

BENCHMARK DOSE (BMD) MODELING OF IMAGE-BASED PHENOTYPIC PROFILING DATA YIELDS MORE POTENT ESTIMATES OF CHEMICAL BIOACTIVITY COMPARED TO CELL VIABILITY AND APOPTOSIS ASSAYS

Clinton Willis^{1,2}, Johanna Nyffeler^{2,3}, Joshua Harrill²

¹Oak Ridge Associated Universities (ORAU), Oak Ridge, TN.

²US Environmental Protection Agency, National Center for Computational Toxicology, Office of Research and Development, Research Triangle Park, NC.

³Oak Ridge Institute for Science and Education (ORISE), Oak Ridge, TN.

Clinton Willis | willis.clinton@epa.gov

Abstract

High-throughput imaging-based phenotypic profiling (HTPP) is a high-throughput chemical screening method that combines automated microscopy and image analysis to measure a large variety of morphological features at the single cell level. Here we describe workflows for concentration-response screening and image analysis using an HTPP assay that quantitatively evaluates changes in organelle morphology (i.e. Cell Painting) and calculation of *in vitro* points-of-departure (POD_{HTPP}) using high-throughput concentration-response modeling with the BMDExpress 2.2 software package. A set of 16 reference chemicals were tested in six human cell lines (U-2 OS, MCF-7, HTB-9, A549, ARPE-19, HepG2). Cells were plated in 384-well plates and after 24 h treated with 7 concentrations (semi-log spacing, $n = 3/\text{plate}$, 3 cultures) in a randomized pattern. After 48 h, cells were live labeled with MitoTracker (mitochondria), fixed, permeabilized and labeled with Hoechst-33342 (nuclei), SYTO14 (nucleoli) and fluorescent conjugates of concanavalin A (ER), phalloidin (cytoskeleton), and wheat germ agglutinin (Golgi/plasma membrane). A multiplexed cell viability (CV) and apoptosis (AP) assay was run in parallel. Confocal images were acquired using an Opera Phenix HCS system and analyzed using Harmony software, yielding ~1300 features per cell. Cell-level data were median absolute deviation (MAD) normalized to DMSO controls. BMD modeling was performed on well-level median values. Most chemicals ($n=14$) affected cell morphology. Distinct patterns of affected cellular features were observed across the chemical set and, in most cases, were consistent with observations from the literature. In general, the chemicals produced similar patterns, with highly correlated potency estimates, across the six different cell lines. For all compounds, Cell Painting BMDs (BMD_{CP}) were at least as sensitive as Cell Viability BMDs (BMD_{CV}). In some cell lines, BMD_{CP} were > 10x lower than BMD_{CV}. Screening of a larger set of chemicals ($n=420$) also demonstrated marked differences in BMD_{CP} and BMD_{CV} potency estimates. In summary, testing of diverse compounds yielded distinct patterns of affected features below the threshold for cytotoxicity, indicating that this profiling method could be used to derive *in vitro* potency estimates for screening level risk assessments. *This abstract does not necessarily reflect USEPA policy.*

