# Practical Application of New Approach Methods in Developmental and Reproductive Toxicity Testing

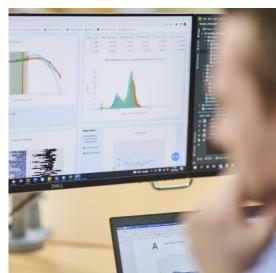
Dr Predrag Kukic Unilever Safety and Environmental Assurance Centre

Theme: NAMs as Problem-Solvers 23<sup>th</sup> October 2023











### **Outline**

- Overview of Unilever's NGRA Framework for DART testing
- Biological relevance of the NGRA Framework for DART testin
- Case studies / fit for purpose validation, next steps





# Unilever Policy & Approach Safe & Sustainable Products without Animal Testing

#### What we believe

- Every Unilever product must be safe for people and our environment
- Non-animal testing to assess ingredient & product safety – there are a wide range of non-animal alternatives grounded in modern science and new technology

#### How we do it







70+ collaborations



600+ publications



















# A paradigm shift is underway as use of non-animal safety science increases & safety assessment frameworks evolve to embed NAMs

> Non-animal safety science is increasingly being used to make decisions on **consumer safety**, **safety of workers**, and safety of **people and non-human species** in the **environment**.

Regulatory Animal Testing of Chemicals is increasingly seen as unjustifiable / unethical by the majority of society

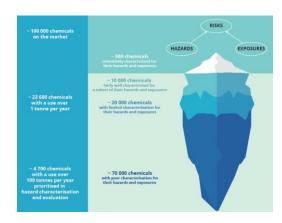
Aug 2021 – Aug 2022: 1.4M+ signatures





NAMs to fully replace the need for chemical regulatory animal testing

High throughput – more testing before the chemical is put on the market, data reuse, etc.



Potential to address information requirements for all substances in the market

Move to more sustainable sources of chemicals (e.g. bio-based) is transforming chemical innovation & use



Potential to ensure new chemicals are Safe & Sustainable by Design



# A paradigm shift is underway as use of non-animal safety science increases & safety assessment frameworks evolve to embed NAMs

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High throughput – more testing before the chemical is put on the market, data

**Human-relevant** 

Move to more sustainable sources of chemicals (e.g. bio-based) is

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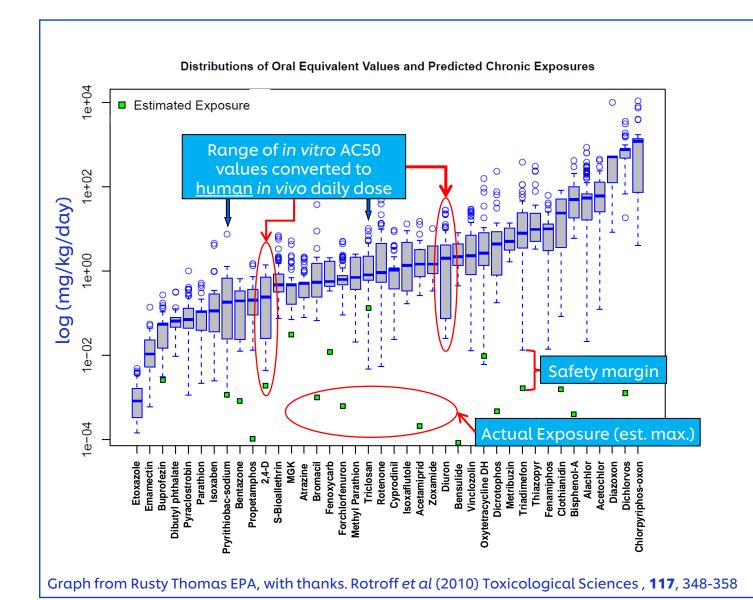


NAMs to fully replace the need for chemical regulatory animal testing Potential to address information requirements for all substances in the market

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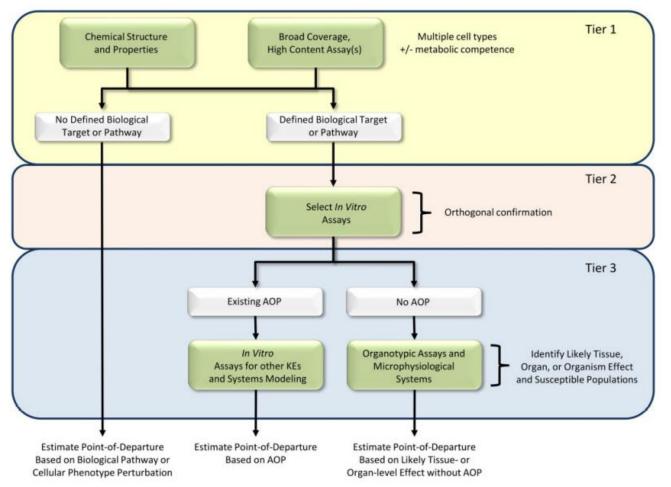
# **US EPA Next Generation Blueprint Tiered Testing Framework**



- NGRA is defined as an exposureled, hypothesis-driven risk assessment approach that integrates New Approach Methodologies (NAMs) to assure safety without the use of animal testing
- If there is no bioactivity observed at consumer-relevant concentrations, there can be no adverse health effects.
- If there is bioactivity observed at consumer-relevant concentrations, follow up testing is required to establish if that could result in an adverse effect
- At no point does NGRA attempt to predict the results of high dose toxicology studies in animals.



