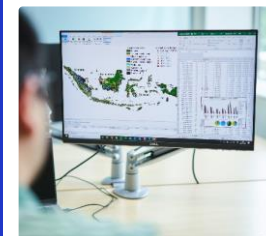
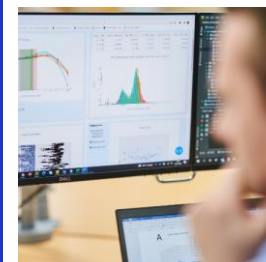


# A Decade of Progress with NAMs & NGRA

**Prof. Dr. Paul L. Carmichael**

**Safety, Environmental & Regulatory Science  
Unilever, UK**

**SERS**  
Safety, Environmental  
& Regulatory Science



# A Decade of TATT

2014线粒体毒性与基于毒性通路的安全性评价新策略学术研讨会暨  
中国毒理学会毒理学替代法与转化毒理学专业委员会成立大会  
2014 Workshop on Mitochondrial Toxicity and Pathway-Based  
Chemical Safety Assessment  
An Inaugural Symposium of the Society of Toxicological  
Alternatives and Translational Toxicology, CSOT

**会议日程与论文集**  
**Program and Proceedings**

主办单位：中国毒理学会  
HOST: Chinese Society of Toxicology (CSOT)  
承办单位：军事医学科学院疾病预防控制中心  
联合利华  
ORGANIZERS: The Institute of Disease Control and Prevention,  
Academy of Military Medical Sciences  
Unilever

2014年10月12 - 14日 • 北京  
October 12 - 14, 2014 • Beijing

**2025 (第八届)**  
**毒性测试替代方法与转化毒理学(国际)学术研讨会**  
2025 (the 8<sup>th</sup>) International Conference on  
Toxicity Testing Alternatives and Translational Toxicology

距离大会开始还有 034 天  
July 2-6, 2025 | Chengdu, Sichuan

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# 2011 to 2025 – From TT21C to NAMs/NGRA



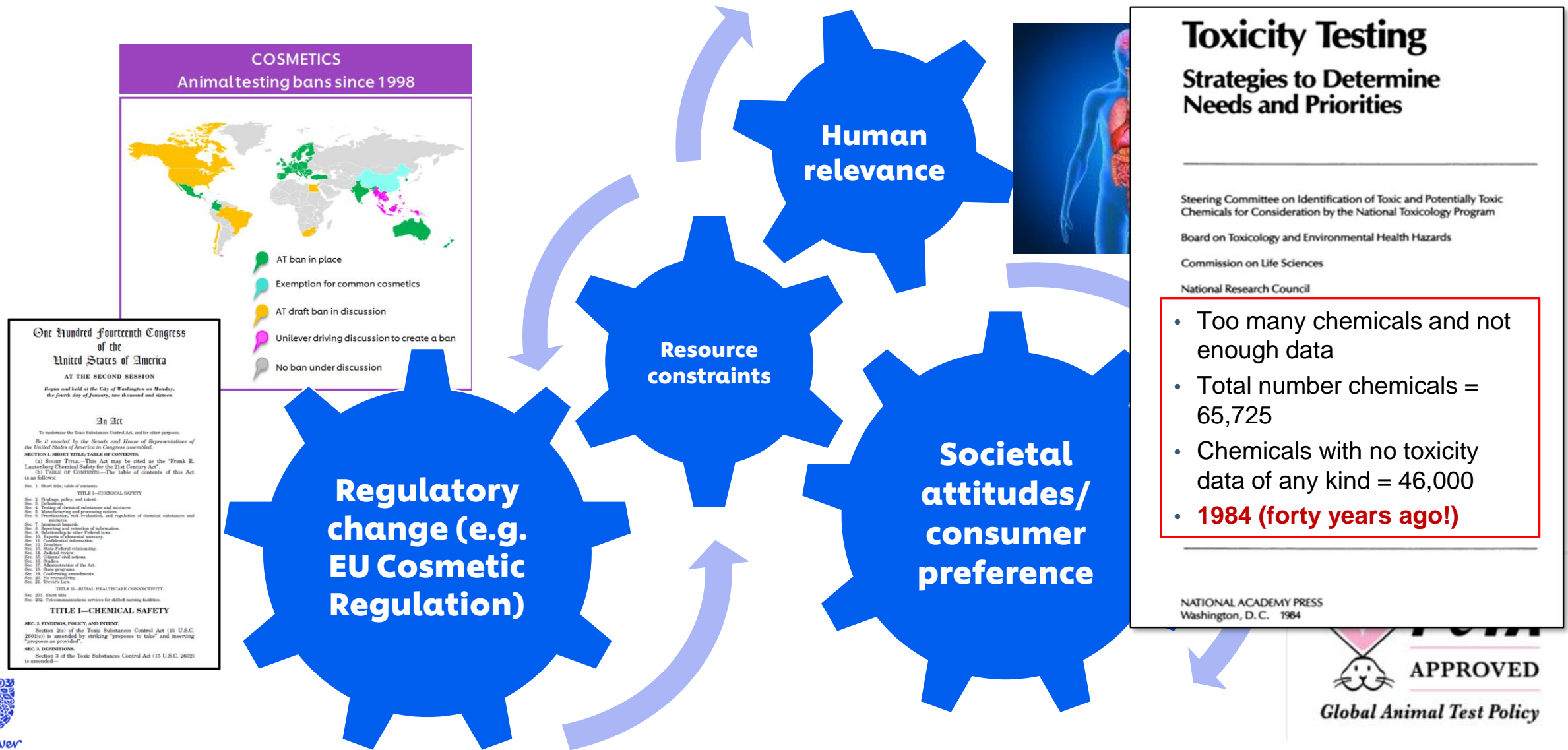
# A Decade of: Next Generation Risk Assessment (NGRA) New Approach (non-animal) Methodologies (NAMs)

**NGRA** is an exposure-led, risk assessment approach that integrates New Approach Methodologies (NAMs) to assure (human & environmental) safety without the use of animal testing

**NAMs** are any *in vitro* or computational (*in silico*) method that enables mechanistically based chemical safety assessment and contributes to the replacement of animals

讨论争议点：下/新一代风险评估；Next/New？

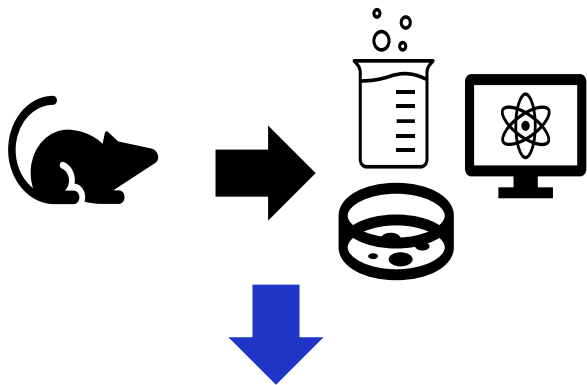
# The Need for Implementation of NAM-Based Safety Assessments





# Non-Animal Frameworks for Safety Decisions:

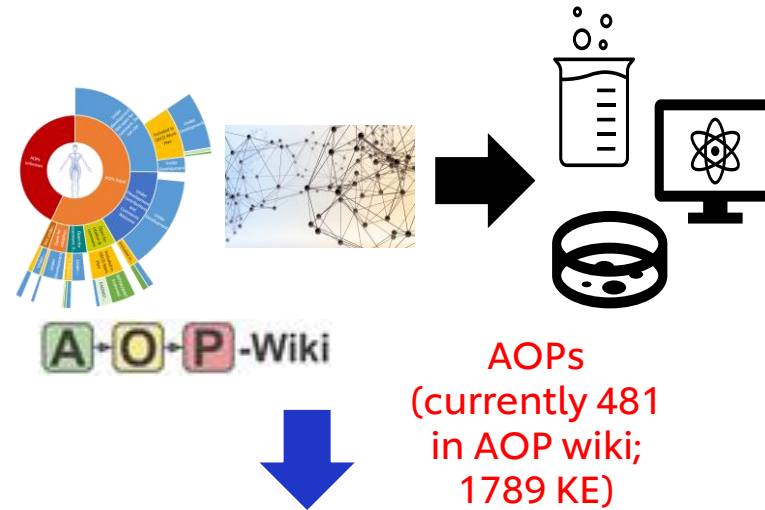
## Non-animal NAMs strategies for 1-2-1 replacement – prediction of animal outcome



Prediction of an animal test is **not** necessarily relevant to assess human safety

Rodent studies have been used in a protective manner with the use of uncertainty factors rather than in a predictive way

## Development of battery of assays aligned to AOPs



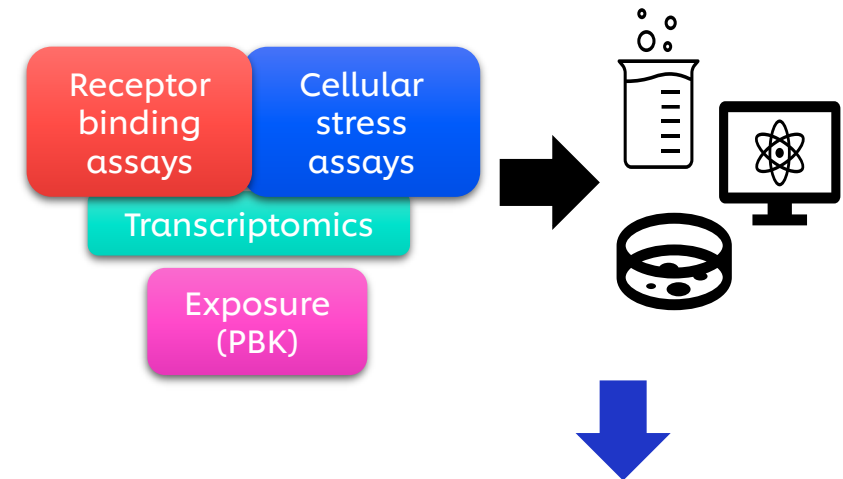
~ Multiple 1000s of assays need to be if multiple AOPs are covered

How to identify the relevant AOP?

**Not** feasible for initial safety testing

Value in bespoke safety assessment when differentiation between bioactivity & adversity is needed (higher tier)

## Development of high-throughput & broad coverage set of non-animal NAMs



### Protection Hypothesis:

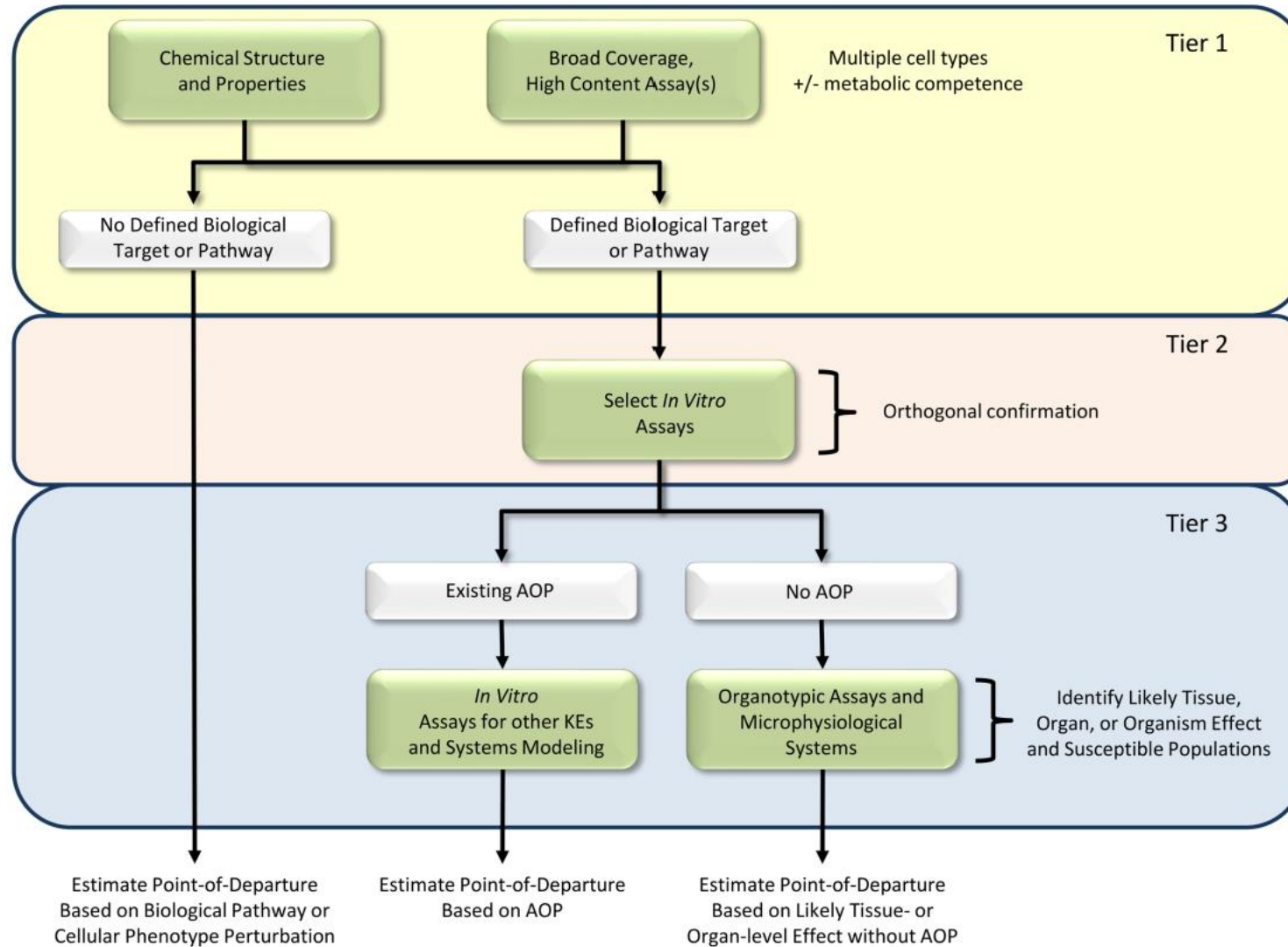
If biological activity measured using a broad suite of human-relevant test systems is above the predicted exposure in humans, then systemic adverse effects are highly unlikely

