** base-by-base compare, record mismatches



# 1. Can we screen chemicals “blinded”?

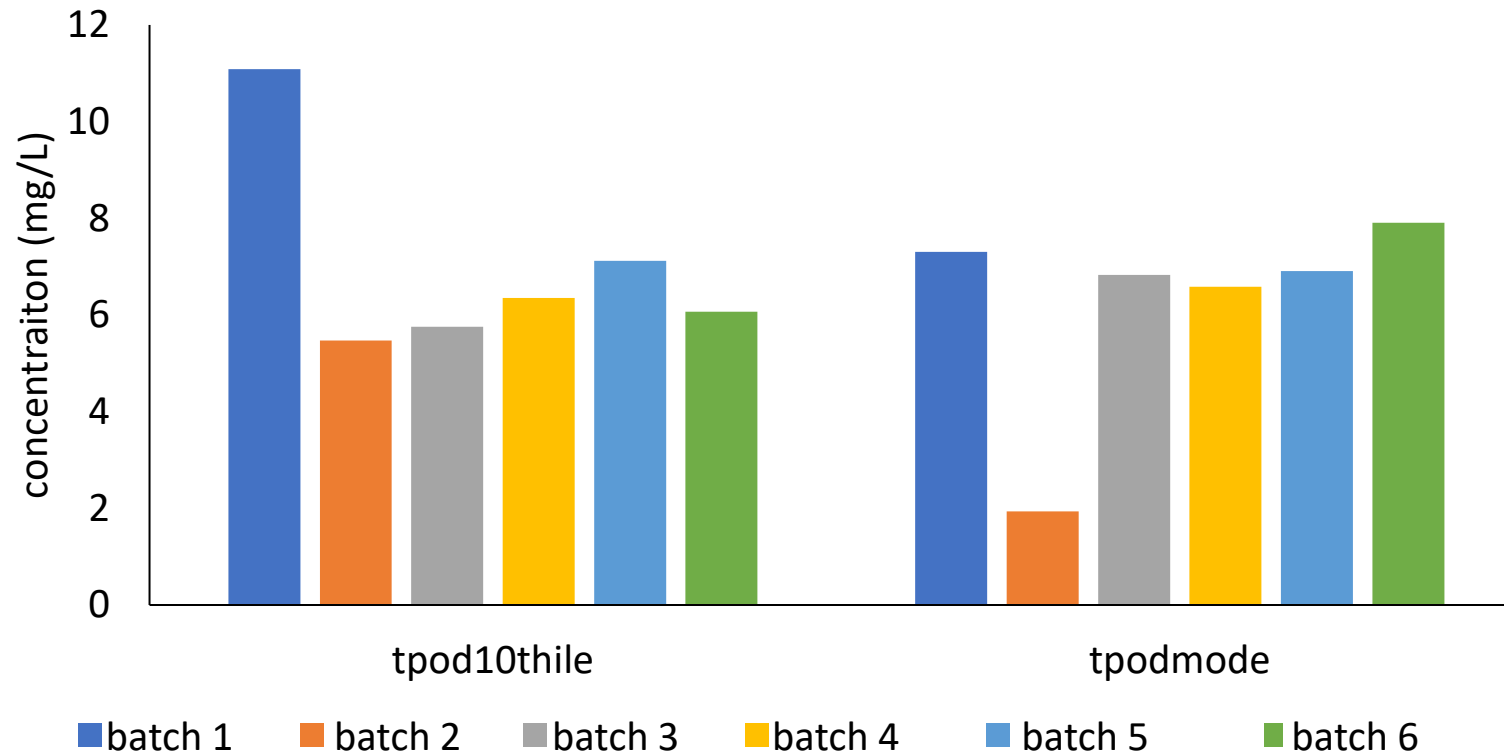
Zone #	# of chem.	Procedure
Zone 1	24	proceed directly to transcriptomics exposure at 100 $\mu$ M
Zone 2	9	conduct rangefinder test at 100 $\mu$ M on a semi-log scale over an eight-point dilution series to determine EC20.
Zone 3		
Zone 4	9	conduct rangefinder test starting at 100 $\mu$ M over an eight-point 10-fold dilution series to determine EC20

