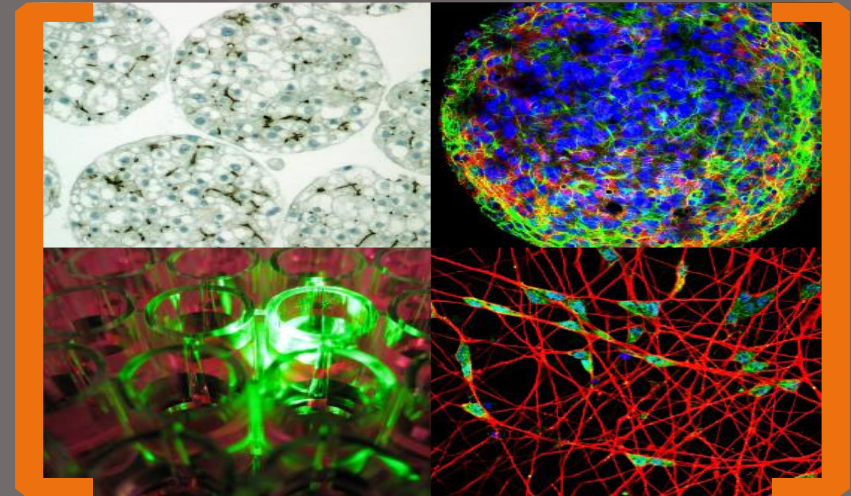


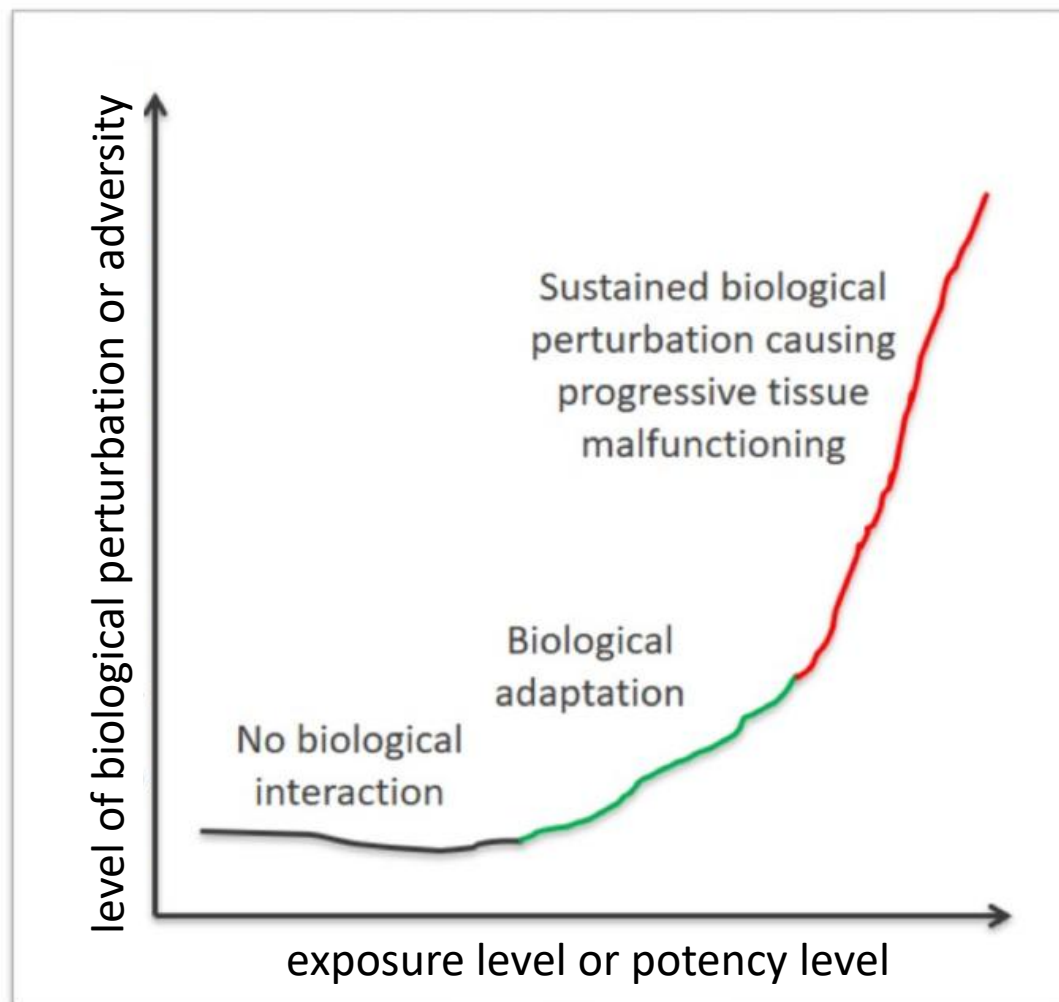
# The case of a high-concern toxicity profile screening

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# Integration of NAMs into prioritization of low tonnage compounds



Low tonnage compounds do have a limited set of in vivo studies.