## Why Tier NAMs in a **Protective NGRA?**

- Can then implement the available and valid tools in a measured and logical way
- Maintain a core dynamic system relevant to the risk assessment/safety question
  - Be ever prepared to improve and update
- Can lead from early-tier rapid computational tools to potentially more complex testing systems in later tiers
  - But only as necessary
- Helps give greater confidence to Regulators?
  - Improving
- Early Tiers come at a 'utility' cost – the loss of otherwise useful chemicals?
  - That's what we have encountered but we must build confidence



# The EPA Blueprint: A 'Tiered' Approach



SOT | Society of Toxicology  
www.toxsci.oxfordjournals.org

TOXICOLOGICAL SCIENCES, 169(2), 2019, 317-332  
doi: 10.1093/toxsci/tfz058  
Advance Access Publication Date: March 5, 2019  
Forum

FORUM

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

Russell S. Thomas,<sup>\*,1</sup> Tina Bahadori,<sup>†</sup> Timothy J. Buckley,<sup>‡</sup> John Cowden,<sup>\*</sup> Chad Deisenroth,<sup>\*</sup> Kathie L. Dionisio,<sup>‡</sup> Jeffery B. Frithsen,<sup>§</sup> Christopher M.





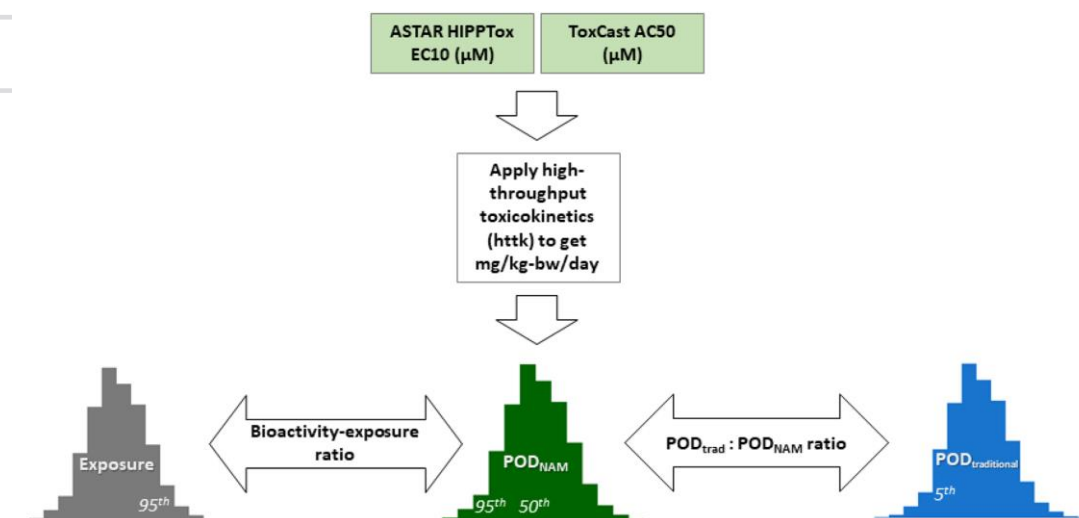
# NAMs are Generally *More* Protective than Animal Tests

## Accelerating the Pace of Chemical Risk Assessment (APCRA)



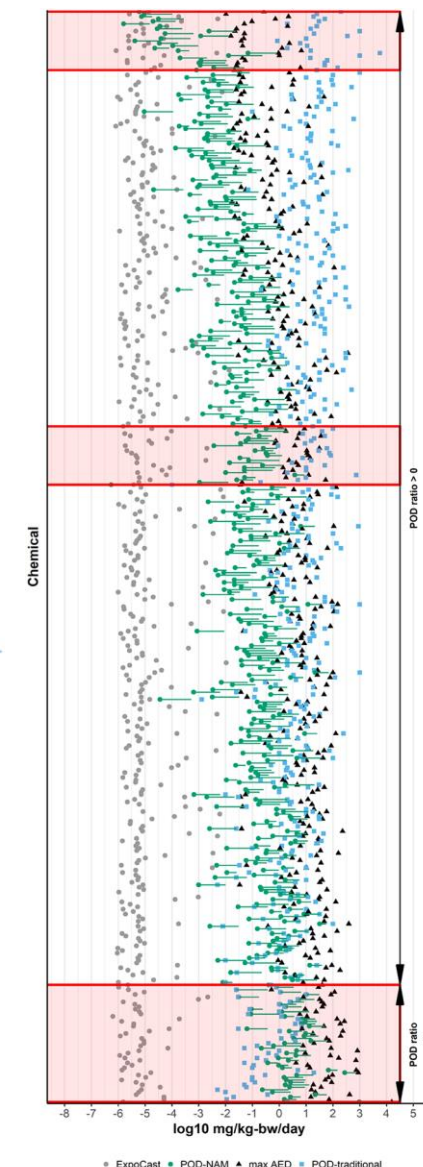
### Utility of In Vitro Bioactivity as a Lower Bound Estimate of In Vivo Adverse Effect Levels and in Risk-Based Prioritization

Katie Paul Friedman <sup>\*,1</sup> Matthew Gagne, <sup>†</sup> Lit-Hsin Loo, <sup>‡</sup> Panagiotis Karamertzanis, <sup>§</sup> Tatiana Netzeva, <sup>§</sup> Tomasz Sobanski, <sup>§</sup> Jill A. Franzosa, <sup>¶</sup> Ann M. Richard, <sup>\*</sup> Ryan R. Lougee, <sup>\*,||</sup> Andrea Gissi, <sup>§</sup> Jia-Ying Joey Lee, <sup>‡</sup> Michelle Angrish, <sup>|||</sup> Jean Lou Dome, <sup>|||</sup> Steven Foster, <sup>#</sup> Kathleen Raffaele, <sup>#</sup> Tina Bahadori, <sup>||</sup> Maureen R. Gwinn, <sup>\*</sup> Jason Lambert, <sup>\*</sup> Maurice Whelan, <sup>\*\*</sup> Mike Rasenberg, <sup>§</sup> Tara Barton-Maclaren, <sup>†</sup> and Russell S. Thomas <sup>\*,\*</sup>

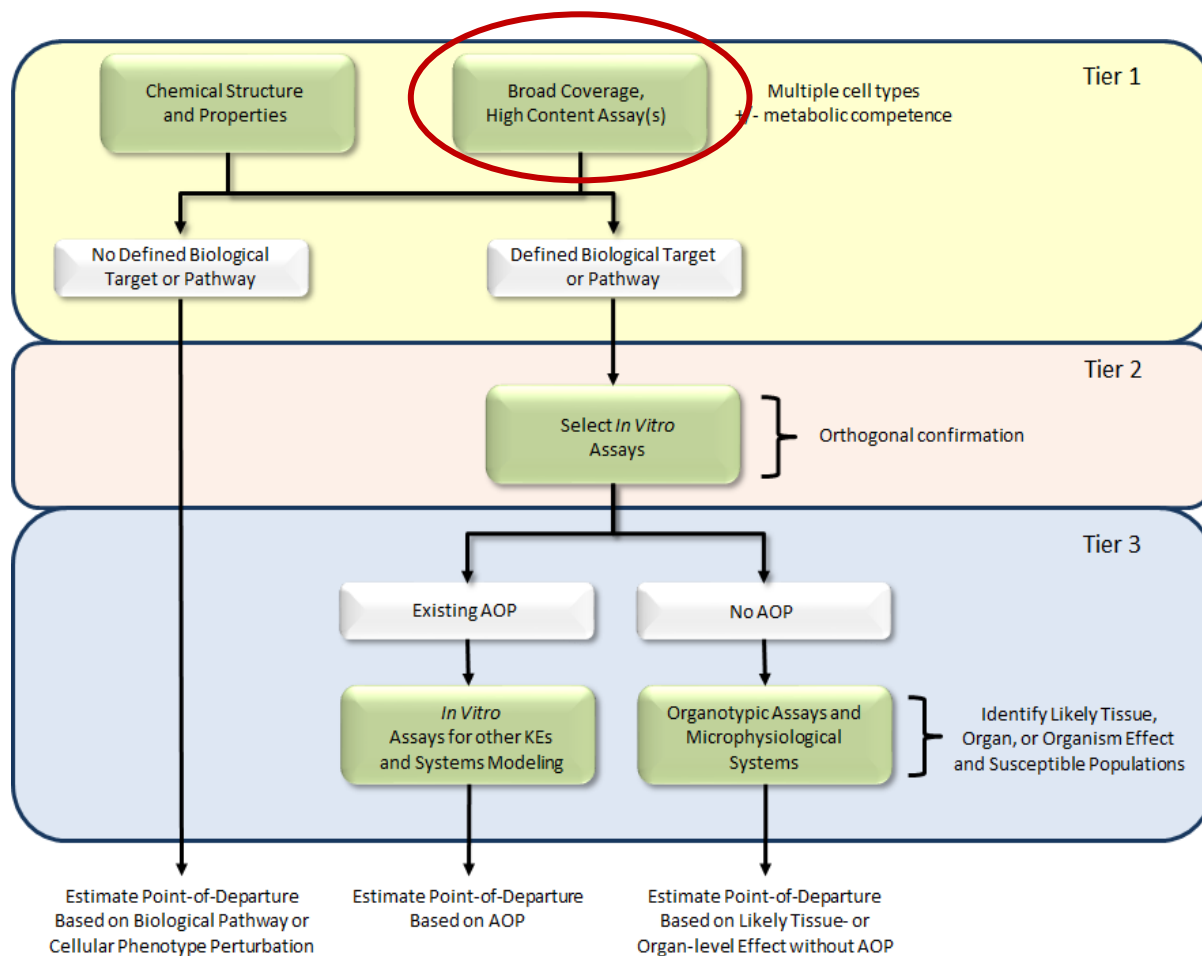


Of the 448 substances, ~**90%** had a POD<sub>NAM,95</sub> that was less than the traditional POD (POD<sub>traditional</sub>) value

**Bioactivity:exposure ratios (BERs)**, useful for identification of priority substances, demonstrated that high-throughput exposure predictions were greater than the POD<sub>NAM,95</sub> for 11 substances



# EPA Transition from ToxCast to Broad Coverage NAM 'Product'



**High throughput profiling (HTP) assays are proposed as the first tier in a NAMs-based hazard evaluation approach**

HTP Assay Criteria:

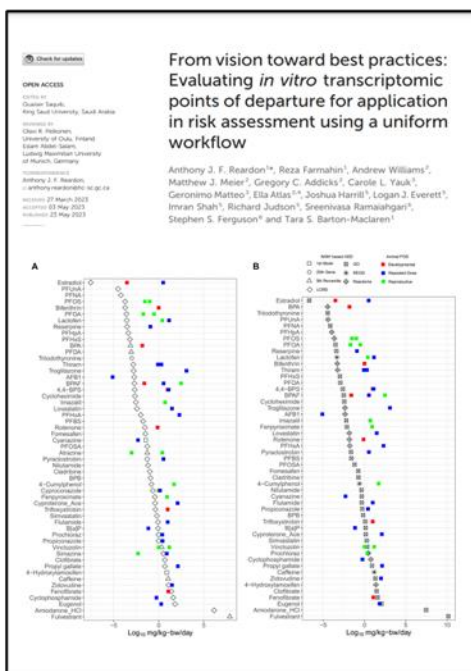
1. Yield bioactivity profiles that can be used for potency estimation, mechanistic prediction and evaluation of chemical similarity
2. Compatible with multiple human-derived culture models
3. Concentration-response screening mode
4. Potential to detect specific and non-specific bioactivity

To date, EPA has identified and implemented two HTP assays that meet this criteria:

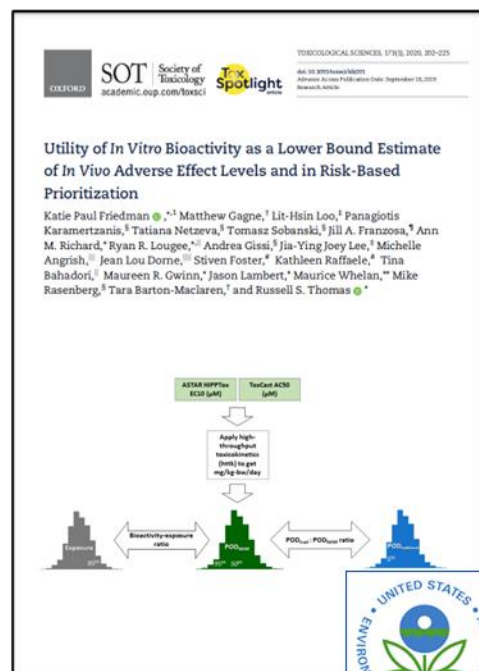
- **High-Throughput Transcriptomics [HTTr]**
- **High-Throughput Phenotypic Profiling [HTPP]**

# Building NAMs/NGRA Confidence: End-to-End Case Studies

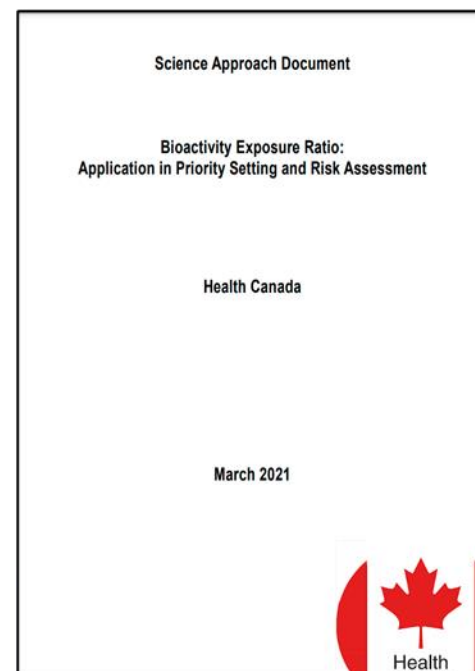
≈40 compounds



448 compounds

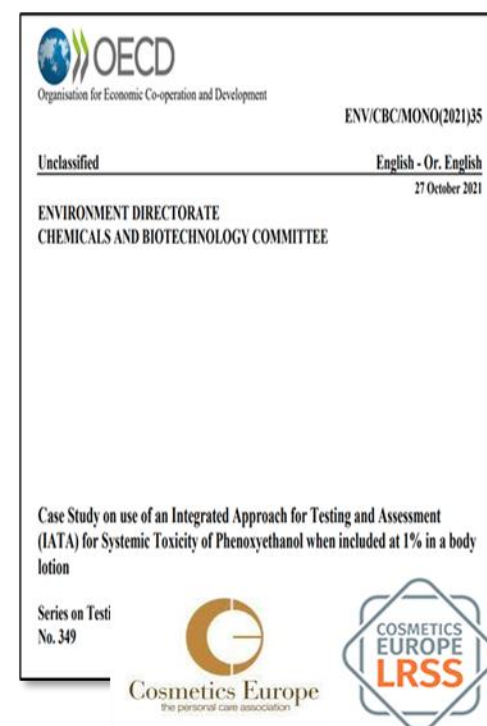


46 compounds

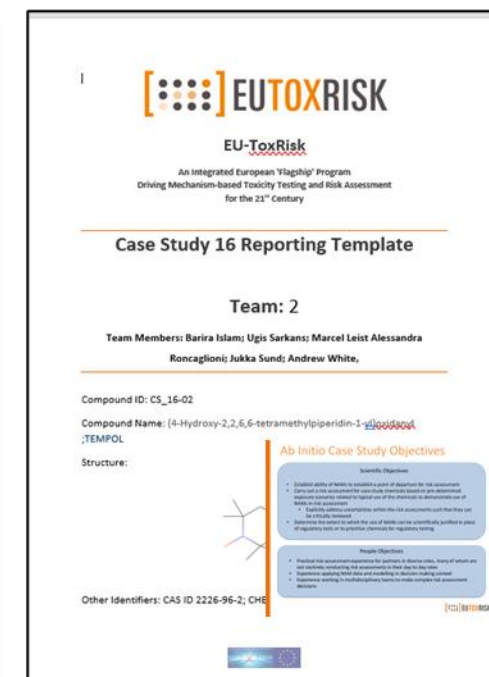


<https://www.canada.ca/en/environment-climate-change/services/evaluating-existing-substances/science-approach-document-bioactivity-exposure-ratio-application-priority-setting-risk-assessment.html>

30 compounds

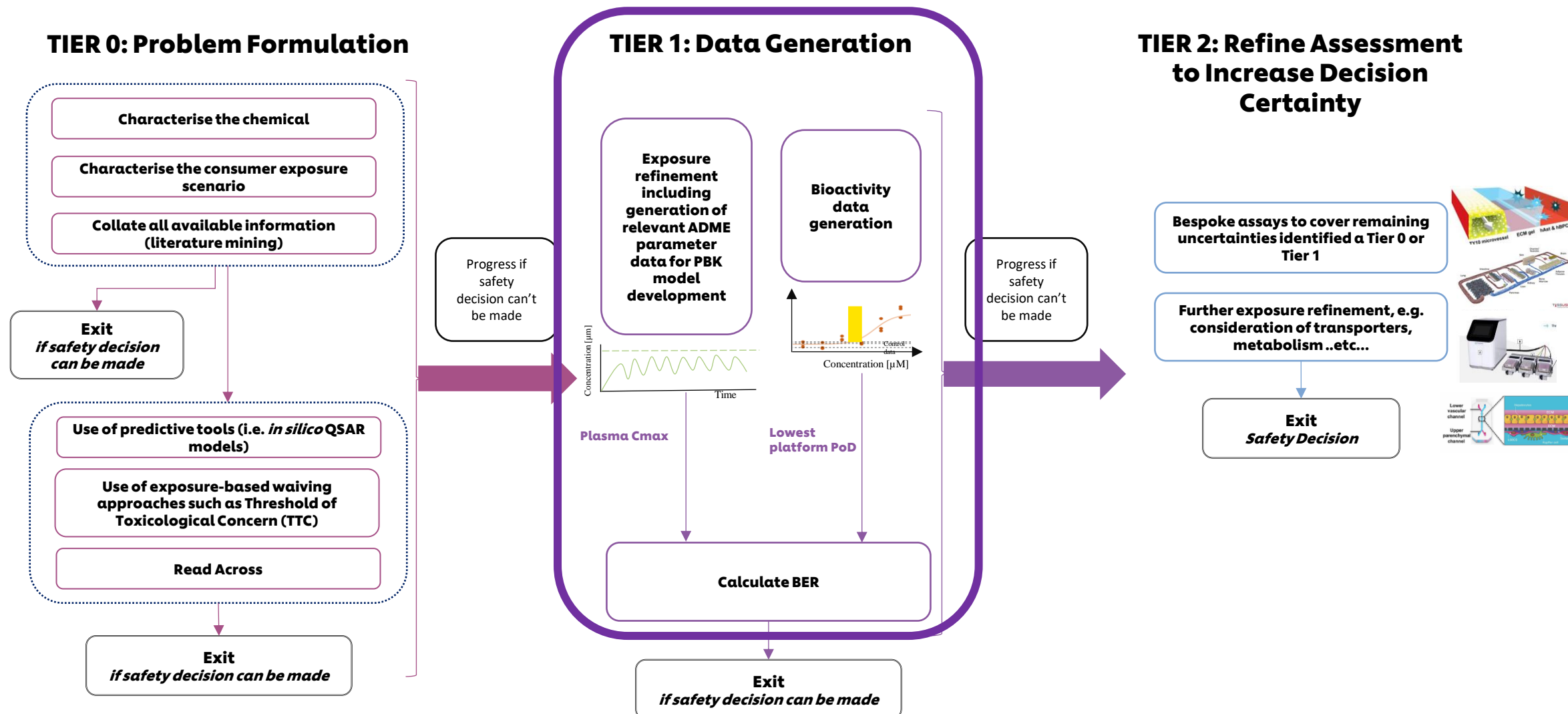


>70 compounds



# Unilever: A NAMs/NGRA Tiered Framework Approach:

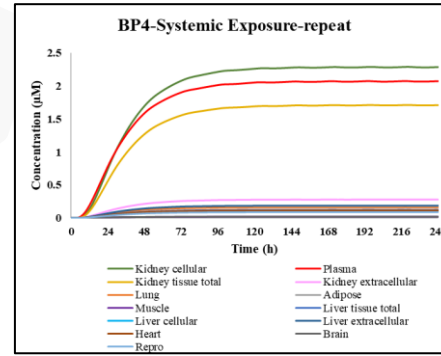
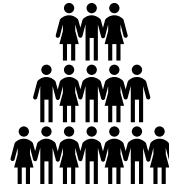
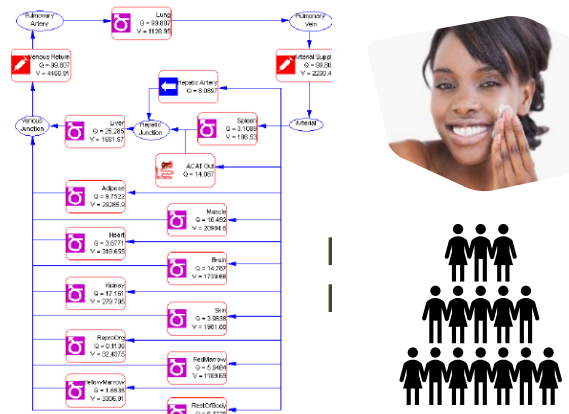
The overall goal is a human safety risk assessment





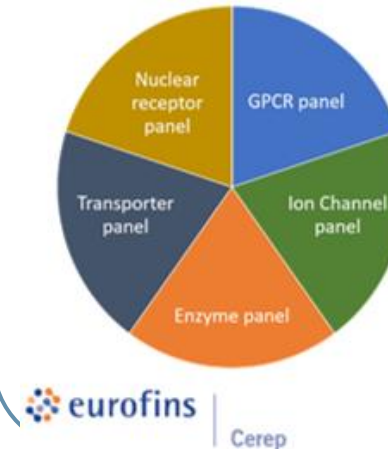
# Unilever: Our Key NAMs

## Internal exposure - PBK modelling



Moxon TE et al., 2020. Toxicology In Vitro, 63, 104746

## In vitro pharmacological profiling



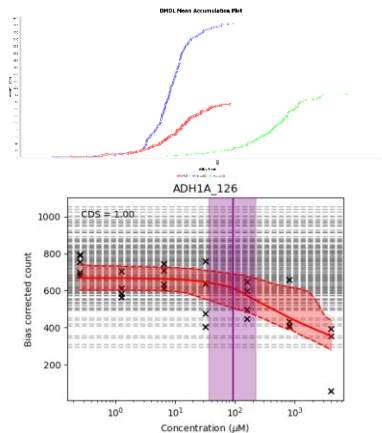
**~79 targets**



Bowes et al. 2012. Nat Rev Drug Discov 11(12): 909-22

## High-Throughput transcriptomics (HTTr)

- TempO-seq technology – full gene panel
- 24hr exposure
- 7 concentrations
- Various cell models (e.g. HepG2, MCF7, HepaRG)
- Dose-response analysis using BMDExpress2 and BIFROST model



Reynolds et al. 2020. Comp Tox 16: 100138  
Baltazar et al. 2020. Toxicol Sci 176(1): 236-252