Cytotoxicity QC, OECD249

- Background no-cell control variation:
  - Across all plates difference between no-cell control wells < 20%
- % cell viability in solvent control
  - % cell viability >90% relative to unexposed cells; averaged 99%



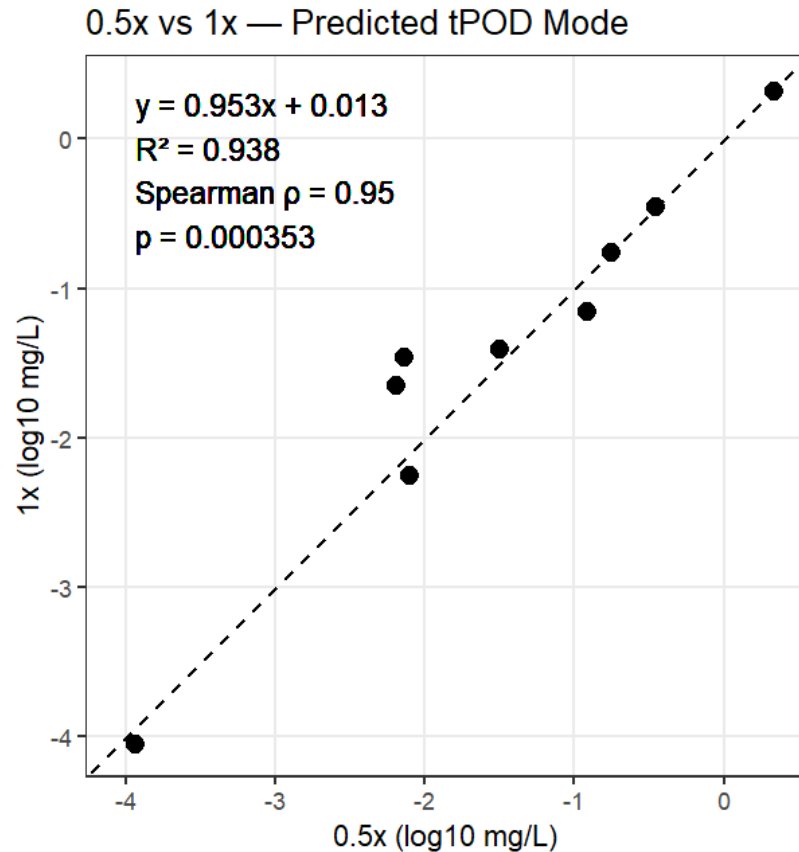
## 2. Are in vitro tPODs reproducible?