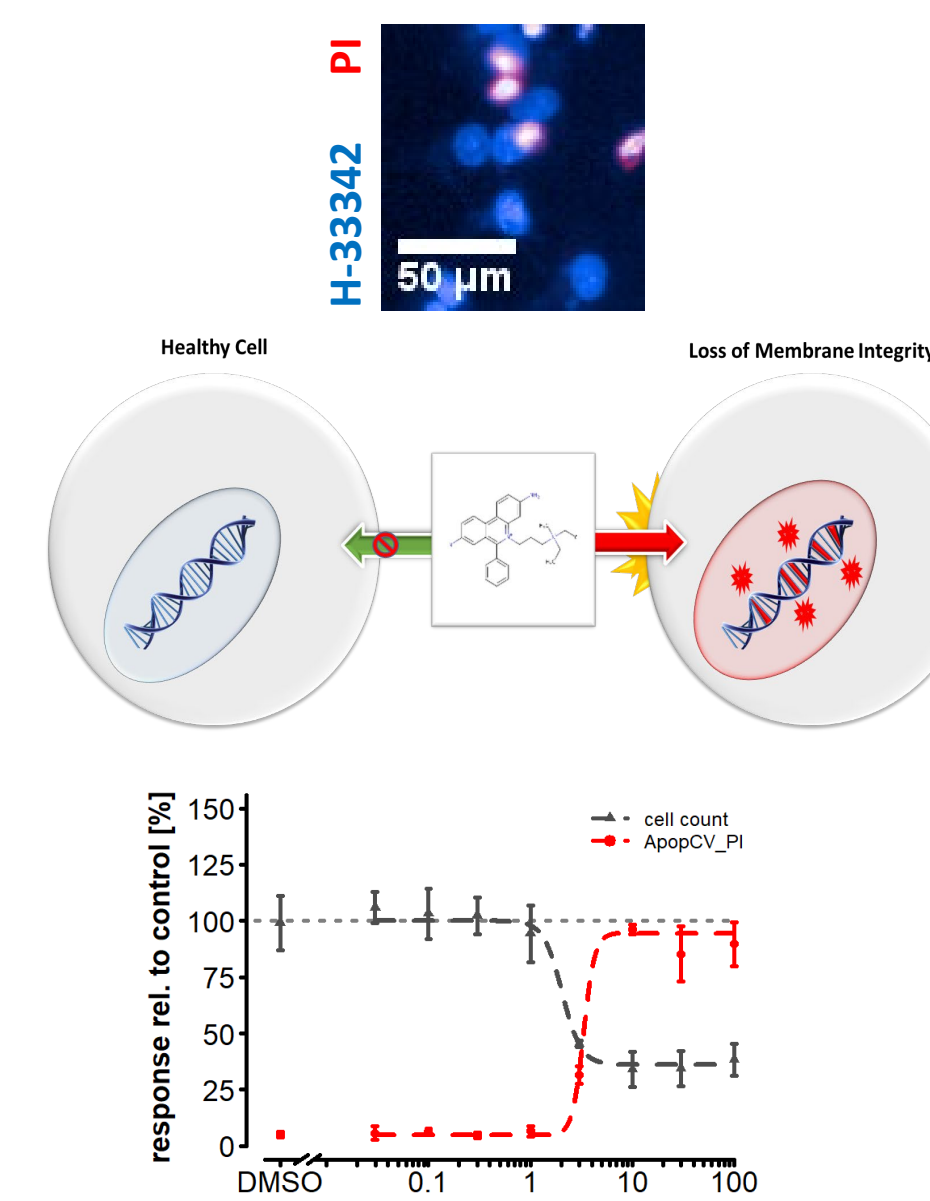
Background and Experimental Design

- HTPP is a chemical screening method that measures a large variety of morphological features of individual cells in *in vitro* cultures.
- Successfully used for functional genomic studies and in the pharmaceutical industry for compound efficacy and toxicity screening.
- May be used as an efficient and cost-effective method for evaluating the bioactivity of environmental chemicals.
- May be used to determine effect thresholds (i.e. *in vitro* point-of-departure, POD) for chemical bioactivity.