## Cell stress panel (CSP)

- 36 biomarkers covering 10 cell stress pathways
- HepG2
- 24hr exposure
- 8 concentrations
- Dose-response analysis using BIFROST model

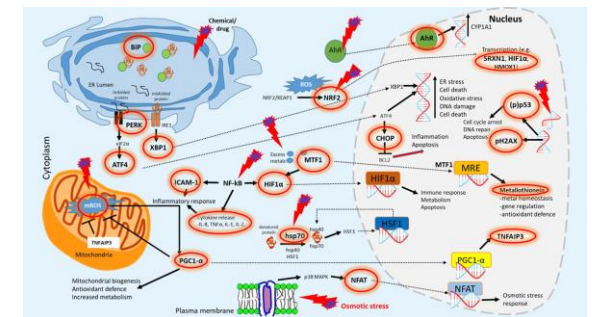


Image kindly provided by Paul Walker (Cyprotex)

Hatherell et al. 2020. Toxicol Sci 176(1): 11-33

# Unilever: Testing the Performance of NAMs in an NGRA

## Benchmarking using chemical-exposure scenarios

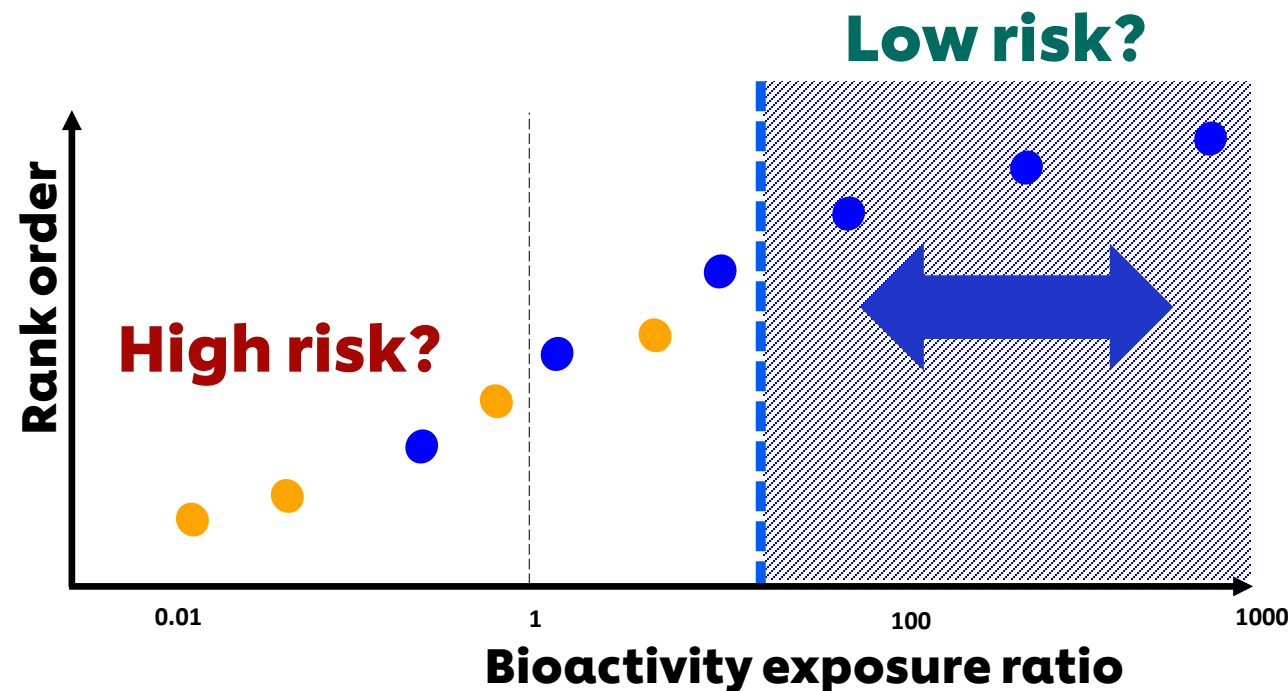
- Chemicals with well-defined human exposures
- Traditional safety assessment available
- High certainty in the risk classification for each chemical-exposure scenario from a consumer goods perspective
- Risk class is relative to consumer health (N.B. drugs = high-risk)



'Low' risk for consumers from systemic perspective



'High' risk for consumers from systemic perspective



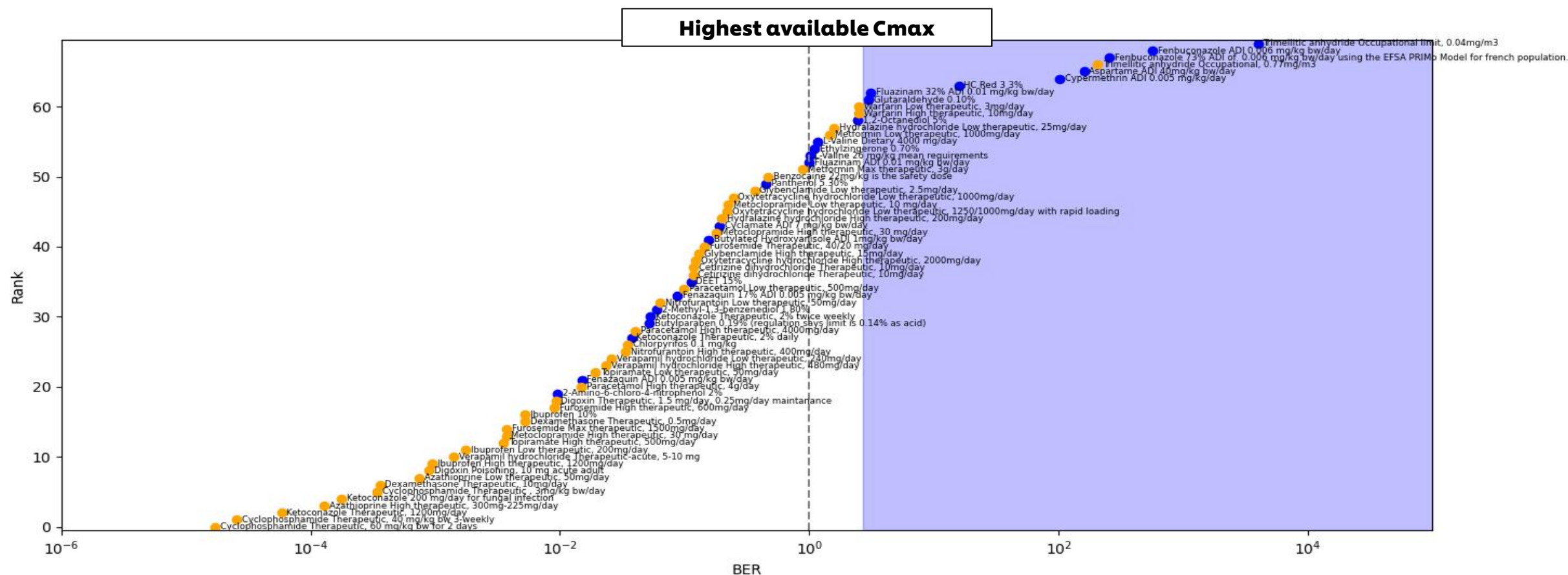
### Protectiveness

How many of the high-risk exposure scenarios are identified as uncertain/high risk?  
(i.e.  $BER < \text{threshold}$ )

### Utility

How many of the low-risk scenarios are identified as low-risk at this early tier stage in a risk assessment framework?  
(i.e.  $BER > \text{threshold}$ )

# Results for 38 Test Chemicals and 70 Exposure Scenarios



**Protectiveness**

**98% (45 out of 46)**

**Utility**

**33% (8 out of 24)**

High-risk exposure scenarios are identified as uncertain/high risk (i.e. BER < threshold)

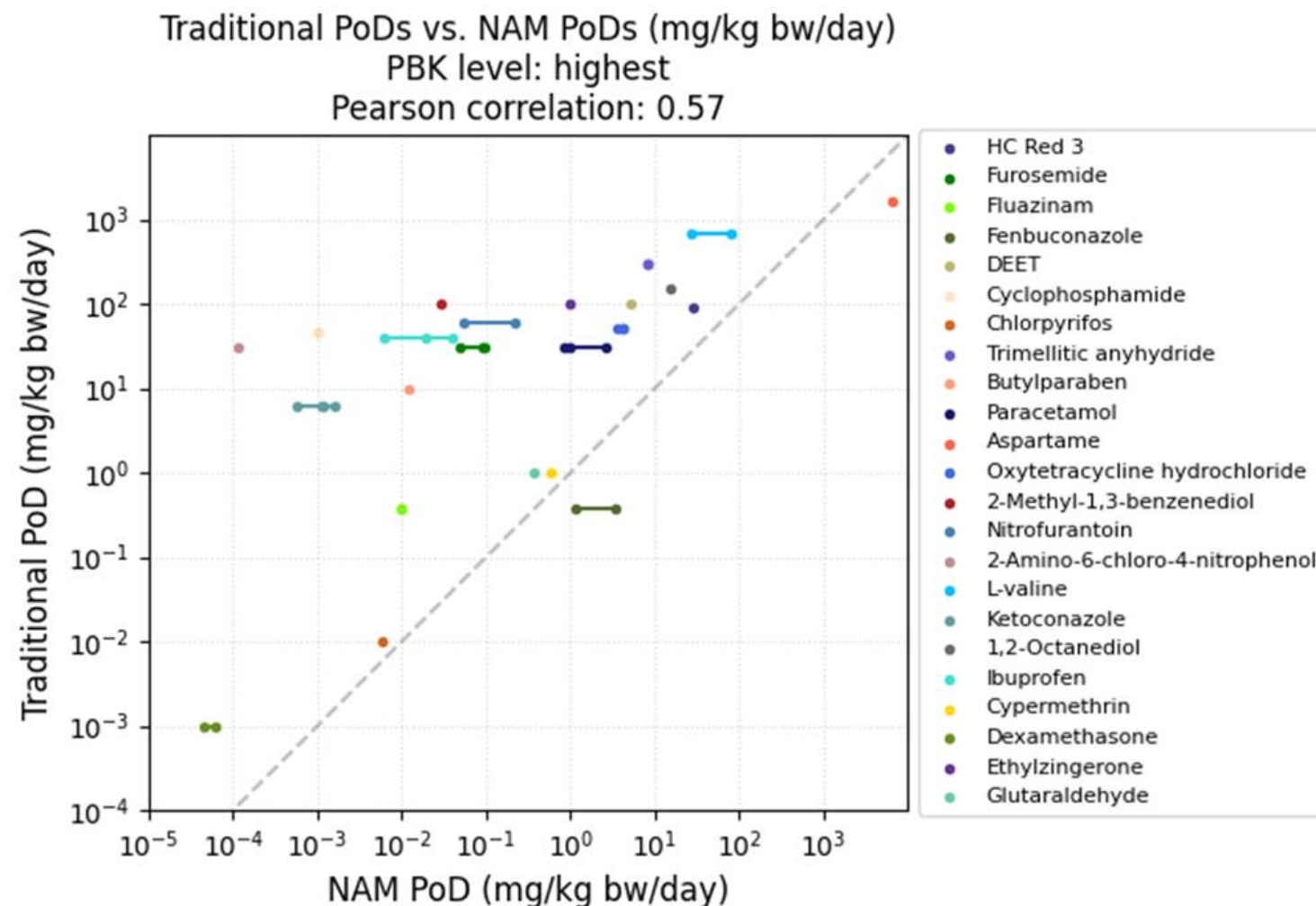
Low-risk scenarios are identified as low risk at this early tier stage in a risk assessment framework (i.e. BER > threshold)

# Comparison of NAM-based Tier 1 Toolbox with Decisions Made Using *in vivo* Data – Protective not Predictive

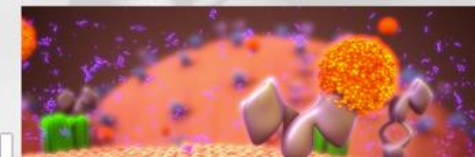
## What if we took the same approach with *in vivo* data?