# **US EPA Next Generation Blueprint Tiered Testing Framework**



Unilever

Figure 2. Tiered testing framework for hazard characterization. Tier 1 uses both chemical structure and broad coverage, high content assays across multiple cell types for comprehensively evaluating the potential effects of chemicals and grouping them based on similarity in potential hazards. For chemicals from Tier 1 without a defined biological target / pathway, a quantitative point-of-departure for hazard is estimated based on the absence of biological pathway or cellular phenotype perturbation. Chemicals from Tier 1 with a predicted biological target or pathway are evaluated Tier 2 using targeted follow-up assays. In Tier 3, the likely tissue, organ, or organism-level effects are considered based on either existing adverse outcome pathways (AOP) or more complex culture systems. Quantitative points-of-departure for hazard are estimated based on the AOP or responses in the complex culture system.







TOXICOLOGICAL SCIENCES, 169(2), 2019, 317-332

doi: 10.1093/toxsci/kfz058 Advance Access Publication Date: March 5, 201

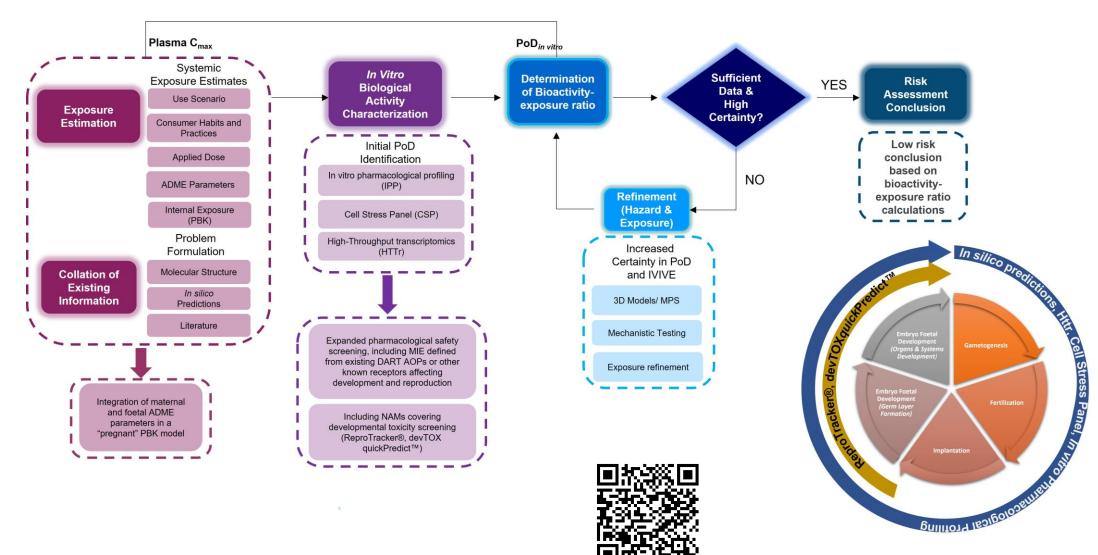
FORUM

# The Next Generation Blueprint of Computational Toxicology at the U.S. Environmental Protection Agency

Russell S. Thomas,\*,¹ Tina Bahadori,† Timothy J. Buckley,‡ John Cowden,\* Chad Deisenroth,\* Kathie L. Dionisio,‡ Jeffrey B. Frithsen,§ Christopher M. Grulke,\* Maureen R. Gwinn,\* Joshua A. Harrill,\* Mark Higuchi,¶ Keith A. Houck,\* Michael F. Hughes,¶ E. Sidney Hunter, III,¶ Kristin K. Isaacs,‡ Richard S. Judson,\* Thomas B. Knudsen,\* Jason C. Lambert,∥ Monica Linnenbrink,\* Todd M. Martin,∥ Seth R. Newton,‡ Stephanie Padilla,¶ Grace Patlewicz,\* Katie Paul-Friedman,\* Katherine A. Phillips,‡ Ann M. Richard,\* Reeder Sams,\* Timothy J. Shafer,¶ R. Woodrow Setzer,\* Imran Shah,\* Jane E. Simmons,¶ Steven O. Simmons,\* Amar Singh,\* Jon R. Sobus,‡ Mark Strynar,‡ Adam Swank,‡ Rogelio Tornero-Valez,‡ Elin M. Ulrich,‡ Daniel L. Villeneuve,∥∥ John F. Wambaugh,\* Barbara A. Wetmore,‡ and Antony J. Williams\*

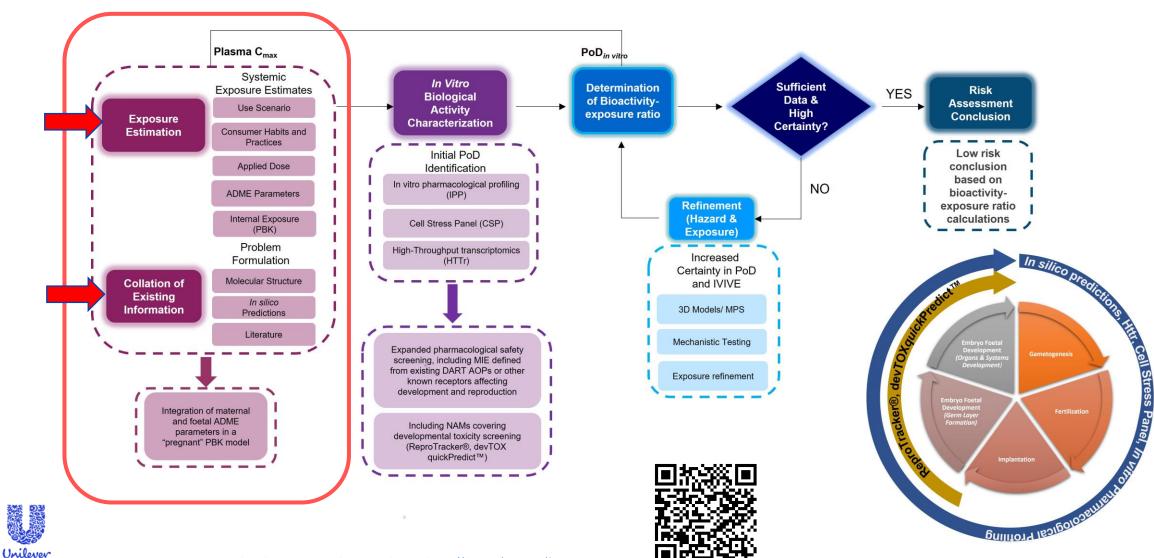
National Center for Computational Toxicology, U.S. Environmental Protection Agency, Research Triangle Park, NC 27711, 'National Center for Environmental Assessment, U.S. Environmental Protection Agency, Washington, D.C. 20004, 'National Exposure Research Laboratory, U.S. Environmental Protection Agency, Research Triangle Park, NC 27711, <sup>\$</sup>Chemical Safety for Sustainability National Research Program, U.S. Environmental Protection Agency, Washington, D.C. 20004, <sup>\$\$</sup>National Health and Environmental Effects Research Laboratory, U.S. Environmental Protection Agency, Research Triangle Park, NC 27711, <sup>\$\$</sup>National Center for Environmental Assessment, U.S. Environmental Protection Agency, Cincinnati, OH 45220,