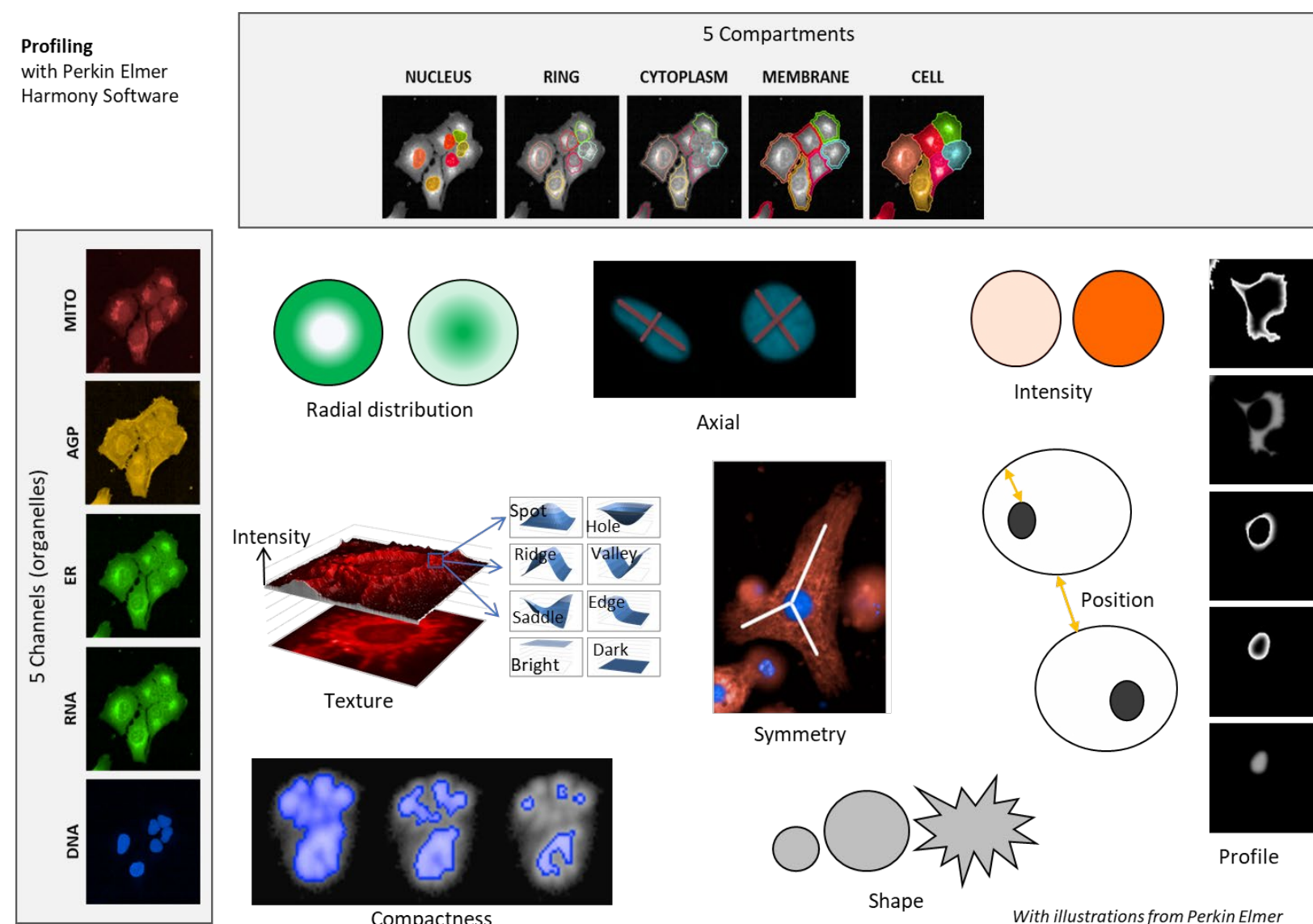
Experimental Design

Cell Types	U-2 OS, MCF-7, A549, HepG2, HTB-9, ARPE-19	Solvent controls / plate	3
Exposure Duration	48 hours	Replicates / plate	24
# Chemicals	14 phenotypic reference chemicals; 2 negatives	# Independent experiments	3
# Concentrations	7 (1/2 log ₁₀ spacing)		