How can we best use NAM data to identify those that are of concern for human health?

# Case study hypothesis

## Main hypothesis:

**High throughput NAMs do properly rank order ECHA substances based on mode-of-action in relation to CMR/ED as well as overall potency on perturbation of human biological systems and can thereby add in the prioritization of ECHA chemicals for further testing.**

## Impact:

**Ability to classify compounds and determine a performance metric on concordance would lead to further confidence on the application of NAMs.**

# Objective

**Demonstrate the overall feasibility to identify substances with a liability for a high toxicity profile based on high throughput NAM hazard information.**

Hazard information based on qualitative and quantitative mechanistic mode-of-action assessment using both *in silico* approaches and *in vitro* human test systems.

# Overall approach

- **TRAINING** set: available pools of chemicals (high tonnage compounds/in vivo safety data) → selected from high tonnage ECHA list
- **Toolbox:** EU-ToxRisk *in silico* and *in vitro* NAMs with focus on high throughput
- **TEST** set: substances based on Annex 8 compounds (10-100 tonnes) that will have 28 day repro and basic mutagenicity (*optional*)

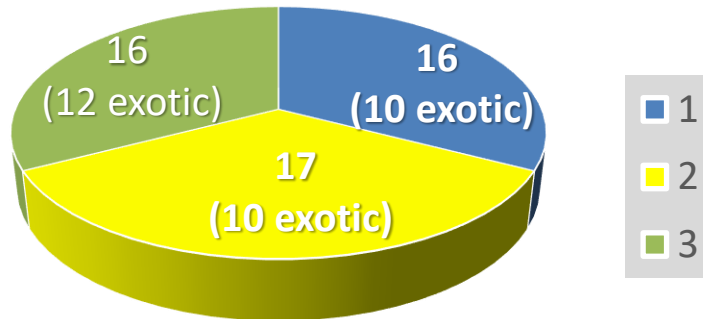
# Proof of concept study chemical selection

150 ECHA compounds  
From high tonnage ECHA list  
(data rich)

Filter for compounds with  
available high quality repeated  
dose toxicity data

44 compounds with  
RDT data in high quality  
databases analysed

Classification based on high  
quality databases



1. (blue) = toxic (NOEL < 10 mg/kg bw/d)
2. (yellow) = moderate
3. (green) = low toxic (NOEL > 100 mg/kg bw/d)