# NGRA Framework for DART – tiered approach

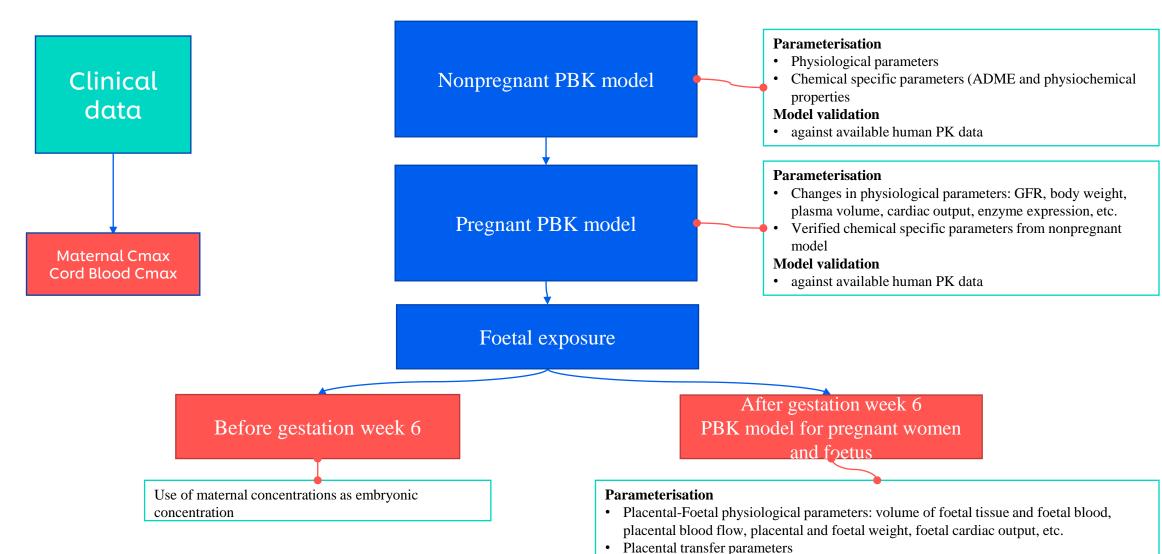




# NGRA Framework for DART - exposure module



# NGRA Framework for DART – exposure module



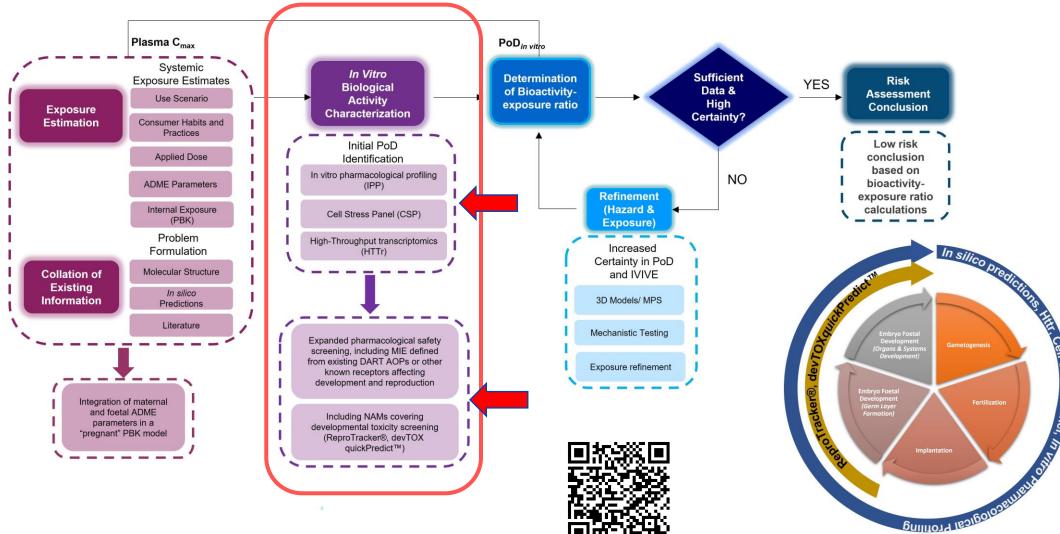


Rajagopal et al., Front. Toxicol., 07 March 2022 https://doi.org/10.3389/ftox.2022.838466