Assay 1: Cell Viability

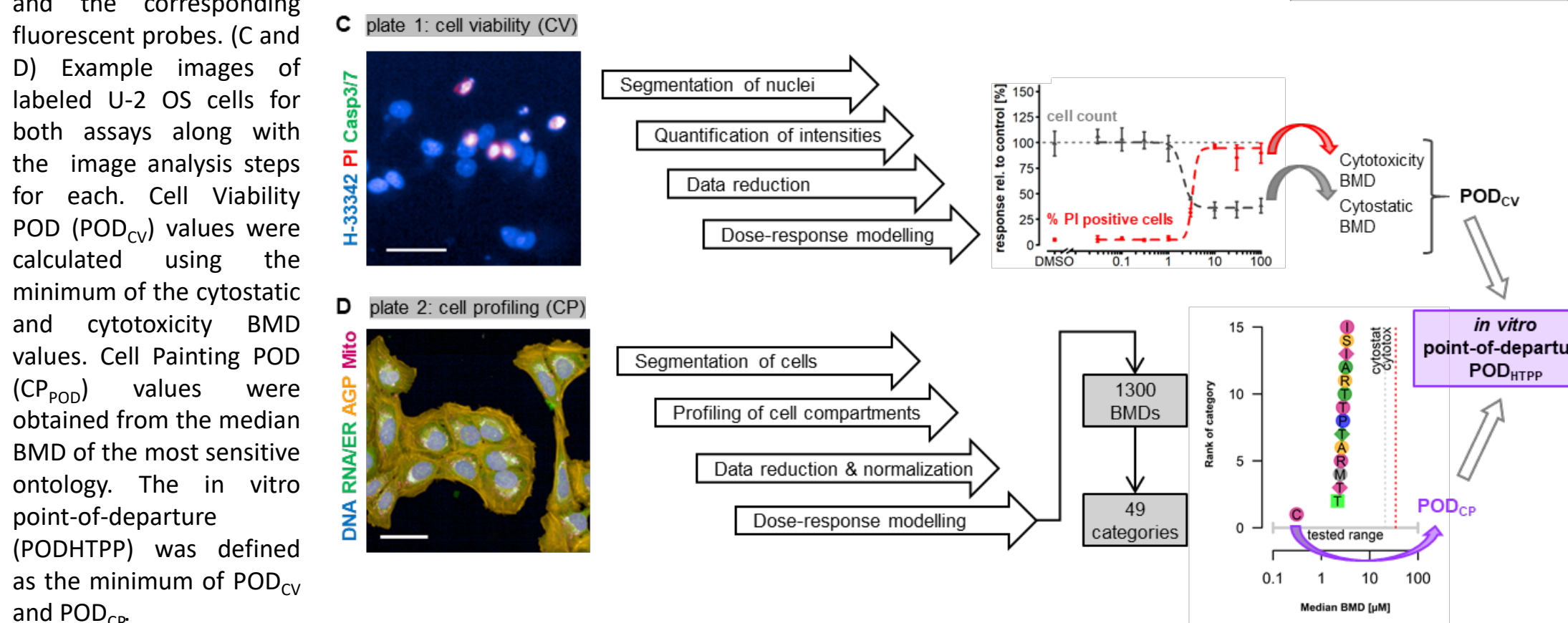


Assay 2: High Throughput Phenotypic Profiling (i.e. Cell Painting)