Toxicodynamics

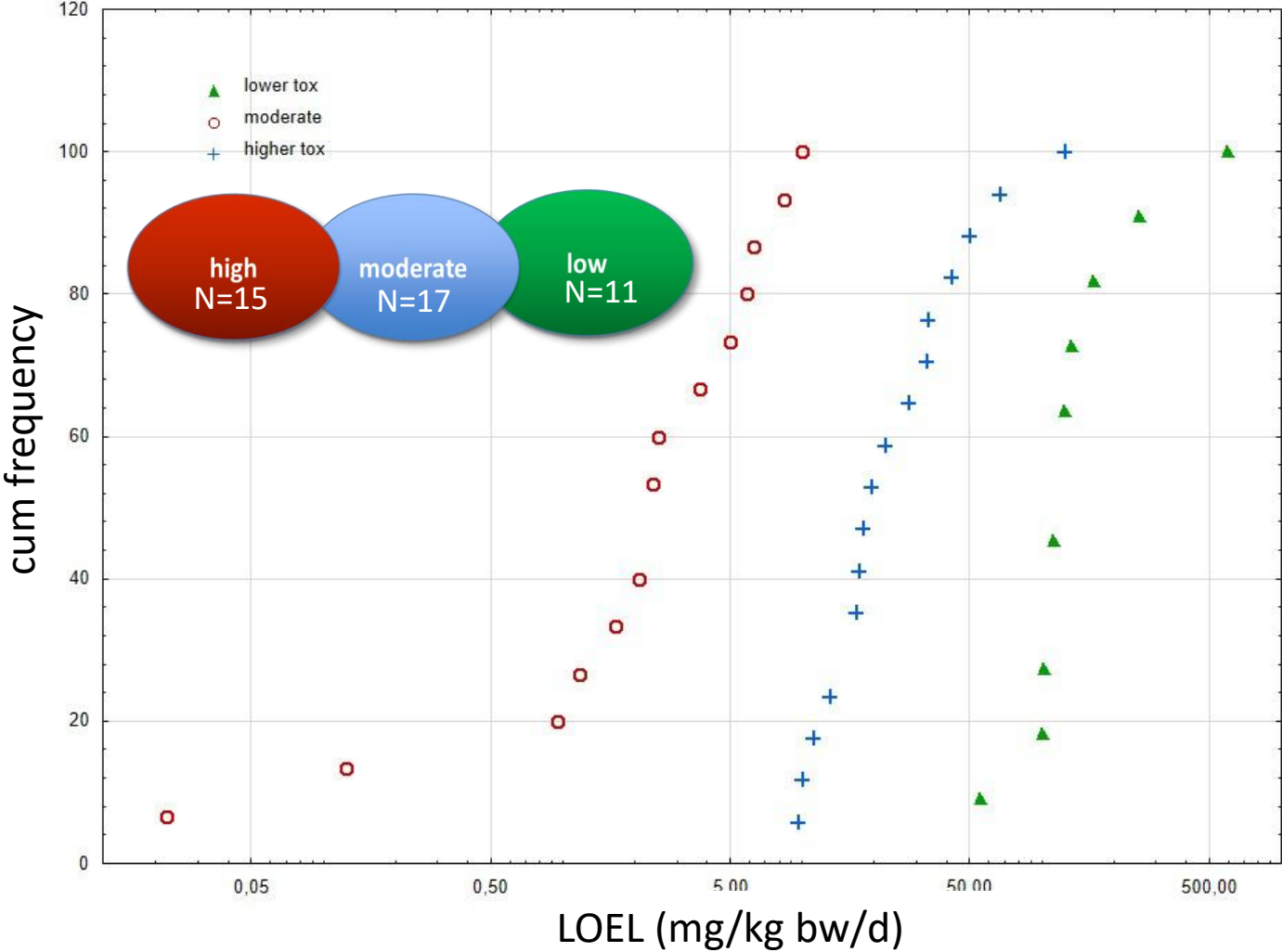
*In silico*

*In vitro*

Toxicokinetics

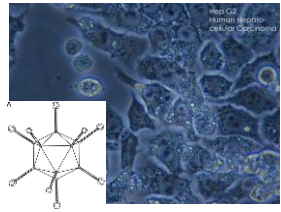
*In silico*

# Case study chemicals from different severity classes.



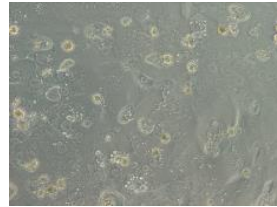
# Test systems involved in the case study to assess toxicodynamics.

## HepG2 AdvCYP



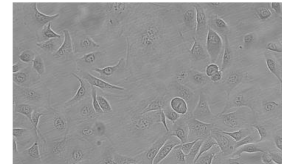
Organ system liver  
(metabolism)  
Partner: HULAFE

## PHH



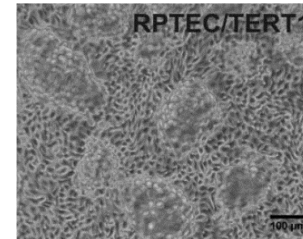
Organ system liver  
Partner: Cyprotex

## PBEC



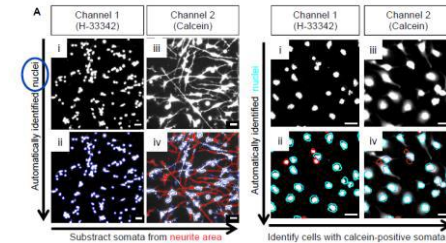
Organ system lung  
Partner: LUMC

## RPTEC



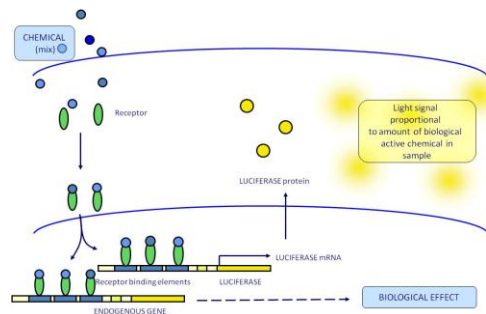
Organ system kidney  
Partner: VUA

## LUHMES

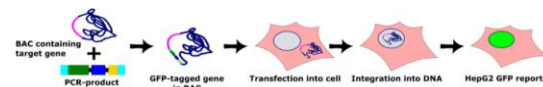


Organ system neuron  
Partner: UKN

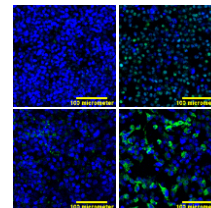
## CALUX



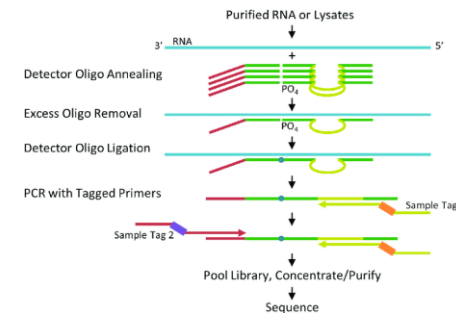
## HepG2 BAC-GFP



Oxidative stress \*SRXN1  
DNA damage  
ER stress  
Inflammatory stress  
Heat shock