Model validation

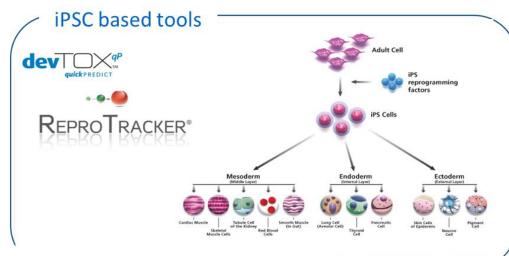
against available human PK data

# NGRA Framework for DART - bioactivity module





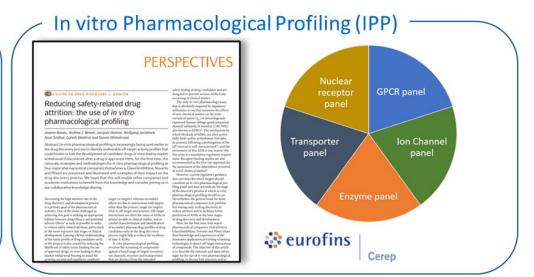
# NGRA Framework for DART - bioactivity module



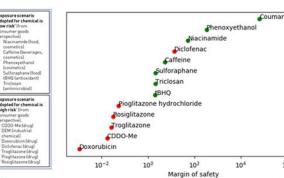
-Toxicology in Vitro (2020), 63, 104746

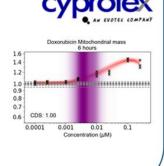
Calculated BMD mean value (µM)

# High-throughput Transcriptomics (HTTr) Use of full human gene panel ~ 21k 24 hrs exposure 7 concentrations 3 cell lines HepG2/ HepaRG/MCF7 3D HepaRG spheroid BMDexpress 2 BMDexpress 2 HepaRG 2D — HepaRG 2D — HepG2 Cytochrome P450arranged by substrate type Phase I Functionalization of compounds





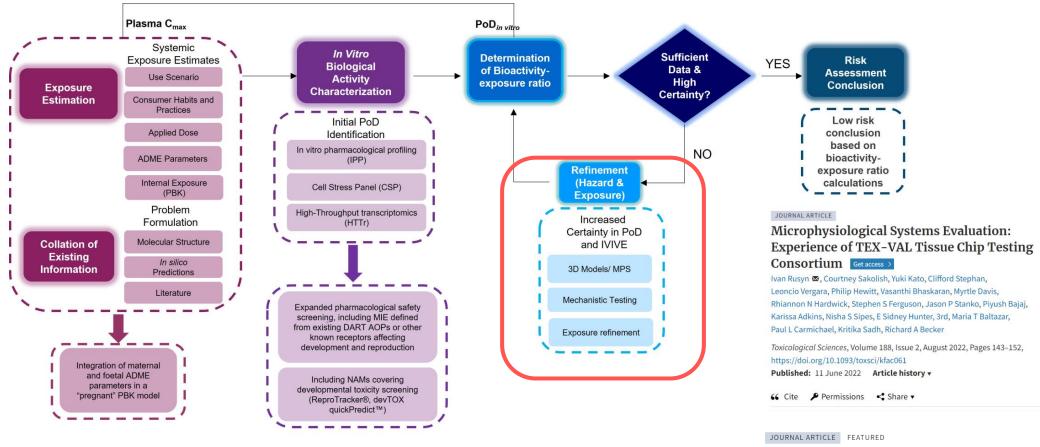






Toxicol Sci (2020), 176, 11-33

## Refinement of Biological Activity and Exposure



- > Tex-Val: public-private collaboration established for testing of diverse microphysiological system
- > Use of metabolically competent models (cell lines, alginate immobilization, etc)



The Alginate Immobilization of Metabolic Enzymes Platform Retrofits an Estrogen Receptor Transactivation Assay With Metabolic Competence

Chad Deisenroth 

, Danica E DeGroot 

, Todd Zurlinden, Andrew Eicher,
James McCord, Mi-Young Lee 

, Paul Carmichael, Russell S Thomas

Toxicological Sciences, Volume 178, Issue 2, December 2020, Pages 281–301, https://doi.org/10.1093/toxsci/kfaa147

Published: 29 September 2020

# NGRA Framework for DART – Scientific and Technical challenges

- > Metabolic capacity of the framework (cell models, MPS, alginate technology, etc.)
- > Spatio-temporal complexity of developmental and reproductive processes
- Short duration exposures and extrapolation to chronic effects
- > Ability to generate reliable and consistent reproducible results (HTTr, cell line variability, cell stress, IPP, reprotracker)
- > Complex data interpretation and uncertainty analysis
- > Coverage of important cellular and intercellular processes biological relevance
- > Chemical domain of applicability / case studies need for a flexible and fit for purpose validation



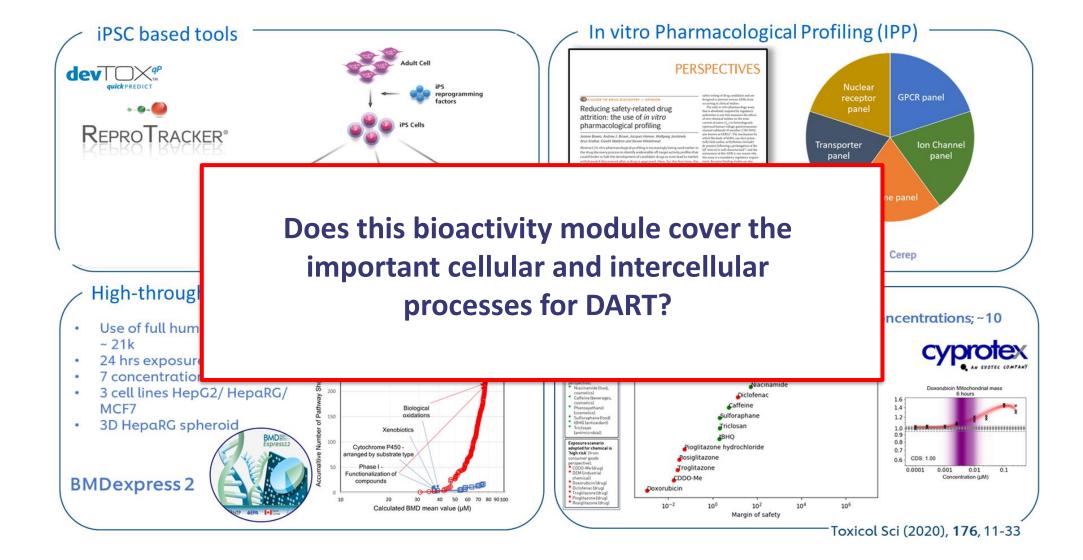
# Biological relevance of the NGRA Framework for DART