- Repeat dose *in vivo* data identified for 27 chemicals of the 38 tested.
- In most cases NAM PoDs are more conservative than traditional PoDs

*In agreement with Paul-Friedman et al. (2020)*







**SOT** | Society of  
Toxicology  
[academic.oup.com/toxsci](https://academic.oup.com/toxsci)


Toxicological Sciences, 2025, 204(1), 79–95

<https://doi.org/10.1093/toxsci/kfae159>

Advance Access Publication Date: December 18, 2024

Research article

# Advancing systemic toxicity risk assessment: Evaluation of a NAM-based toolbox approach

Sophie Cable\*, Maria Teresa Baltazar, Fazila Bunglawala, Paul L. Carmichael, Leonardo Contreas, Matthew Philip Dent, Jade Houghton, Predrag Kukic, Sophie Malcomber, Beate Nicol, Katarzyna R. Przybylak, Ans Punt, Georgia Reynolds, Joe Reynolds, Sharon Scott, Dawei Tang, Alistair M. Middleton 

Safety and Environmental Assurance Centre (SEAC), Unilever, Colworth Science Park, Sharnbrook MK44 1LQ, United Kingdom

physiologically based biokinetics



# A Decade of Progress for TATT with NAMs/NGRA/TT21C





# And Yet, Today, Nearly All Regulatory Safety Testing is **Still** Based on:



Despite the current toxicity testing paradigm not sufficiently serving our needs... woefully limited data, high testing costs, extremely long timelines, lack of human relevance, etc.

# What does the USA FDA think?



Home / Regulatory Information / Search for FDA Guidance Documents  
/ Considerations for the Use of Artificial Intelligence To Support Regulatory Decision-Making for Drug and Biological Products

## GUIDANCE DOCUMENT

### Considerations for the Use of Artificial Intelligence To Support Regulatory Decision-Making for Drug and Biological Products

Draft Guidance for Industry and Other Interested Parties

JANUARY 2025

THE AMERICAN JOURNAL  
of MEDICINE  
Official Journal of the Alliance for Academic Internal Medicine

#### The FDA Modernization Act 2.0: Drug Testing in Animals is Rendered Optional

Eli Y. Adashi, MD, MS<sup>a</sup> · Daniel P. O'Mahony, MSLS<sup>b</sup> · I. Glenn Cohen, JD<sup>c</sup>

Affiliations & Notes Article Info

The FDA Modernization Act 2.0 (S.5002), a bipartisan bill co-authored by Senators Cory Booker (D-NJ) and Rand Paul (R-KY), was passed by the Senate on September 29, 2022, by unanimous consent and without amendments. Later ratified by the US House of Representatives, the FDA Modernization Act 2.0, now part of the Consolidated Appropriations Act, 2023, was enacted into law on December 29, 2022.<sup>1</sup> The new law amends the Federal Food, Drug, and Cosmetic Act by authorizing sponsors of novel drugs to make use of "certain alternatives to animal testing, including cell-based assays and computer models, to obtain an exemption from the Food and Drug Administration to investigate the safety and efficacy of a drug."<sup>2</sup> The new law also "removes a requirement to use animal studies as part of the process to obtain a license for a biological product that is biosimilar or interchangeable with another biological product."<sup>3</sup> In this Commentary we review the history of legislative efforts to curtail the mandated use of animals in the testing of drugs for safety and efficacy, discuss the emergence of nonanimal drug testing technology, and review the future prospects thereof.

Congressional expressions of concern over the reliance of the US Food and Drug Administration (FDA) on animals for safety and efficacy testing have been a matter of record since 1998. It was in the course of hearings of the Senate Committee on Appropriations that the *Doris Day Animal League* lamented the discontinuation of "all assessments of in vitro, or non-animal test methods, to substantiate the safety of products."<sup>2</sup> Congressional consideration of the authorization of an *Interagency Coordinating Committee on the Validation of Alternative Methods* followed before too long. However, it was not until 2019 that a flurry of Congressional bills were introduced with an eye toward limiting the use of animals in the testing of new drugs on humane grounds. Examples include, but are not limited to, the *Reducing Animal Testing Act* (S.4288), a bill introduced by Senator Ben Ray Lujan (D-NM) on May 19, 2022. Perhaps the most compelling illustration of the leanings of the Senate is illustrated by the "Statement of the Senate on Animal Testing" the central message of which was that "animal testing should not be used for the purposes of safety testing or



## Roadmap to Reducing Animal Testing in Preclinical Safety Studies

### Executive Summary

This roadmap outlines a strategic, stepwise approach for FDA to reduce animal testing in preclinical safety studies with scientifically validated new approach methodologies (NAMs), such as organ-on-a-chip systems, computational modeling, and advanced *in vitro* assays. By partnering with federal agencies like NIH and VA through ICCVAM, FDA can accelerate the validation and adoption of these human-relevant methods, improving predictive accuracy while reducing animal use. This transition will enhance public health by streamlining drug development and ensuring safer therapies reach patients faster, while positioning FDA as a national leader in modern regulatory science and innovation.

### FDA NEWS RELEASE

## FDA Announces Plan to Phase Out Animal Testing Requirement for Monoclonal Antibodies and Other Drugs

Human health and receive FDA approval have been particularlyheimer's (3) andmans, such as aspirin, appeared safe in animal differences between

For Immediate Release: April 10, 2025

Today, the U.S. Food and Drug Administration is taking a groundbreaking step to advance public health by replacing animal testing in the development of monoclonal antibody therapies and other drugs with more effective, human-relevant methods. The new approach is designed to improve drug safety and accelerate the evaluation process, while reducing animal experimentation, lowering research and development (R&D) costs, and ultimately, drug prices.

The FDA's animal testing requirement will be reduced, refined, or potentially replaced using a range of approaches, including AI-based computational models of toxicity and cell lines and organoid toxicity testing in a laboratory setting (so-called New Approach Methodologies or NAMs data).



# Reflected in Funding at USA National Institutes of Health?

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*Turning Discovery Into Health*

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NEWS RELEASES

Tuesday, April 29, 2025

## NIH to prioritize human-based research technologies

*New initiative aims to reduce use of animals in NIH-funded research.*

The National Institutes of Health (NIH) is adopting a new initiative to expand innovative, human-based science while reducing animal use in research. Developing and using cutting-edge alternative nonanimal research models aligns with the U.S. Food and Drug Administration's (FDA) [recent initiative](#) to reduce testing in animals. While traditional animal models



Combining microfabrication techniques with

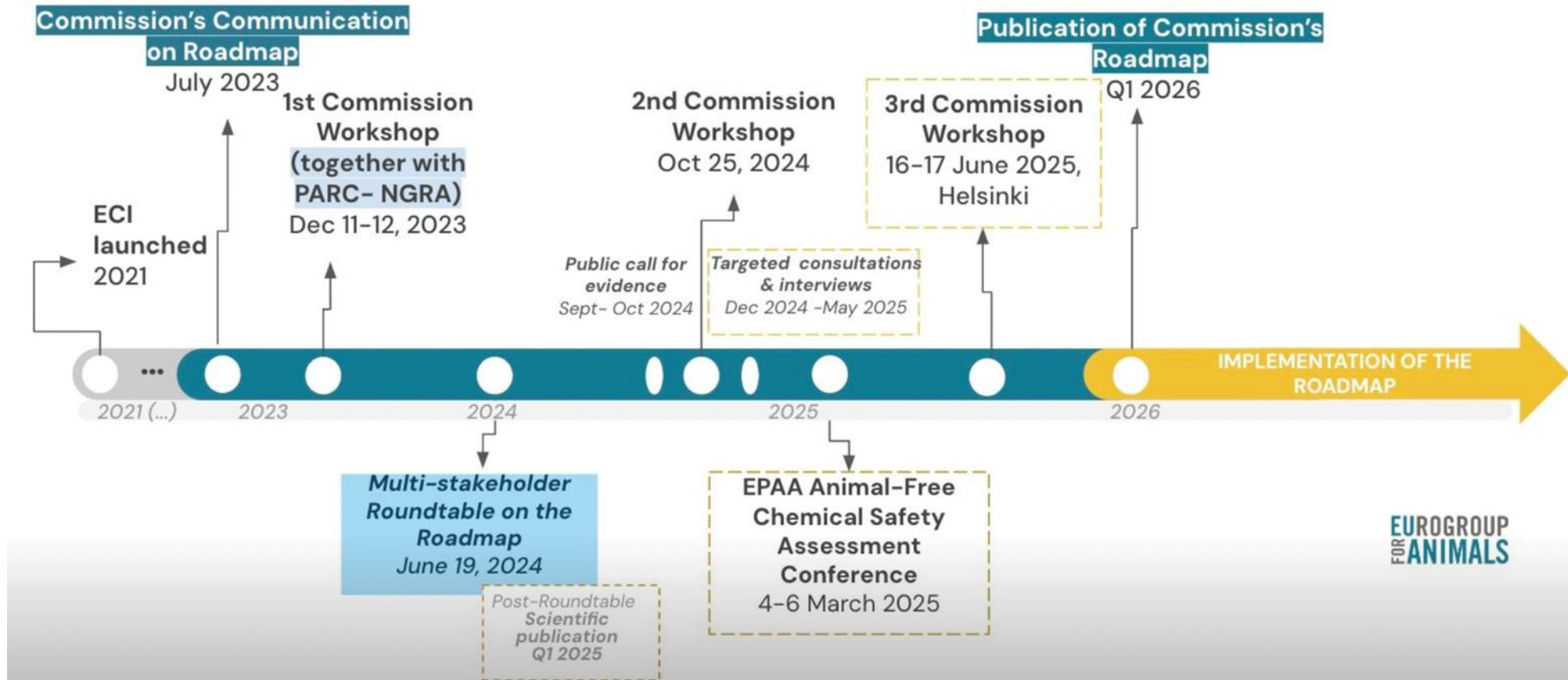
**Institute/Center**  
National Institutes of Health (NIH)

**Contact**  
NIH Office of Communications and Public Liaison   
301-496-5787

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# In Europe, A Roadmap? Talk or Action? Lots to Decide On

## COMMISSION ROADMAP TOWARDS PHASING OUT ANIMAL TESTING FOR CHEMICAL SAFETY ASSESSMENT



# In Our Own Work: Selection of Cells?

- Initially generated HTTr data in **MCF-7**, **HepG2** and **HepaRG** cell lines for all chemicals, and the Cell Stress Panel is conducted in HepG2 cells.
- Now focussed on using **hTERT** cell lines in place of cancer cell lines for new data generation
- Factors that have been considered:
  - Biological diversity. Using content maximisation methods and baseline gene expression data to rank cells
  - Metabolic capacity. Defining HepaRG 3D as the most competent and using the baseline gene expression data to identify the most and least competent cell lines
  - Presence of DART related genes. Making use of the baseline gene expression data and published lists of DART genes
  - Tissue of origin. Consideration of including most likely target organs or barrier tissues

## Already tested:

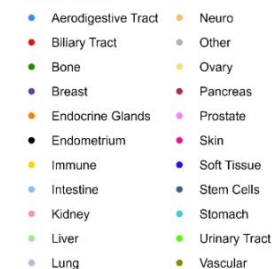
**CHON-001** → Bone  
**HBEC3-KT** → Lung  
**HepG2** → Liver  
**hNP1** → Brain  
**TeloHAEC** → Vascular  
**U-2-OS** → Bone

## In progress:

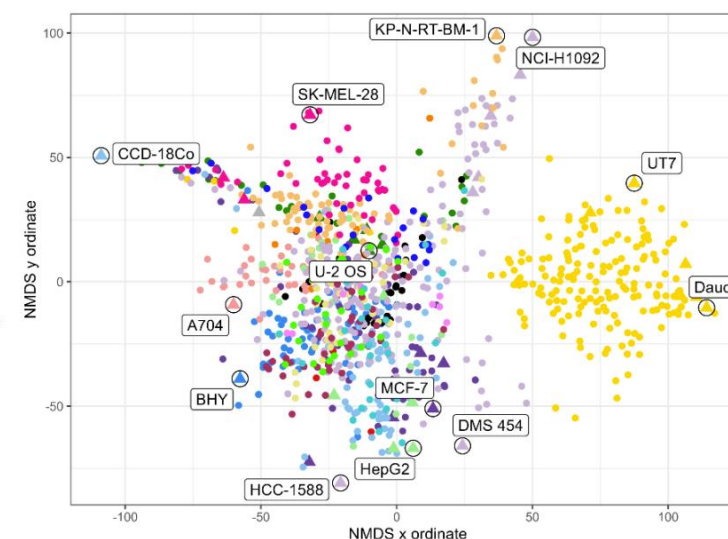
**RPTEC** → Kidney  
**Ker-CT** → Skin  
**CCD-18Co** → Fibroblast  
**ASC52telo** → Mesenchymal stem cell  
**Two zebrafish cell lines**