## High throughput transcriptomics TempO-Seq (HTTr)



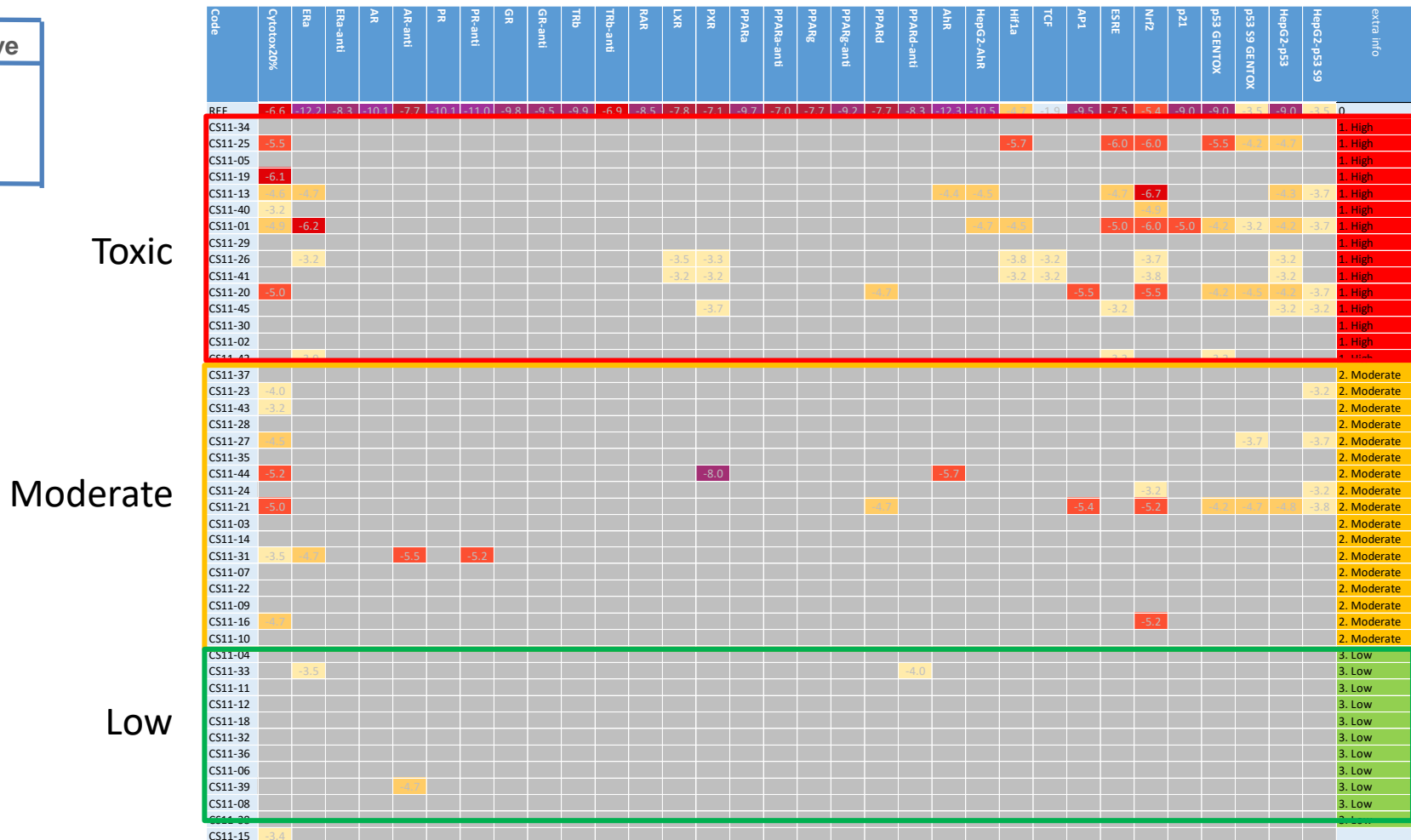
1. HepG2
2. PHH
3. PBEC
4. RPTEC
5. LUHMES