van der Zalm et al. Archives of Toxicology (2022) 96:2865–2879



## Coverage of important cellular and intercellular processes for DART





# Key Biomarkers for DART - Systematic literature search

# AOPs based approach

- 11 DART-related Adverse Outcome Pathways (AOPs)
- At present, a decision framework based only on AOPs is not feasible. However, AOPs can used as a knowledge base for enhancing a testing strategy

List of key stages, morphogen etic events, organ or organ systems

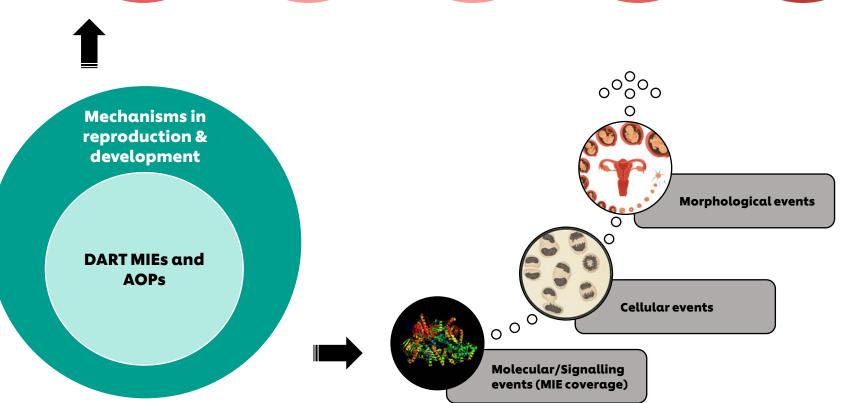
Targeted
literature
search for
cellular and
molecular
mechanism
s

Extraction of key biomarker terms for each stage, including any related to xenobiotic stress

Pooling all biomarker terms to generate master content

master content, evaluation of biological coverage of the NAMs and potential gaps

Using the





# Key Stages, Morphogenetic Events and Derivatives Organs & Systems in Human Reproduction and Development

#### Sex determination

#### Gametogenesis

#### **Fertilization**

#### **Zygote formation**

#### **Implantation**

#### **Blastulation**

#### Gastrulation

#### Placenta formation

#### **Neurulation**

#### Ectoderm formation and its derivatives

- Central nervous system
- Peripheral nervous system
- Autonomous nervous system
- Integumentary system

#### Mesoderm formation and its derivatives

- Somitogenesis
- Hematopoiesis
- Heart and circulatory system
- Immune system
- Spleen
- Urinary system and urethra
- Reproductive system testis
- Reproductive system ovary
- Skeletal system
- Limbs

#### **Endoderm formation and its derivatives**

- Digestive system
- Respiratory system
- Thymus
- Parathyroid
- Thyroid

#### Structures developing from mesenchyme or multiple germ layers

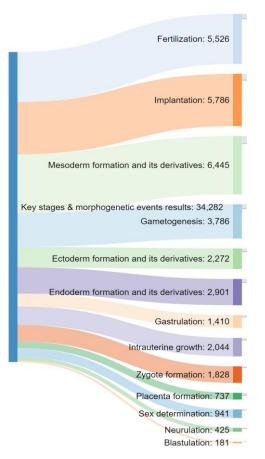
- Adrenal glands
- Eyes
- Ears
- Face and neck



# Overview of Literature Search and Extraction of Key Markers Information

Literature search MeSH Ontology 37 million Articles

Validation and quality check of results; finalising the articles



34,308 articles on key stages and morphogenetic events 69,299 articles on organs and organ systems development

Central nervous system: 6,755

Autonomic nervous system: 2,119

Heart and circulatory system: 1,711

Peripheral nervous system: 831

Skeletal system: 1,175

Respiratory system: 2,012

Digestive system: 2,244

Somitogenesis: 1,389

Adrenal glands: 2,193

Ears: 2,733

Immune system: 2,212

Urinary system: 1,470

Parathyroid: 910 -

Face and neck: 868
Thyroid: 1,002

Reproductive system – Testis: 6,078

Limbs: 6,061

Integumentary system: 4,282

Hematopoiesis: 5,157

Spleen: 2,935

Reproductive system – Ovary: 5,575

Derivative organs & organ systems: 69,412

Thymus: 4,087

Semantic enrichment using HGNC, miRNA and biological processes ontologies

Abstracts extracted and collated

#### Summary

PAXIP1 Potentiates the Combination of WEE1 Inhibitor AZD1775 and Platinum Agents in Lung Cancer The DNA damage response (DDR) involves a complex network of signaling events mediated by modular protein domains such as the BRCA1 C-terminal (BRCT) domain. Thus, proteins that interact with BRCT domains and are a part of the DDR constitute potential targets for sensitization to DNA-damaging chemotherapy agents. We performed a pharmacologic screen to evaluate 17 kinases, identified in a BRCT-mediated interaction network as targets to enhance platinum-based chemotherapy in lung cancer. Inhibition of mitotic kinase WEE1 was found to have the most effective response in combination with platinum compounds in lung cancer cell lines. In the BRCT-mediated interaction network, WEE1 was found in complex with PAXIP1, a protein containing six BRCT domains involved in transcription and in the cellular response to DNA damage. We show that PAXIP1 BRCT domains regulate WEE1-mediated phosphorylation of CDK1. Furthermore, ectopic expression of PAXIP1 promotes enhanced caspase-3 mediated apoptosis in cells treated with WEE1 inhibitor AZD1775 (formerly, MK-1775) and cisplatin compared with cells treated with AZD1775 alone. Cell lines and patient-derived xenograft models expressing both PAXIP1 and WEE1 exhibited synergistic effects of AZD1775 and cisplatin. In summary, PAXIP1 is involved in sensitizing lung cancer cells to the WEE1 inhibitor AZD1775 in combination with platinum-based treatment. We propose that WEE1 and PAXIP1 levels may be used as mechanism-based biomarkers of response when WEE1 inhibitor AZD1775 is combined with DNA-damaging agents.