**Tested DCA 6x: tPOD Coefficient of variance ~30%**



### 3. Can the workflow be more cost-effective?



**ExpressAnalyst**  
-- a unified platform for gene expression data analysis

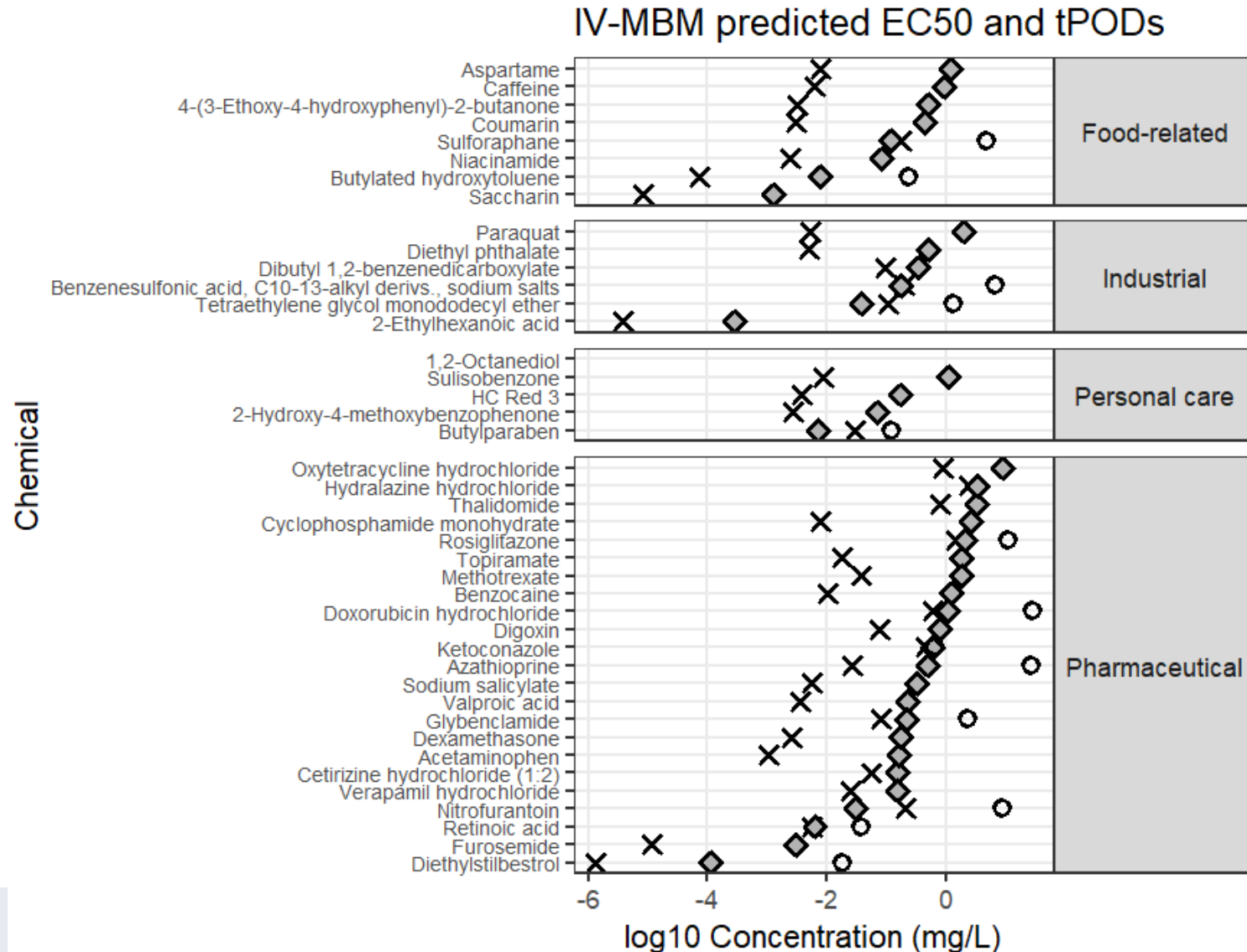
[Start Here](#)