# CALUX – reporter assays

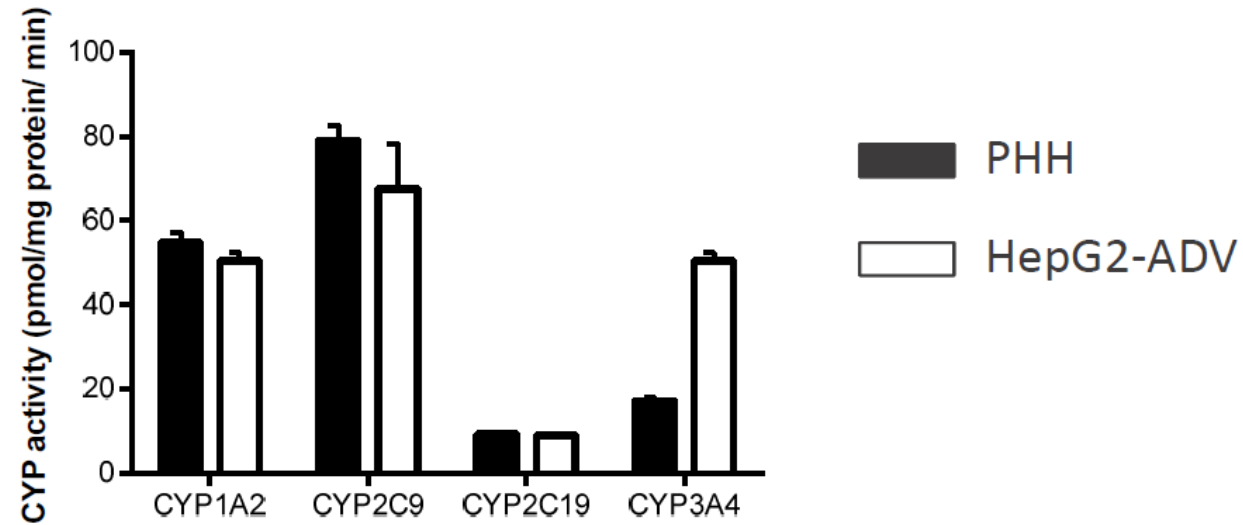
- Lowest observed effect concentrations (LOECs) in LogM

Category	Hits/c	% active
Toxic	3.8	66.7 %
Moderate	1.5	47.1 %
Low	0.3	18.2 %



## Metabolic retrofitting of HepG2 HCl assay

- We have successfully transfected simultaneously HepG2 cells with 4 different ADV encoding for CYP1A2, CYP2C9, CYP2C19 and CYP3A4

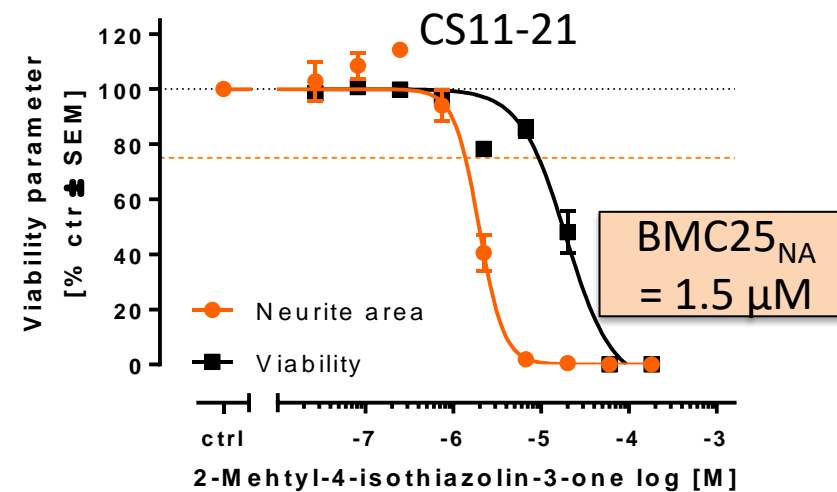
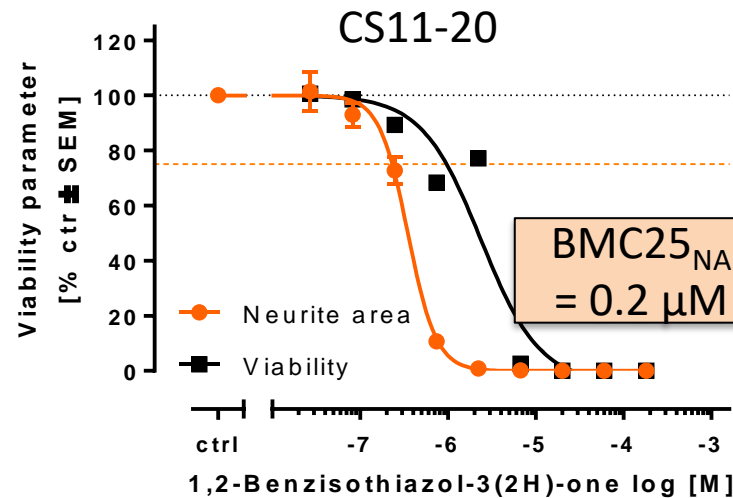
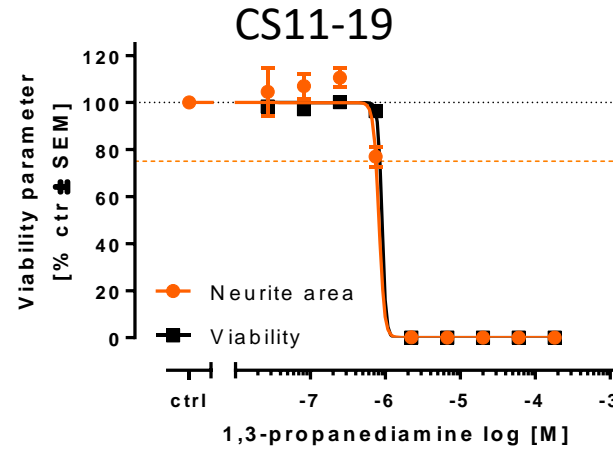