Query run: ("CNS") AND (embryonic development OR fetal development) AND (cell physiology OR nervous system physiology) OR (signalling OR pathway OR gene OR protein) AND (human OR mammalian) NOT (infections)



## Pooled List of DARS biomarkers

#### 3551 DARS Genes

| 4  | А           | В   | С        |
|----|-------------|---|----------|
| 1  | Gene symbol | Name  | HitCount |
| 2  | CGA         | glycoprotein hormones, alpha polypeptide      | 11924    |
| 3  | SHH         | sonic hedgehog                                | 6622     |
| 4  | WNT1        | Wnt family member 1                           | 6428     |
| 5  | TGFB1       | transforming growth factor beta 1             | 6056     |
| 6  | IGF1        | insulin like growth factor 1                  | 4556     |
| 7  | INS         | insulin                                       | 4395     |
| 8  | GNRH1       | gonadotropin releasing hormone 1              | 3943     |
| 9  | CTNNB1      | catenin beta 1                                | 3912     |
| 10 | VEGFA       | vascular endothelial growth factor A          | 3777     |
| 11 | SRY         | sex determining region Y                      | 3479     |
| 12 | POMC        | proopiomelanocortin                           | 3454     |
| 13 | EGF         | epidermal growth factor                       | 3396     |
| 14 | KIT         | KIT proto-oncogene receptor tyrosine kinase   | 3380     |
| 15 | POU5F1      | POU class 5 homeobox 1                        | 3307     |
| 16 | CD4         | CD4 molecule                                  | 3152     |
| 17 | PAX6        | paired box 6                                  | 3124     |
| 18 | LIF         | LIF, interleukin 6 family cytokine            | 3070     |
| 19 | BMP4        | bone morphogenetic protein 4                  | 3027     |
| 20 | CD34        | CD34 molecule                                 | 3027     |
| 21 | ESR1        | estrogen receptor 1                           | 2946     |
| 22 | SOX9        | SRY-box 9                                     | 2649     |
| 23 | TNF         | tumor necrosis factor                         | 2620     |
| 24 | TP53        | tumor protein p53                             | 2520     |
| 25 | PTHLH       | parathyroid hormone like hormone              | 2436     |
| 26 | AMH         | anti-Mullerian hormone                        | 2431     |
| 27 | NR5A1       | nuclear receptor subfamily 5 group A member 1 | 2341     |
| 28 | IGF2        | insulin like growth factor 2                  | 2290     |
| 29 | LEP         | leptin  | 2058     |
| 30 | AKT1        | AKT serine/threonine kinase 1                 | 1977     |
| 31 | FGF2        | fibroblast growth factor 2                    | 1912     |

# 474 DARS Biological Processes

|    | Α           | В                                    | C        |
|----|-------------|--------------------------------------|----------|
| 1  | HitID       | Name                                 | HitCount |
| 2  | GO_0023052  | signaling                            | 21733    |
| 3  | GO_0007049  | cell cycle                           | 3228     |
| 4  | GO_0008219  | cell death                           | 2514     |
| 5  | GO_0006306  | DNA methylation                      | 2440     |
| 6  | GO_0001837  | epithelial to mesenchymal transition | 2422     |
| 7  | GO_0016310  | phosphorylation                      | 2372     |
| 8  | GO_0030154  | cell differentiation                 | 2262     |
| 9  | GO_0048468  | cell development                     | 2248     |
| 10 | GO_0001556  | oocyte maturation                    | 1973     |
| 11 | GO_0022008  | neurogenesis                         | 1567     |
| 12 | GO_0006412  | translation                          | 1541     |
| 13 | NCIT_C17741 | Oxidative Stress                     | 1449     |
| 14 | GO_0048477  | oogenesis                            | 1243     |
| 15 | GO_0001171  | reverse transcription                | 1235     |
| 16 | GO_0016477  | cell migration                       | 1209     |
| 17 | GO_0007165  | signal transduction                  | 1146     |
| 18 | GO_0030218  | erythrocyte differentiation          | 1134     |
| 19 | GO_0016049  | cell growth                          | 1041     |
| 20 | GO_0006914  | autophagy                            | 1021     |

#### 338 DARS miRNA

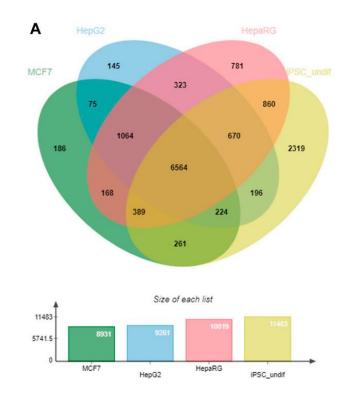
|    | А         | В        |
|----|-----------|----------|
| 1  | HitID     | HitCount |
| 2  | LET7      | 155      |
| 3  | MIR-21    | 127      |
| 4  | MIR-145   | 85       |
| 5  | MIR-125B  | 73       |
| 6  | MIR-17    | 73       |
| 7  | MIR-17-92 | 65       |
| 8  | MIR-1     | 64       |
| 9  | MIR-302   | 62       |
| 10 | MIR-124   | 56       |
| 11 | MIR-29B   | 55       |
| 12 | MIR-34C   | 52       |
| 13 | MIR-34A   | 51       |
| 14 | MIR-130B  | 51       |
| 15 | MIR-375   | 49       |
| 16 | MIR-200C  | 46       |
| 17 | MIR-24    | 45       |
| 18 | MIR-29A   | 44       |
| 19 | MIR-429   | 41       |
| 20 | MIR-223   | 41       |