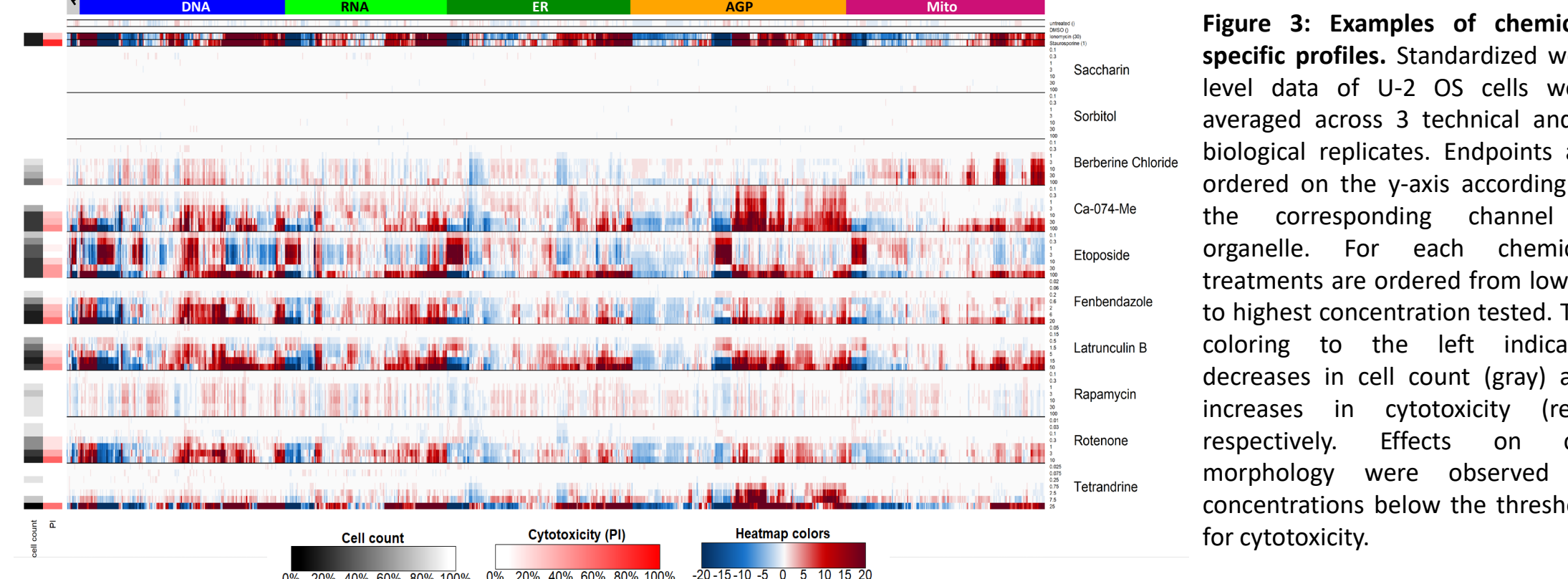
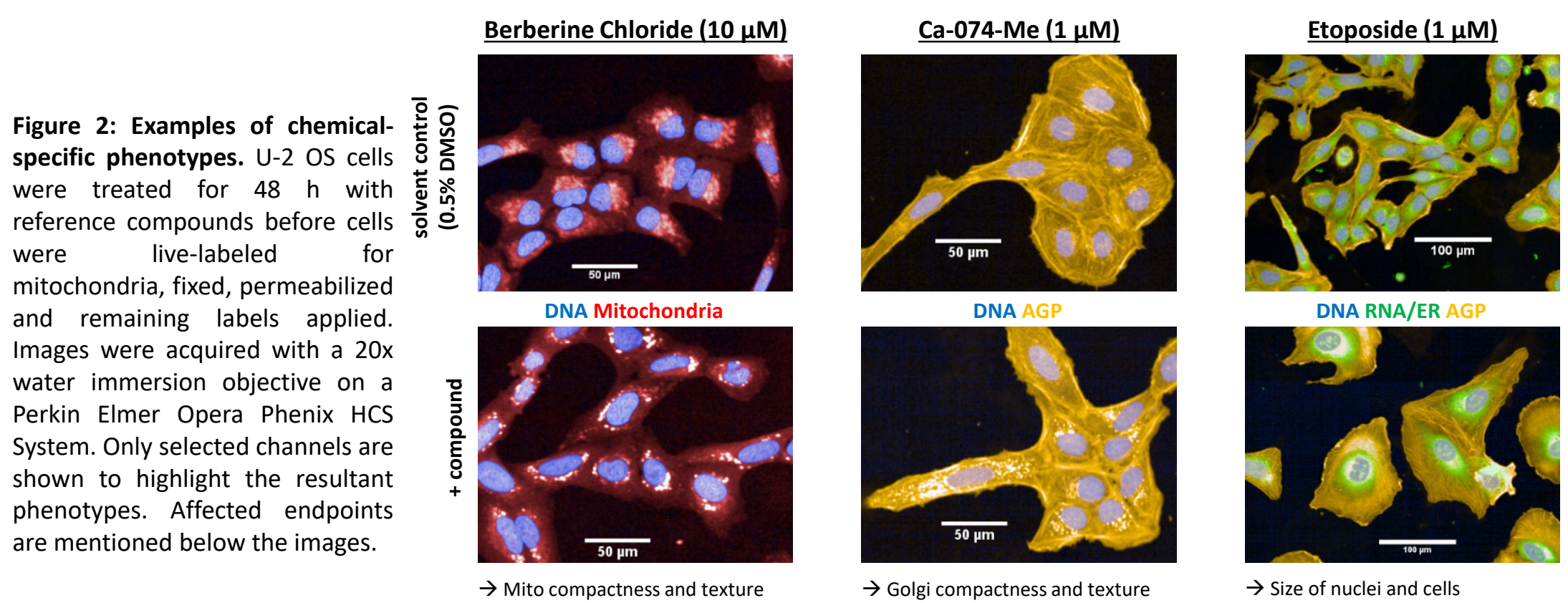