# Metabolic retrofitting of HepG2 HCl assay

			HepG2	HepG2 + Adv-CYP
CS11-19	2372-82-9	1,3-propanediamine, N-(3-aminopropyl)-N-dodecyl-	3	3
CS11-20	2634-33-5	1,2-Benzisothiazol-3(2H)-one	60	180
CS11-21	2682-20-4	2-Methyl-4-isothiazolin-3-one	60	180
CS11-22	2695-37-6	sodium 4-vinylbenzenesulphonate	>180	>180
CS11-23	2855-13-2	(5-Amino-1,3,3-trimethyl-1-cyclohexylmethyl)amine	180	180
CS11-24	2871-01-4	2-(4-amino-2-nitroanilino)ethanol	>180	60
CS11-25	3811-73-2	1-Oxide-2-pyridinethiol, Sodium salt	10	>180
CS11-26	3926-62-3	2-chloroacetic acid sodium salt (1:1)	>180	>180
CS11-27	4719-04-4	1,3,5-Triazine-1,3,5(2H,4H,6H)-triethanol	180	>180
CS11-28	552-30-7	1,2,4-benzenetricarboxylic anhydride	>180	60
CS11-29	56-93-9	benzyltrimethylammonium chloride	>180	>180
CS11-30	58-55-9	1,3-dimethyl-2,3,6,7-tetrahydro-1H-purine-2,6-dione	>180	>180

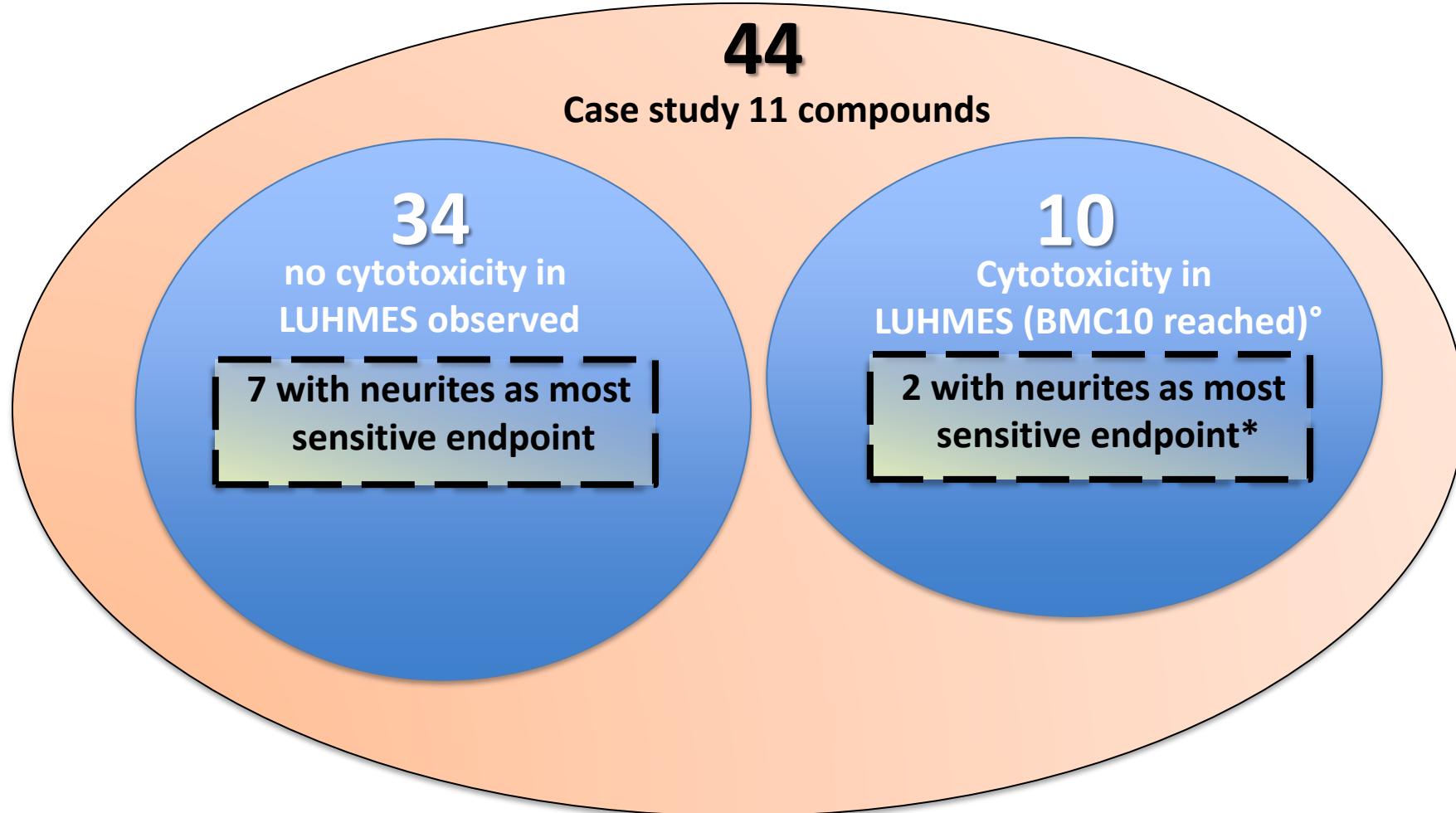
# Neurite outgrowth effects CS11 compounds