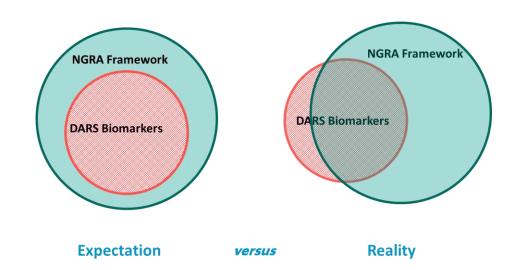


# Coverage of important DART biomarkers using Literature Search

HepG2, MCF-7, HepaRG, hiPSCs







Differentiated hiPSCs not included in this study but in scope for future work

#### Gaps

- 41 GPCRs (6 present in IPP)
- 60 HTH transcription factors (mainly homeobox transcription factors)
- Intercellular signal molecules (chemokines, cytokines, growth factors, neurotropic factors, peptide hormones)



Filling the gaps – work in progress: placenta transfer measurements, DNT, DIT, studying epigenetics in germline development, advanced cell models for refinement.

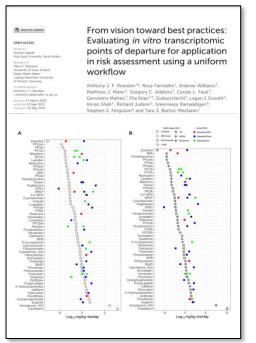
# Case studies / fit for purpose validation, next steps

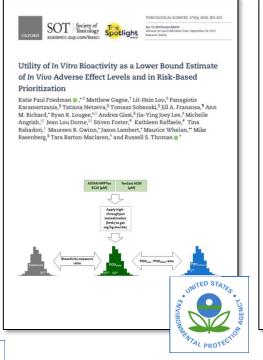


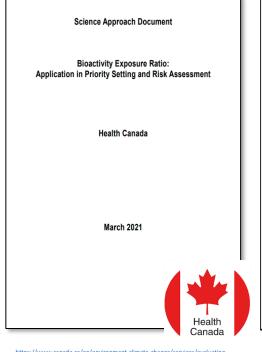
van der Zalm et al. Archives of Toxicology (2022) 96:2865–2879



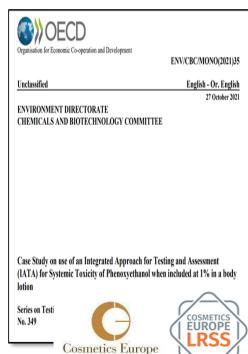
# Examples of ongoing or completed case studies for NAM/NGRA BER based risk assessment or prioritisation

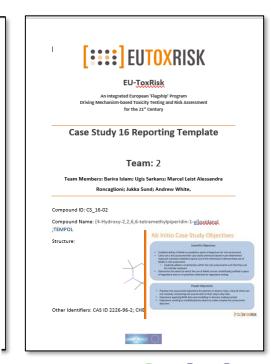












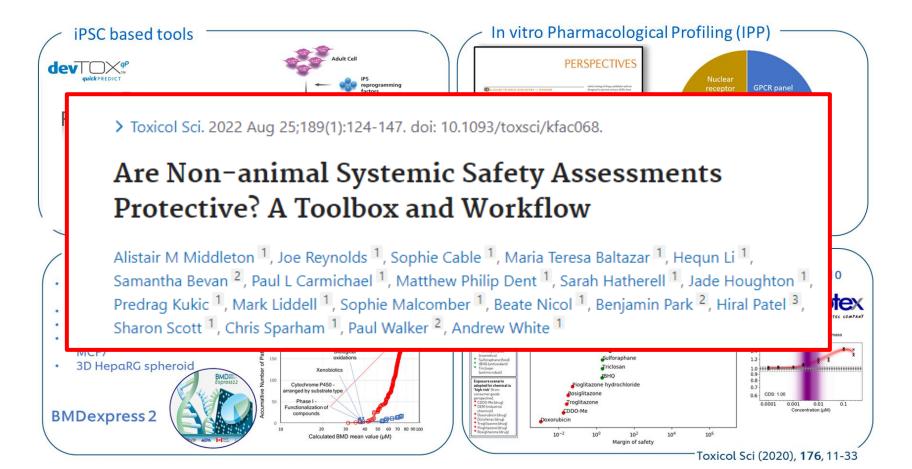








- > Aim: evaluate protectiveness of the NGRA Framework for DART for a given chemical-exposure scenario
- > Each chemical-exposure scenario is classified as "high" or "low" risk for pregnancy
- > For each chemical-exposure scenario we generate NAM data using NGRA Framework

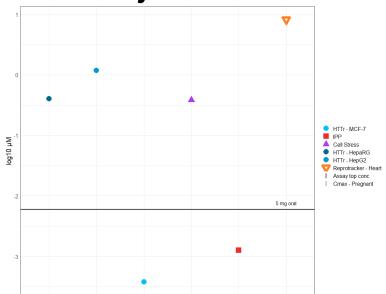






**Exposure Scenario**: Oral 0.5 mg tablet daily during pregnancy = risk for pregnancy





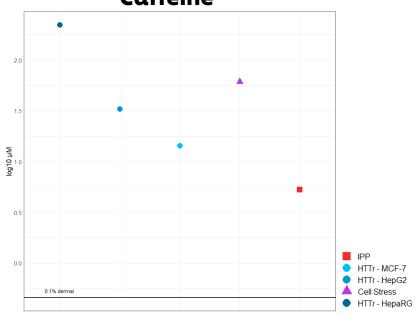
Outcome: Bioactivity detected at or below the plasma Cmax = <u>risk for pregnancy</u>