**-half-volume reactions successful**

**-free online bioinformatics tool**



# 4. in vitro tPODs vs cytotoxicity EC50s?



**EC50: 13/44 chemicals**  
**tPOD: 43/44 chemicals**  
**-tPOD ~23x lower than EC50**

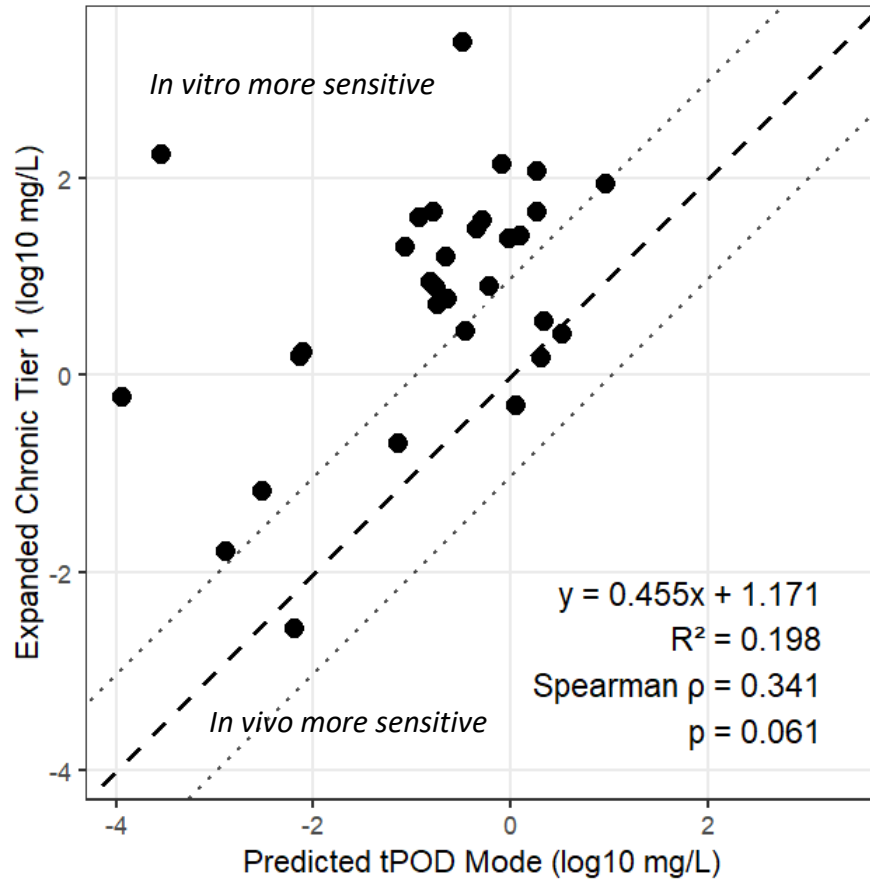
# 5. In vitro tPODs vs. in vivo fish chronic

- Compared to the following datasets obtained from Schumann et al. as per next talk by Dr. Rivetti:
  - Tier 1 dataset – fish chronic apical endpoints, mortality
  - Tier 2 dataset – fish chronic, non-mortality endpoints (e.g., biochemical)
  - Expanded criteria dataset (Exp) – not quality filtered
  - Quality filtered dataset (QualFilt) – quality filtered (filtered for chemical purity < 85%, non-relevant exposure routes, chemical formulations rather than pure substances, non-statistically significant effects, non-standard endpoints, records with insufficient controls, records missing test duration information, mixture studies, studies lacking clear dose-response relationships, and records where NOEC were equal to highest tested concentration).
- Schumann et al. 2025. Zebrafish cell lines and high-throughput transcriptomics: advancing in vitro and bioinformatics methods for supporting environmental risk assessment. *Toxicol Sci.* 208(2):343-356. doi: 10.1093/toxsci/kfaf127