Methods

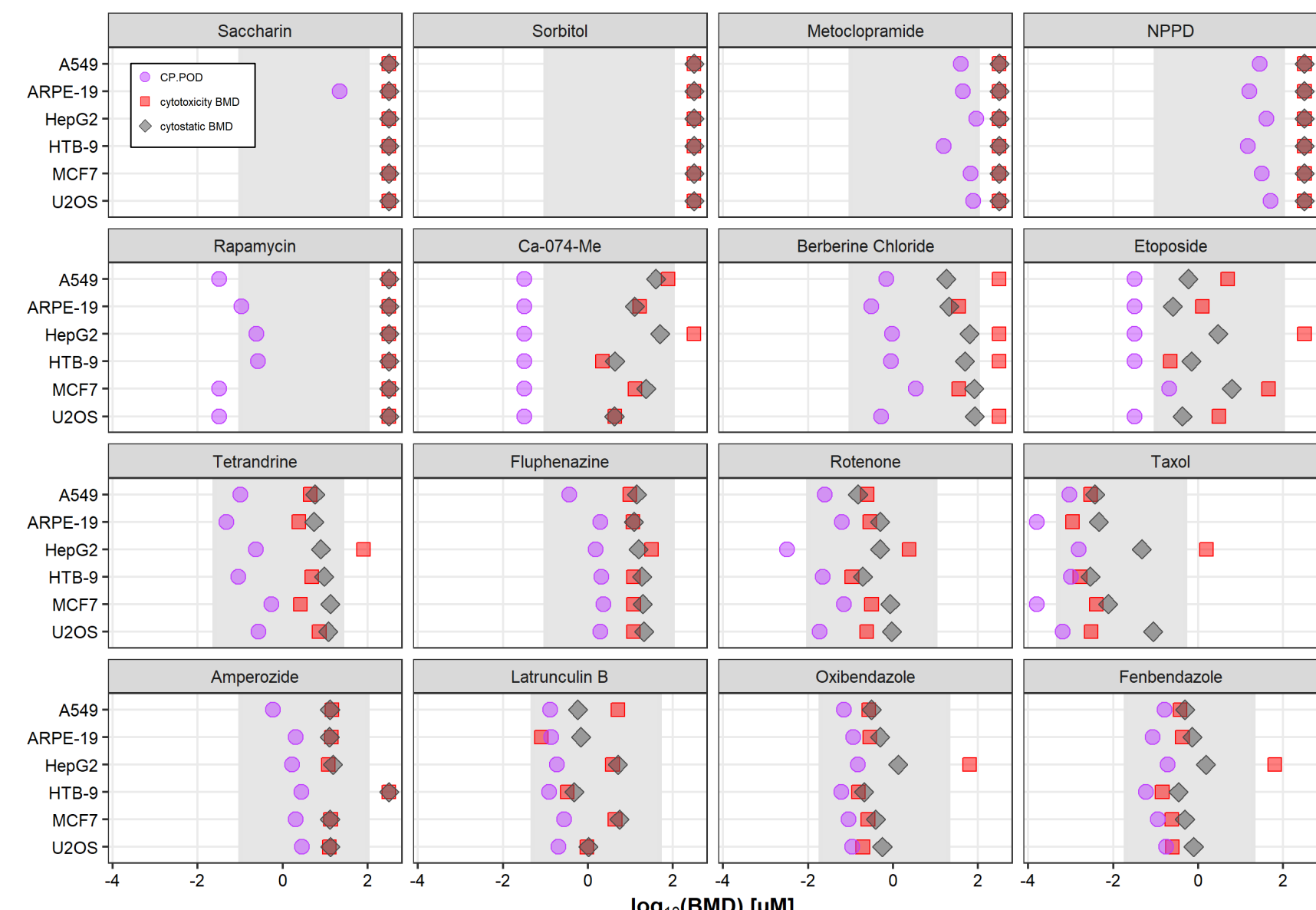
Figure 1: Assay and Data Analysis Workflows. (A) Diagram illustrating plate processing for both assay formats. (B) Table listing organelles labeled using the Cell Painting assay and the corresponding fluorescent probes. (C and D) Example images of labeled U-2 OS cells for both assays along with the image analysis steps for each. Cell Viability POD (POD_{CV}) values were calculated using the minimum of the cytostatic and cytotoxicity BMD values. Cell Painting POD (CP_{POD}) values were obtained from the median BMD of the most sensitive ontology. The *in vitro* point-of-departure (POD_{HTPP}) was defined as the minimum of POD_{CV} and POD_{CP}.