Unilever

The lowest PoD is coming from HTTR data from MCF7 cells expressing the Estrogen receptor, and from IPP (ER binding)

**Exposure Scenario**: Daily dermal application of 0.1% caffeine in a body lotion = low risk for pregnancy





Outcome: Bioactivity across the DART toolbox occurring at much higher concentrations than the plasma  $C_{max} = low risk for pregnancy$ 

The lowest PoD coming from IPP ADORA2A

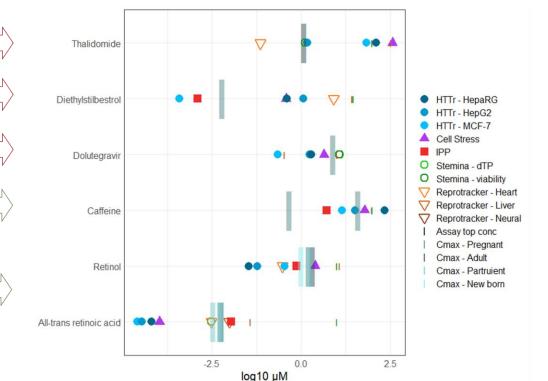
50mg oral application of Thalidomide, high risk, causing dev. toxicity.

5mg oral application of DES, high risk, causing estrogen activity/ED

50mg oral application of Dolutegravir, high risk, causing dev. toxicity

Dermal application of 0.1% caffeine in body lotion (lower Cmax), or oral uptake at recommended TDI of 200mg per days (higher Cmax) of caffeine, both low risk risk.

Uptake of vitamin A/retinol or retinol equivalents in normal diet, low risk.
Cmax concentration of retinol and alltrans retinoic acid (metabolite of retinol) were measured in blood of adult, pregnant and parturient woman as well as in newborns<sup>3)</sup>.



Lowest PoD for Thalidomide is below Cmax value, the toolbox has correctly identified Thalidomide as high risk with lowest PoD coming from ReproTracker® assay.

Lowest PoD for DES is below Cmax value, the toolbox has correctly identified DES as high risk, lowest POD coming from MCF7 HTTr and estrogen receptor binding (IPP).

Lowest PoD for Dolutegravir is below Cmax value of exposure scenario, the toolbox has correctly identified it as high risk. Refinement for hazard classification as dev. Toxicant would be needed, if requested, as there are indications on dev. tox. but above Cmax values. Cell models like gastroloid systems can detect effects at relevant conc.<sup>4</sup>.

Cmax for dermal application of caffeine is below lowest PoD, the toolbox has correctly identified it as low risk. For oral uptake of caffeine, the lowest PoD is below Cmax values indicating risk. Refinement for risk assessment would be needed.

Lowest PoD for retinol as well as all-trans retinoic acid is below Cmax values indicating high risk. Further tools would be needed to refine between bioactivity versus adversity of the compound.



50mg oral application of Thalidomide, high risk, causing dev. toxicity.

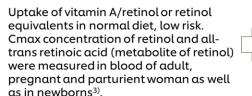
5mg oral application of DES, high risk, causing estrogen activity/ED



50mg oral application of Dolutegravir, high risk, causing dev. toxicity



Dermal application of 0.1% caffeine in body lotion (lower Cmax), or oral uptake at recommended TDI of 200mg per days (higher Cmax) of caffeine, both low risk risk.





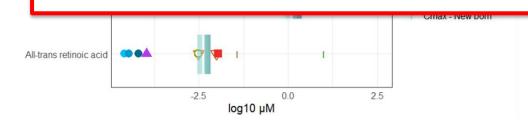
Lowest PoD for Thalidomide is below Cmax value, the toolbox has correctly identified Thalidomide as high risk with lowest PoD coming from ReproTracker® assay.

Lowest PoD for DES is below Cmax value, the toolbox has correctly identified DES as

Preliminary data is encouraging, we are protective for some key known high risk exposure scenarios. Lots more data to analyse (40 compounds total, ~60+ different exposure scenarios) but a promising start!

lutegravir is below Cmax value of exposure scenario, the toolbox ified it as high risk. Refinement for hazard classification as dev. needed, if requested, as there are indications on dev. tox. but ss. Cell models like gastroloid systems can detect effects at

application of caffeine is below lowest PoD, the toolbox has d it as low risk. For oral uptake of caffeine, the lowest PoD is below ating risk. Refinement for risk assessment would be needed.

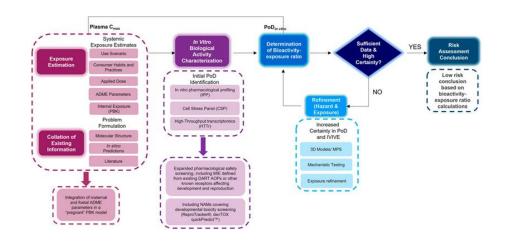


indicating high risk. Further tools would be needed to refine between bioactivity versus adversity of the compound.



## **Next Steps**

- > Evaluation of DART NGRA across many chemistries
- > ReproTracker assay
  - Development and evaluation of an osteoblast differentiation protocol



Rajagopal et al., Front. Toxicol., 2022

- ➤ Identification and filling of existing gaps (placenta transfer measurements, DNT, DIT, endocrine disruptors, multigenerational effects, studying epigenetics in germline development, advanced cell models for refinement)
- > CLP/GHS hazard classification with NAMs



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70+ collaborations



600+ publications