A

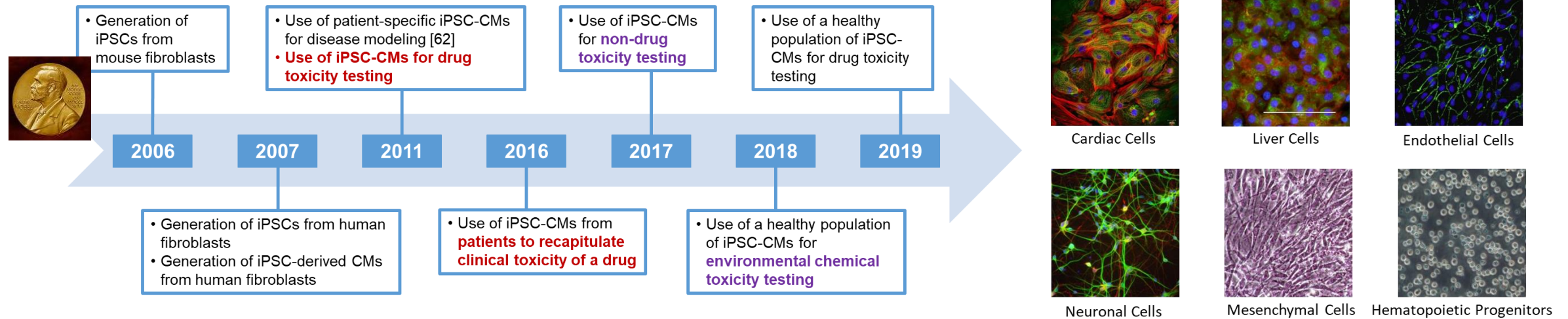
Tissue Origin



TempOSeq



# Are hTERT Cells Likely to be Superseded? YES! By Human iPSCs



Patient/ Healthy Subject

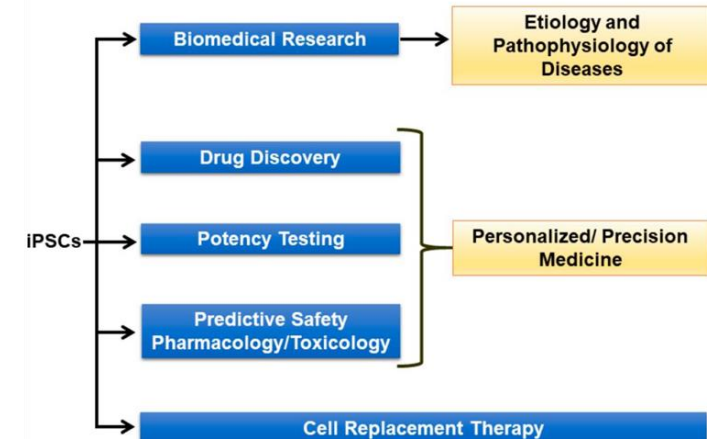


Tissue Sources

Skin Biopsy  
Peripheral Blood  
Hair follicles  
Urine

Cell Types

Fibroblasts  
T Lymphocytes  
Keratinocytes  
Renal Tubular Cells





Grouping of UVCBs to  
waive (or minimize)  
animal testing  
requirements

Research Article  
**Grouping of UVCB Substances with New Approach Methodologies (NAMs) Data**

## Grouping of UVCB Substances with Dose-Response Transcriptomics Data from Human Cell-Based Assays

**A tiered testing strategy based on *in vitro* phenotypic and transcriptomic data for selecting representative petroleum UVCBs for toxicity evaluation *in vivo***

Han-Hsuan Doris Tsai<sup>1,2,3</sup>, John S. House<sup>4</sup>, Fred A. Wright<sup>1,4,5</sup>, Weihsueh A. Chiu<sup>1,2</sup>, Ivan Ruany<sup>1,2,4</sup>

<sup>1</sup>Interdisciplinary Faculty of Toxicology, College Station, Texas 77843, USA  
<sup>2</sup>Department of Veterinary Physiology and Pharmacology, Texas A&M University, College Station, Texas 77843, USA  
<sup>3</sup>National Institute of Environmental Health Sciences, Research Triangle Park, North Carolina 27709, USA  
<sup>4</sup>Department of Statistics and Biostatistics Research Center, North Carolina State University, Raleigh, North Carolina 27695, USA  
<sup>5</sup>Department of Biological Sciences and Biocomputation Research Center, North Carolina State University, Raleigh, North Carolina 27695, USA

To whom correspondence should be addressed. E-mail: [iruan@ncsu.edu](mailto:iruan@ncsu.edu)

1 Department of Veterinary Physiology and Pharmacology, Texas A&M University, College Station, Texas 77843, USA  
2 National Institute of Environmental Health Sciences, Research Triangle Park, North Carolina 27709, USA  
3 Department of Statistics and Biostatistics Research Center, North Carolina State University, Raleigh, North Carolina 27603, USA  
4 Department of Biological Sciences and Biostatistics Research Center, North Carolina State University, Raleigh, North Carolina 27609, USA

\* To whom correspondence should be addressed. E-mail: [inayem@stat.tamu.edu](mailto:inayem@stat.tamu.edu)

**1** **Abstract**

**1** **1** Hazard evaluation of substances of "unknown or variable" composition, complex reaction products and biological materials ("UVChs")

**1** **2** presents a major challenge in regulatory science because their chemical composition is difficult to ascertain. Petroleum substances

**1** **3** are representative UVChs and human cell-based data have been previously used to substantiate their grouping for regulatory

**1** **4** hazard assessment. However, the use of cell-based data for hazard assessment is limited by the lack of mechanistic information

**1** **5** needed to select groups of representative worst-case petroleum UVChs for subsequent toxicity evaluation *in vivo*. We used data obtained

**1** **6** from a comprehensive review of the literature to identify and evaluate the most informative cell types and assays for the

**1** **7** derived hepatocytes, cardiomyocytes, neurons, and endothelial cells, and MCF7 and A549 cell lines. Benchmark doses for genomic

**1** **8** DNA damage, cell death, and cell growth were determined for each cell type and assay. The most informative cell types and

**1** **9** assays were identified and used to select representative worst-case petroleum UVChs for subsequent toxicity evaluation *in vivo*.

**1** **10** Correlation analysis and machine learning were used to assess associations between phenotypic and transcriptional PkGs and to

**1** **11** identify the most informative cell types and assays for representing a cost-effective regulatory testing strategy. We found that 2

**1** **12** cell types—JSC-derived hepatocytes and cardiomyocytes—contributed the most informative and protective PkGs and may be used

**1** **13** for the selection of representative worst-case petroleum UVChs for subsequent toxicity evaluation *in vivo*. The use of these

**1** **14** regulatory methodologies to prioritize UVChs has not been widely adopted. Our study supports a revised testing strategy based on

**1** **15** cell-derived hepatocytes and cardiomyocytes for the selection of representative worst-case petroleum UVChs from complex

**1** **16** manufacturing categories for further toxicity evaluation *in vivo*.

**1** **17** **1** **18** **19** **20** **21** **22** **23** **24** **25** **26** **27** **28** **29** **30** **31** **32** **33** **34** **35** **36** **37** **38** **39** **40** **41** **42** **43** **44** **45** **46** **47** **48** **49** **50** **51** **52** **53** **54** **55** **56** **57** **58** **59** **60** **61** **62** **63** **64** **65** **66** **67** **68** **69** **70** **71** **72** **73** **74** **75** **76** **77** **78** **79** **80** **81** **82** **83** **84** **85** **86** **87** **88** **89** **90** **91** **92** **93** **94** **95** **96** **97** **98** **99** **100** **101** **102** **103** **104** **105** **106** **107** **108** **109** **110** **111** **112** **113** **114** **115** **116** **117** **118** **119** **120** **121** **122** **123** **124** **125** **126** **127** **128** **129** **130** **131** **132** **133** **134** **135** **136** **137** **138** **139** **140** **141** **142** **143** **144** **145** **146** **147** **148** **149** **150** **151** **152** **153** **154** **155** **156** **157** **158** **159** **160** **161** **162** **163** **164** **165** **166** **167** **168** **169** **170** **171** **172** **173** **174** **175** **176** **177** **178** **179** **180** **181** **182** **183** **184** **185** **186** **187** **188** **189** **190** **191** **192** **193** **194** **195** **196** **197** **198** **199** **200** **201** **202** **203** **204** **205** **206** **207** **208** **209** **210** **211** **212** **213** **214** **215** **216** **217** **218** **219** **220** **221** **222** **223** **224** **225** **226** **227** **228** **229** **230** **231** **232** **233** **234** **235** **236** **237** **238** **239** **240** **241** **242** **243** **244** **245** **246** **247** **248** **249** **250** **251** **252** **253** **254** **255** **256** **257** **258** **259** **260** **261** **262** **263** **264** **265** **266** **267** **268** **269** **270** **271** **272** **273** **274** **275** **276** **277** **278** **279** **280** **281** **282** **283** **284** **285** **286** **287** **288** **289** **290** **291** **292** **293** **294** **295** **296** **297** **298** **299** **300** **301** **302** **303** **304** **305** **306** **307** **308** **309** **310** **311** **312** **313** **314** **315** **316** **317** **318** **319** **320** **321** **322** **323** **324** **325** **326** **327** **328** **329** **330** **331** **332** **333** **334** **335** **336** **337** **338** **339** **340** **341** **342** **343** **344** **345** **346** **347** **348** **349** **350** **351** **352** **353** **354** **355** **356** **357** **358** **359** **360** **361** **362** **363** **364** **365** **366** **367** **368** **369** **370** **371** **372** **373** **374** **375** **376** **377** **378** **379** **380** **381** **382** **383** **384** **385** **386** **387** **388** **389** **390** **391** **392** **393** **394** **395** **396** **397** **398** **399** **400** **401** **402** **403** <

**Keywords:** transcriptomic; petroleum; grouping; dose-response; HTTx; NAM

[illegible]

**Rapid Hazard Characterization of Environmental Chemicals Using a Compendium of Human Cell Lines from Different Organs**

### Cumulative Risk Meets Inter-Individual Variability: Probabilistic Concentration Addition of Complex Mixture Exposures in a Population-Based Human In Vitro Model

<sup>1</sup> Interdisciplinary Faculty of Toxicology, College of Veterinary Medicine and Biomedical Sciences, Texas A&M University, College Station, TX 77843, USA

Department of Veterinary  
Texas A&M University  
([cmey@tamu.edu](mailto:cmey@tamu.edu))

**Keywords:** cumulative risk; dose addition; concentration addition; inter-individual variability; toxicodynamics; chemical mixtures; defined mixtures; human health risk assessment; uncertainty factors; new approach methods

ALTEX 37(6), 2020

[illegible]Research A Section 508-conformant HTML version of this article is available at <https://doi.org/10.1289/ehp.2760>

<sup>1</sup>Interdisciplinary Faculty of Toxicology and Department of Veterinary Integrative Biosciences, College of Veterinary Medicine and Biomedical Sciences, Texas A&M University, College Station, Texas, USA

and or expos *Article*

Weihshueh A. Chiu <sup>1,2</sup> and Ivan Rusyn <sup>1,2,\*</sup>

**Abstract:** Human cell-based population-wide in vitro models have been proposed as a strategy to

**Keywords:** population-wide; inter-individual variability; toxicodynamics; chemical mixtures; defined mixtures; human health risk assessment; uncertainty factors; genome-wide association study

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Zunwei Chen \*,†, Dillon Flood ‡§, Vi-Hai Zhou ‡§, Wafar  Chen et al 2021