Example of cytotoxic classified compound:



Neurite endpoint ~5x more sensitive than viability for CS11-20 and CS11-21

# Summary neurite outgrowth phenotypic assay



# High throughput transcriptomics

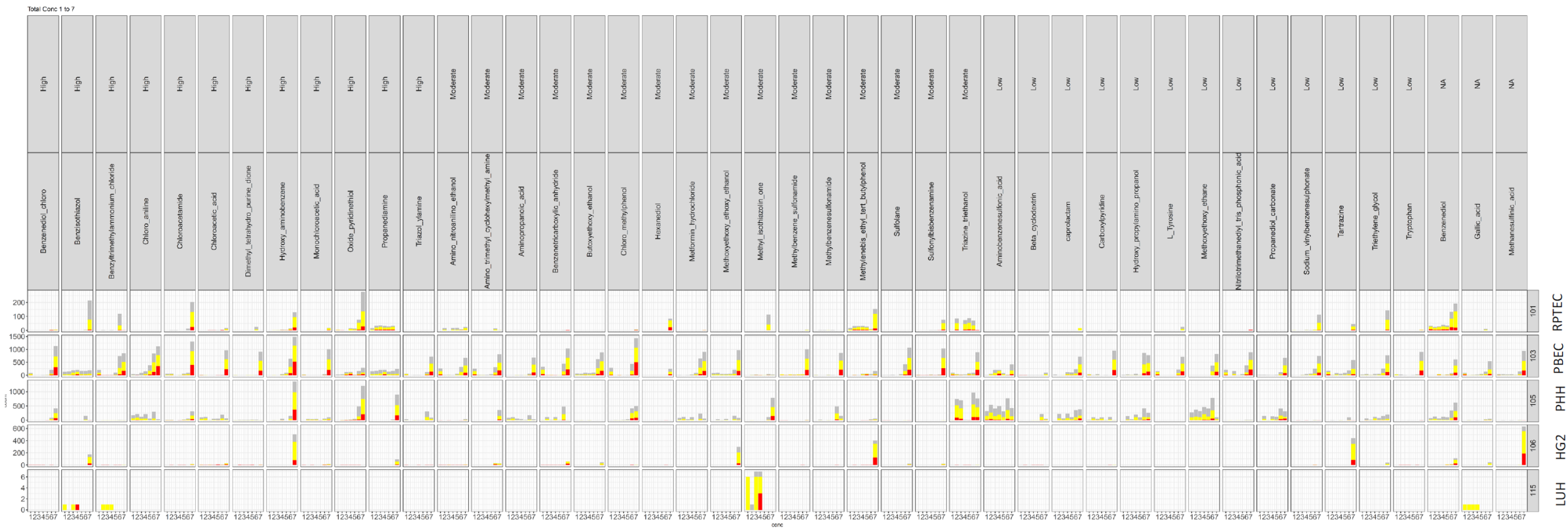
## Toxicodynamics - Hazard characterization

### Detailed MoA analysis

- High throughput transcriptomics (HepG2, PHH, PBEC, RPTEC, LUHMES)
- TempO-seq EU-ToxRisk gene panel (~3500 sentinel genes)