Results: Reference Chemical Phenotypes



Results: Reference Chemical PODs

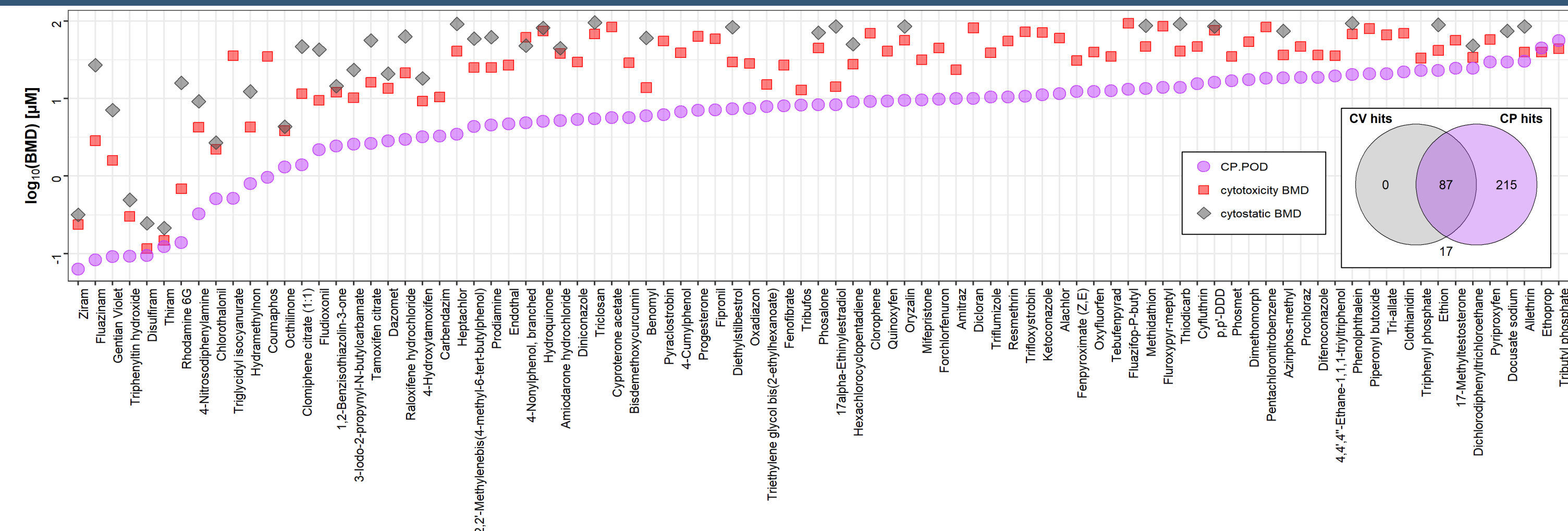


Cell Line	Description
A549	Human lung adenocarcinoma epithelial cells
ARPE-19	Human retinal pigment epithelial cell line
HepG2	Human liver carcinoma cells
HTB-9	Human urinary bladder epithelial cells
MCF-7	Human breast adenocarcinoma cell line
U-2 OS	Human bone osteosarcoma epithelial cells

Figure 4: The 16 reference chemicals were tested in 6 cell types. The PODs for the Cell Painting assay (POD_{CP}), cytotoxicity BMD and cytostatic BMD were compared. In all cases, POD_{CP} was less than the cytotoxicity or cytostatic PODs. In many cases, the POD_{CP} was left-shifted as compared to the cytotoxicity or cytostatic PODs by several orders of magnitude. For each reference chemical, the Cell Painting and cytotoxicity / cell count PODs were remarkably consistent across cell lines.

- ⇒ Cell Painting PODs < Cytotoxicity / Cytostatic PODs
- ⇒ PODs vary less than 2 order of magnitude across cell types.

Results: Chemical Screening



Conclusions and Future Directions

- ⇒ A set of phenotypic reference chemicals with distinct response profiles and comparable potency across multiple cell types was identified.
- ⇒ The morphological phenotypes of the reference chemicals were consistent with those reported in the scientific literature for the Cell Painting assay.
- ⇒ The HTPP assay was able to replicate the morphological phenotypes for a set of phenotypic reference chemicals identified from the scientific literature.
- ⇒ In a screen of several hundred chemicals, HTPP detected concentration-dependent changes in cell morphology with a hit rate of ~95%.
- ⇒ Future directions include expanding the number of chemicals, and the number of cell types, screened using HTPP and incorporating the resulting information into existing approaches for hazard binning and prioritization using *in vitro* screening data.