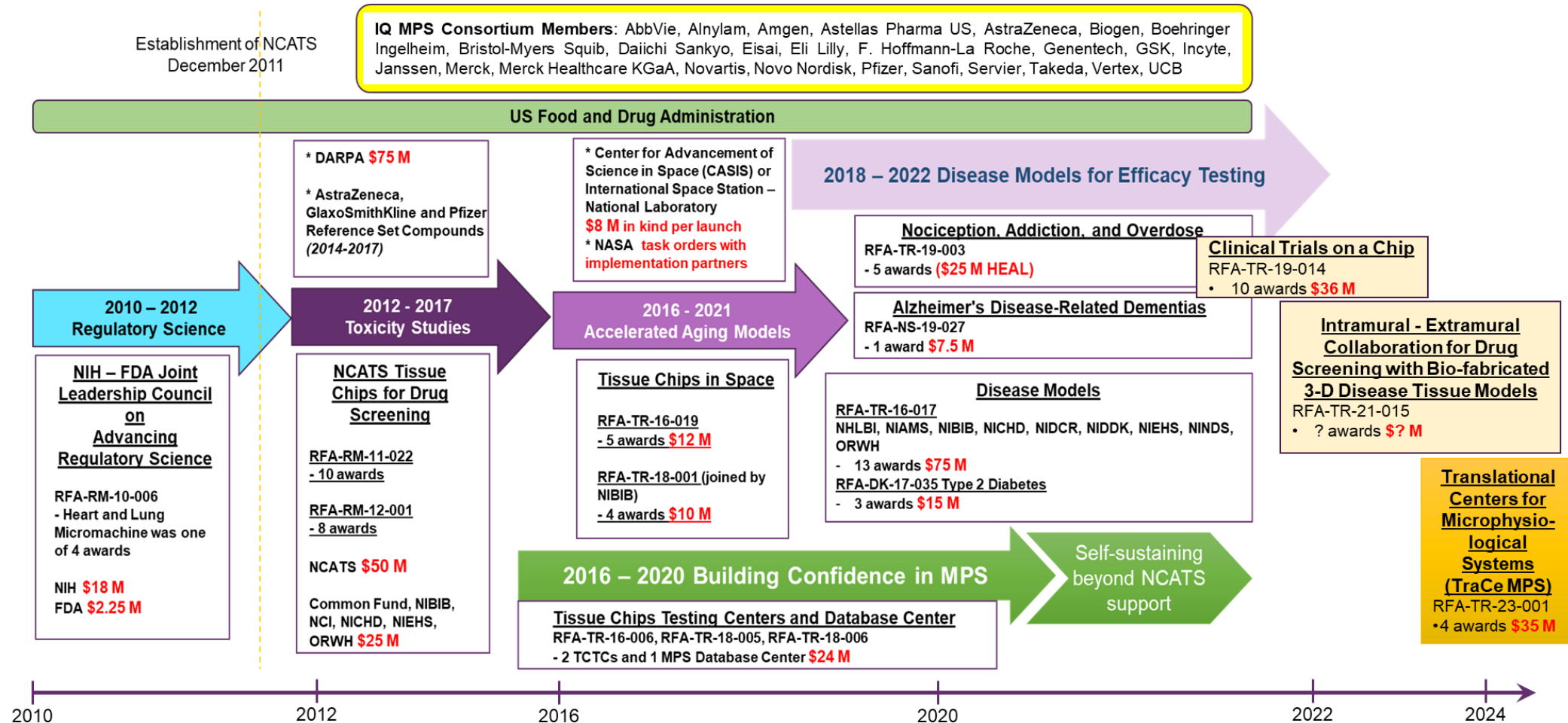
Zunwei Chen<sup>1</sup>, Suji Jang<sup>1</sup>, James M. Kaihatu<sup>2</sup>, Yi-Hui Zhou<sup>3</sup>, Fred A. Wright<sup>3</sup>, Weihsueh A. Chiu<sup>1</sup>  
and Ivan Ruvyn<sup>1,\*</sup>

<sup>1</sup> Civil & Environmental Engineering and Ocean Engineering, Texas A&M University, College Station, TX 77843, USA; [jiazhatao@civl.tamu.edu](mailto:jiazhatao@civl.tamu.edu)

**Keywords:** new approach methods (NAMs); environmental mixtures; disaster research

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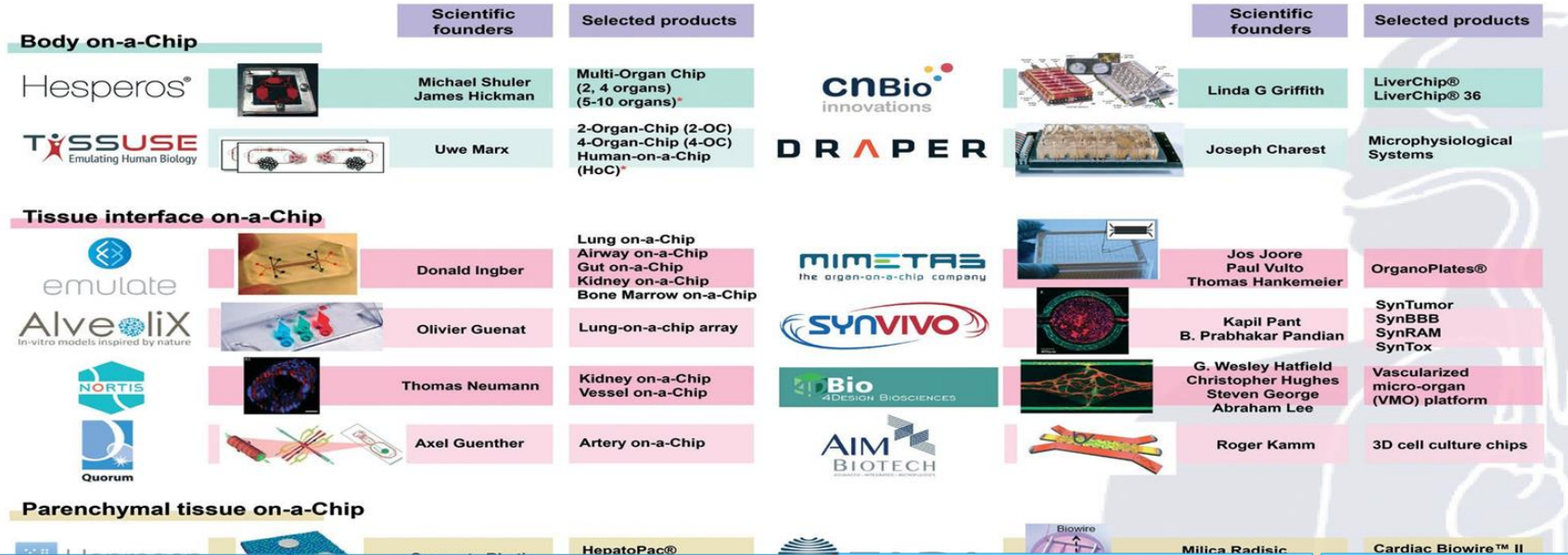
# And That is Without Factoring the MPS/Organ-on-Chip Possibilities!



The USA NIH-funded **'Tissue Chips'** Landscape = **>\$400m** to-date



# Why Not Spin-Off Your Own Organ-on-Chip Company (NIH Supported):



## Tier -1:

### Collaborative research and technology transfer agreements

- Execution of all legal agreements
- Sharing of the protocols
- TAMU staff training with developers

## Tier 0:

### Tissue chip testing without cells

- Assembling of tissue chips
- Testing of the flow and operation
- Testing drug binding to devices
- Development of LC-MS methods

## Tier 1:

### Reproducibility testing of tissue chips

- Replicating published studies
- Evaluation of key findings
- Detailed protocols and SOPs

## Tier 2:

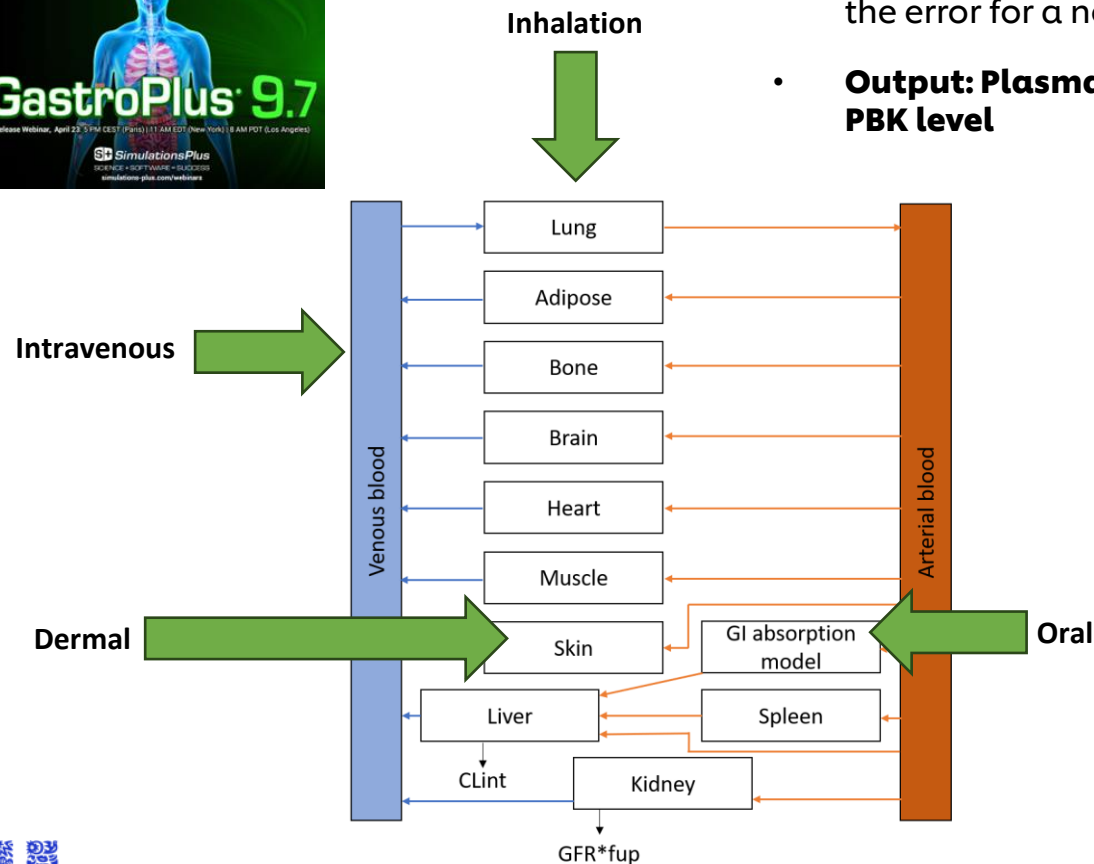
### Extending the utility of the tissue chips

- Defining the “context of use”
- Conducting additional studies
- Depositing data into MPS-Db

4-8 months period of testing for each tissue chip/microphysiological system (MPS)

# PBK Modelling in NGRA – ‘Black Box’ or Fully Understood?

Physiologically based kinetic (PBK) models are used to simulate the behaviour of a chemical in the body for a given exposure scenario



- A Bayesian statistical model to quantifies the error for a novel chemical
- **Output: Plasma C<sub>max</sub> distribution at each PBK level**

- PBK models are composed of multiple coupled ordinary differential equations.
- The model have various parameters that need to be determined.
- Example equation for determining the concentration of chemical in the liver:

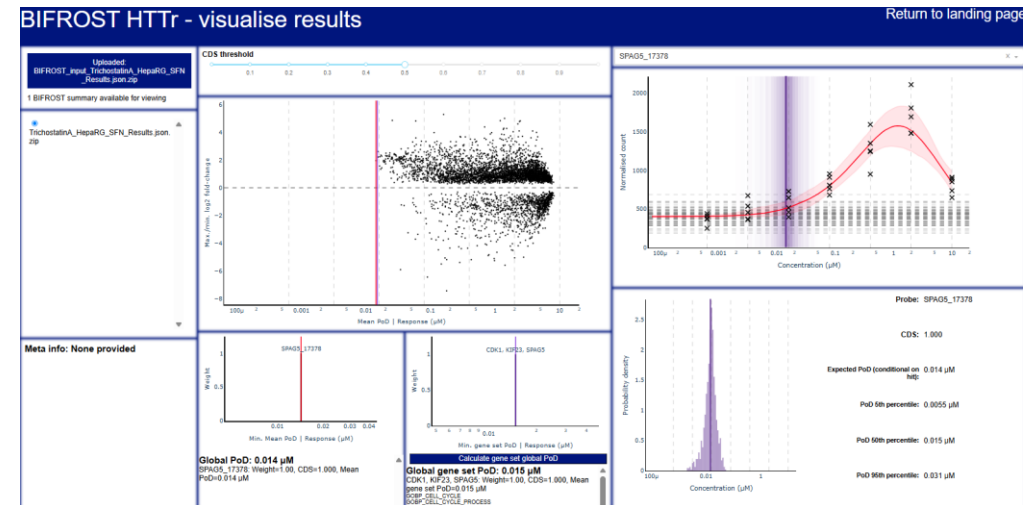
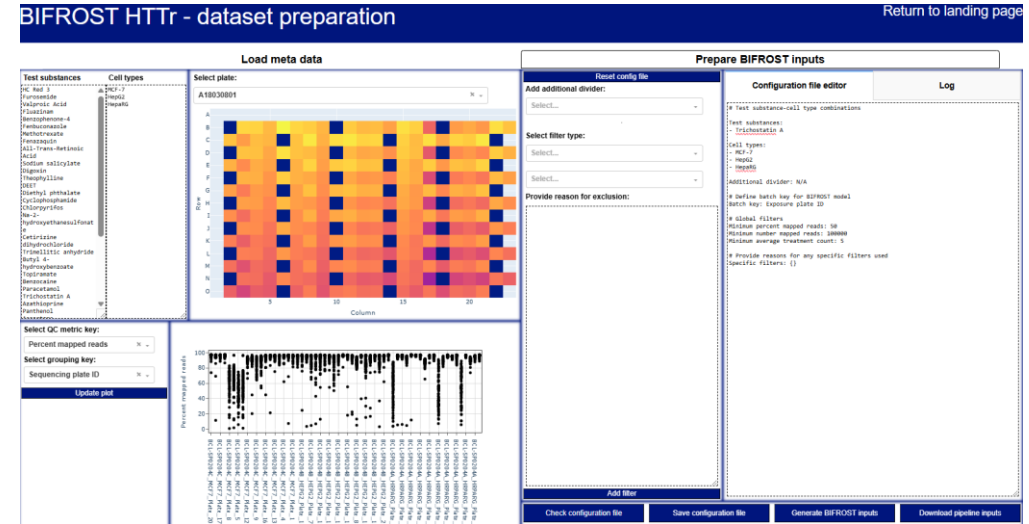
$$V_{\text{Liver}} \frac{dC_{\text{Liver}}}{dt} = Q_{\text{Liver}} \left( C_{\text{Arterial}} - \frac{C_{\text{Liver}}}{P_{\text{Liver}}} \right) - CL_{\text{int}} \left( \frac{C_{\text{Liver}}}{P_{\text{Liver}}} * F_{\text{up}} \right)$$