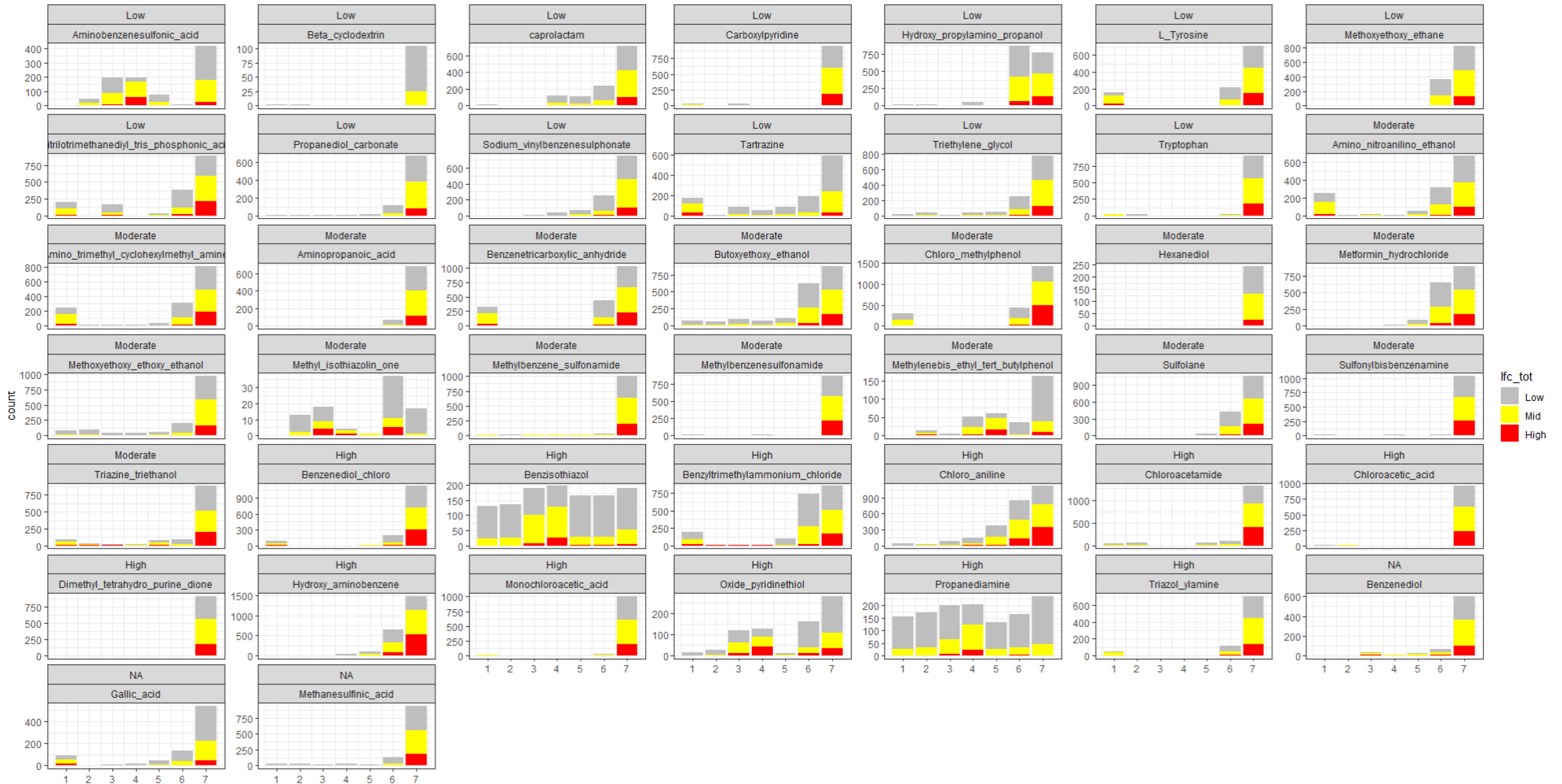
EUT	Partner	Model	Status	Samples
101	VU	RPTEC/TERT	Data analysis	1855
103	LUMC	PBEC	Data analysis	1923
105	CPX	PHH	Data analysis	1839
106	UL	HepG2	Data analysis	2064
115	UKN	LUHMES	Data analysis	945

# High throughput transcriptomics: CS11 compounds



# High throughput transcriptomics: lung PBEC dose response

EUT103 Conc 1 to 7





# Conclusions

- Target prediction models can identify candidates for some chemicals
  - Metabolic retrofitting critical to determine toxicity liability / potency.
  - Highly toxic chemicals show more profound effects in reporter assays.
  - Phenotypic assays (e.g. neurite outgrowth assay) are more sensitive than cytotoxicity.
  - HTTr is a sensitive method to determine biological perturbations.
- *Preliminary conclusion:* HT NAMs can contribute to prioritization of highly toxic chemicals.

Thanks to entire Case Study 11 team!!!

Thank you!!