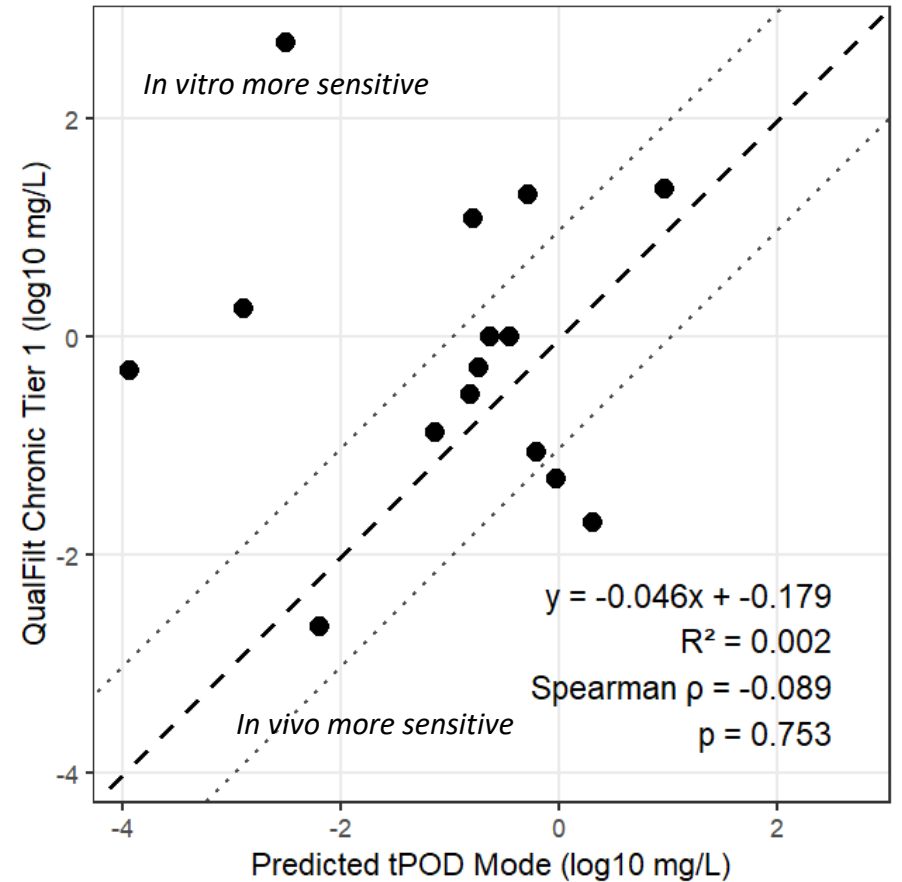


# tPODs vs. in vivo fish chronic (Mortality endpoint)

**Fish Chronic Toxicity (Mortality)**  
**all studies**



**Fish Chronic Toxicity (Mortality)**  
**Higher quality studies**

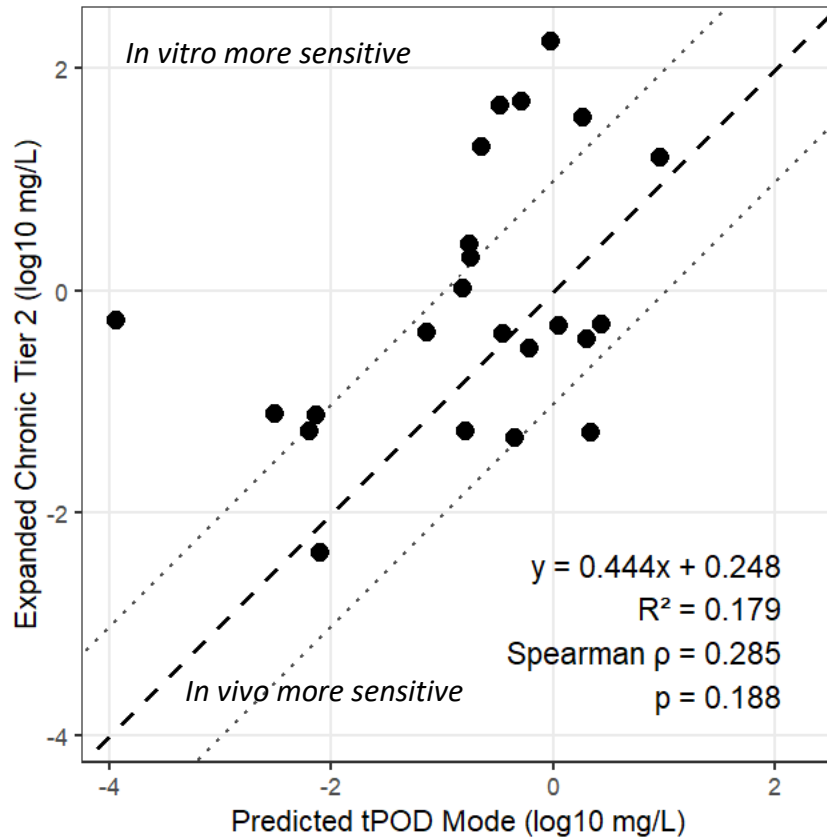


**In Vitro tPOD (RTgill-W1) – Current Study**

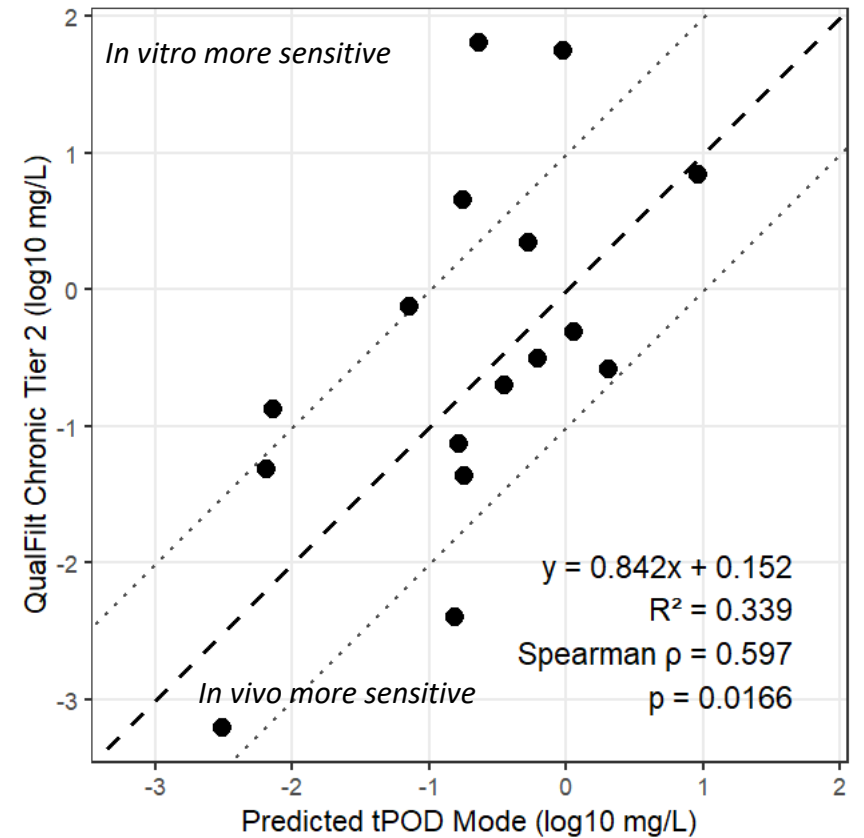


# tPODs vs. in vivo fish chronic (Non-Mortality)

**Fish Chronic Toxicity (Sub-Lethal)  
all studies**



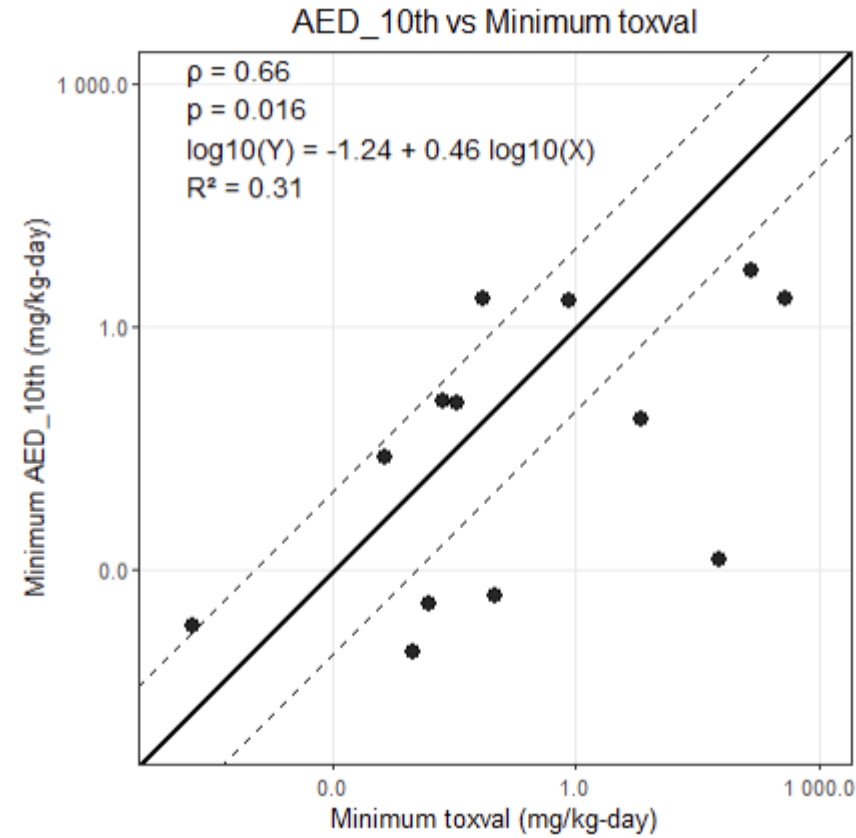
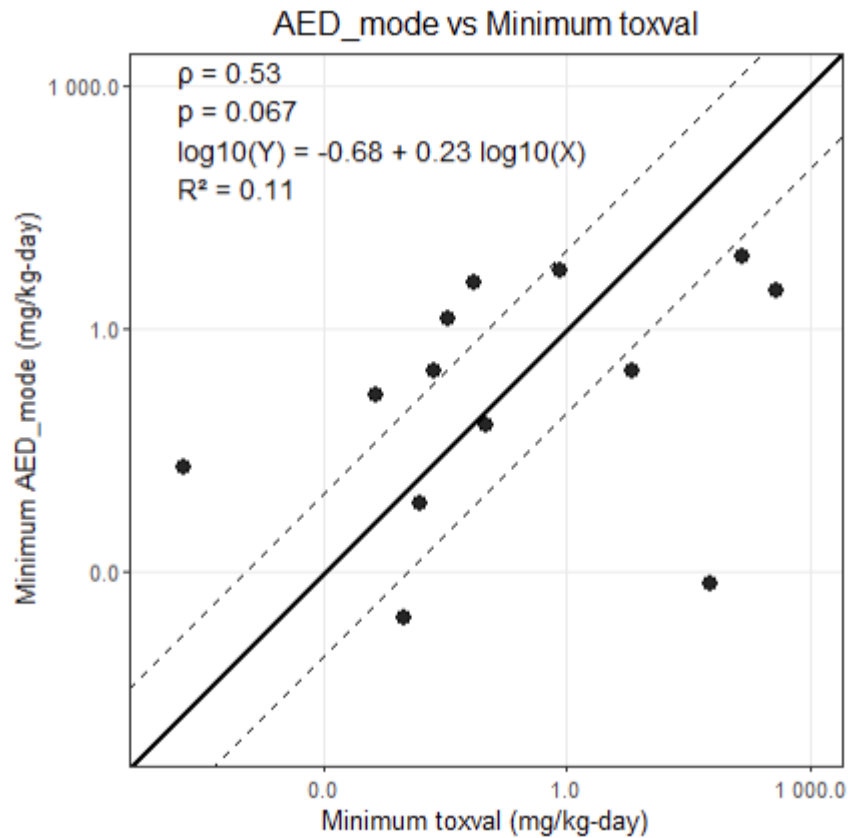
**Fish Chronic Toxicity (Sub-Lethal)  
Higher quality studies**



**In Vitro tPOD (RTgill-W1) – Current Study**



# Aside: In vitro tPOD (Caco-2 cells) vs. animal (rodent) chronic toxicity



Minimum AED (of 4 cell lines) vs minimum chronic animal studies

# Concluding remarks

- results (to date) suggest in vitro tPODs yield data that are comparable to fish chronic toxicity data → 'omics enhanced OECD249
- findings align with our past studies in vitro (e.g., 18 pesticides in RTgill-W1 cells), and also in vivo (US EPA ETAP), among others
- results presented are comparable to those from Caco-2 cells
- next steps:
  - deeper bioinformatics (e.g., biological pathways) and tPODs (e.g., sensitivity tests) analyses, and further examination of fish chronic data; both tPODs and fish chronic data can vary a lot
  - embrace the outlier! To understand domains of applicability, examine outliers (e.g., physico-chem properties, mechanisms of action, etc.)



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- Questions? Nil Basu at [niladri.basu@mcgill.ca](mailto:niladri.basu@mcgill.ca)