[Rate of change of concentration in the liver]      [Blood flow rate]      [Concentration of chemical in the arterial blood]      [Concentration of chemical in the liver plasma]      [Clearance rate via metabolism in the liver]

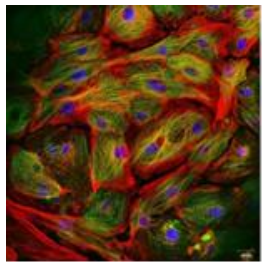


# Defined Approaches for Complex Data Integration: POD in HTTr

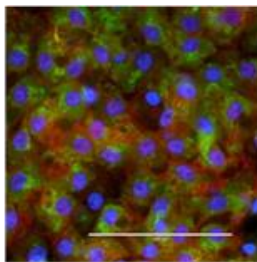
- High throughput transcriptomic (HTTr) studies provide high dimensional datasets for the bioactivity of a chemical
- Downstream results are **highly sensitive** to a plethora of decisions around experimental design, data normalisation, modelling choices, summary statistics and more
- For robust inferences, it is critical that the consequence of each decision, as well as the consequence of having made an alternative choice, is understood



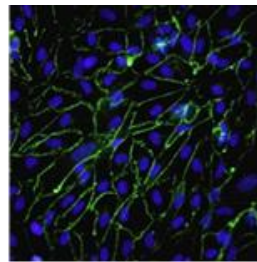
Put Together iPSC of Multiple Lineages, into MPS/OoC, Treat with 'Toxins', Determine POD from HTTr (and other NAMs) by Bayesian Statistics and Compare to Exposure Estimated by Computational PBK with Population Variability... **Oh, and VALIDATE THAT!**



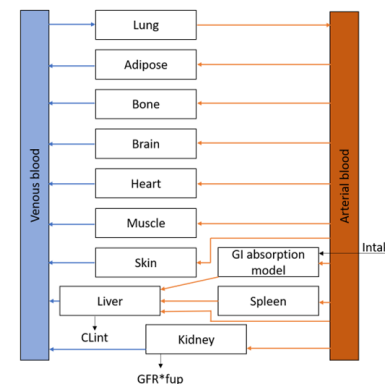
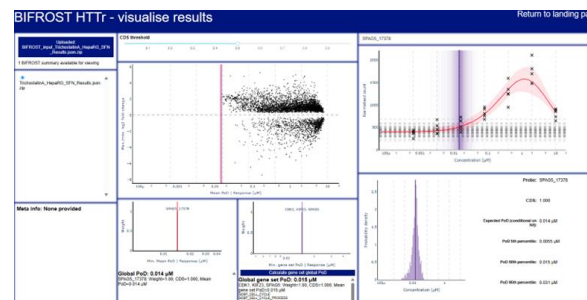
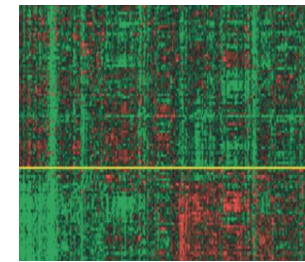
Cardiac Cells



Liver Cells



Endothelial Cells



And let AI make the decision on:  
**Safe or Not-Safe**

# Reproducible? The Regulators Replication Validation Dilemma

nature

News in focus

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NEWS FEATURE | 05 December 2023

Is AI leading to a reproducibility crisis in science?

Scientists worry that ill-informed use of artificial intelligence is driving a deluge of unreliable or useless research.

with repetitive sequences, says Wei.

**Drug design**  
Protein design isn't the only field in which scientists are wielding AIs with words. A slew of models aims to apply a similar approach to designing chemicals.  
Last year, for instance, Gitter's team released a model that designs small molecules in response to text prompts, and showed that it could design drug-like inhibitors of known protein targets. The designs haven't been lab tested, but computational 'docking' tools widely used in drug discovery suggested that some were promising.  
Scientists are also using bio-AIs to 'talk' to cells. Efforts to sequence all the RNA molecules in individual cells have become a bedrock technique in cell biology, revealing unprecedented diversity. But making sense of these data-heavy experiments usually requires intense collaboration between biologists and data scientists, says Christoph Bock, a computational biologist at the Medical University of Vienna.  
As a shortcut, his lab developed an AI chatbot called CellWhisperer. It can take plain English instructions – 'describe these cells in

**AI LINKED TO EXPLOSION IN LOW-QUALITY BIOMEDICAL PAPERS**  
Hundreds of studies seem to follow a template, reporting correlations based on public data sets.

By Miryam Naddaf  
The scientific literature is at risk of becoming flooded with papers that make misleading health claims based on openly available data that are easy to process using artificial intelligence (AI) tools, researchers have warned.  
In a study published in *PLoS Biology* this month, scientists analysed more than 300 papers that used data from the US National Health and Nutrition Examination Survey (NHANES), an open data set of health records (T. Suchak *et al. PLoS Biol.* 23, e3003152; 2025). The papers all seemed to follow a similar template, associating one

conditions having many contributing factors.  
"We have a sudden explosion in publication rates of papers that are extremely formulaic that could easily have been generated by large language models," says study co-author Matt Spick, a biomedical scientist at the University of Surrey in Guildford, UK.  
Spick and his colleagues found that the associations in many of the papers did not hold up to statistical scrutiny, and that some studies seemed to have cherry-picked data.  
"Imagine you're trying to pass an exam that has a particular pass rate, and you add as many questions as you want. You see which ones you got right, and you remove the ones that you got wrong. That's basically what they're doing,"



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Huge reproducibility project fails to validate dozens of biomedical studies

Unique reproducibility effort in Brazil focuses on common methods rather than a single field — and prompts call for reform.

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News Feature | Published: 25 May 2016

1,500 scientists lift the lid on reproducibility

[Monya Baker](#)

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‘Publish or perish’ culture blamed for reproducibility crisis

Survey of more than 1,600 biomedical researchers also flagged small sample sizes and cherry-picking of data as leading causes of reproducibility problems.

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‘Doing good science is hard’: retraction of high-profile reproducibility study prompts soul-searching

A paper by some of the biggest names in scientific integrity is retracted for issues including misstatements about the research plan.



# Are We Even Scratching the Surface of the Wealth of Data in Transcriptomics?



**SOT** | Society of  
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academic.oup.com/toxsci



Toxicological Sciences, 2025, 205(2), 310–325

<https://doi.org/10.1093/toxsci/kfaf036>

Advance Access Publication Date: March 20, 2025

Research article



## A workflow for human health hazard evaluation using transcriptomic data and Key Characteristics-based gene sets

Han-Hsuan D. Tsai <sup>1,2</sup>, King D. Oware <sup>3</sup>, Fred A. Wright<sup>1,4,5</sup>, Weihsueh A. Chiu <sup>1,2</sup>, Ivan Rusyn<sup>1,2,\*</sup>

<sup>1</sup>Interdisciplinary Faculty of Toxicology, Texas A&M University, College Station, TX 77843, United States

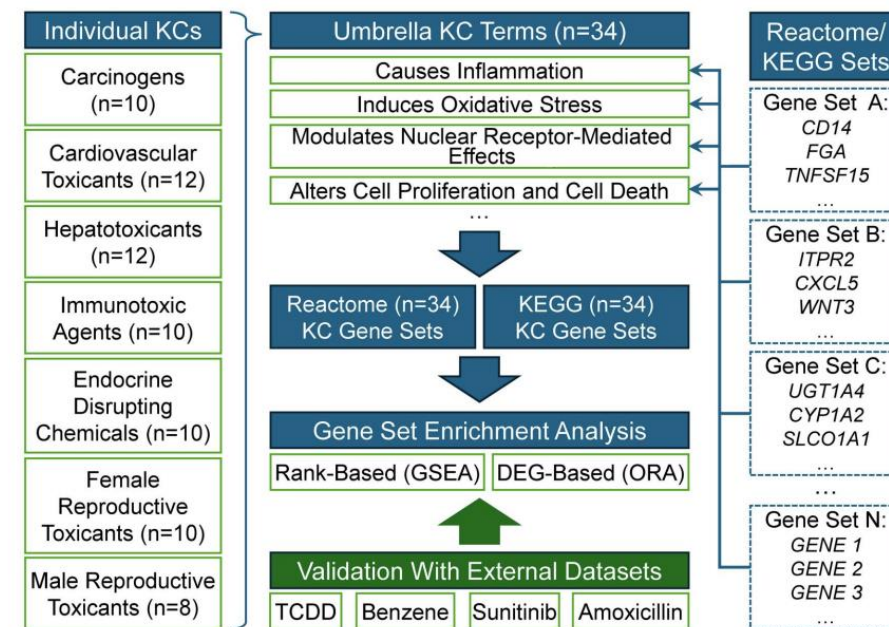
<sup>2</sup>Department of Veterinary Physiology and Pharmacology, Texas A&M University, College Station, TX 77843, United States

<sup>3</sup>School of Public Health, Texas A&M University, College Station, TX 77843, United States

<sup>4</sup>Bioinformatics Research Center, North Carolina State University, Raleigh, NC 27603, United States

<sup>5</sup>Department of Statistics and Bioinformatics Research Center, North Carolina State University, Raleigh, NC 27603, United States

\*Corresponding author: Department of Veterinary Physiology and Pharmacology, TAMU 4466, Texas A&M University, College Station, TX 77843-4466, United States. Email: [irusyn@tamu.edu](mailto:irusyn@tamu.edu)



**Fig. 1.** Overview of the study workflow. Abbreviations: KC, key characteristic; KEGG, Kyoto Encyclopedia of Genes and Genomes; GSEA, gene set enrichment analysis; DEG, differential expressed gene; ORA, over-representation analysis; TCDD, 2,3,7,8-tetrachlorodibenzo-p-dioxin.



# New Collaboration Between Unilever and CMU for 2026+



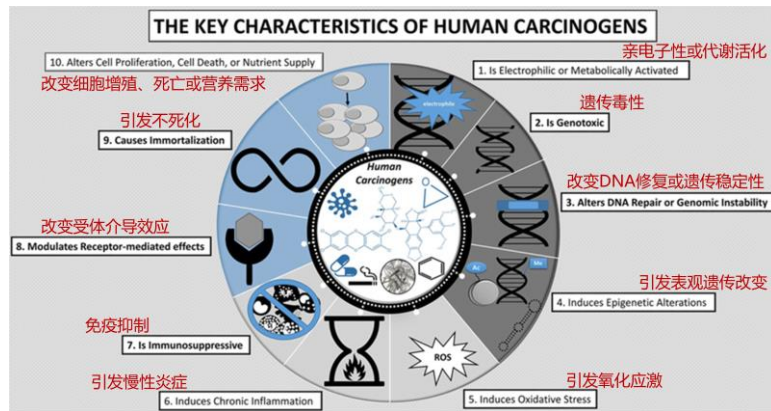
The Significance of Mechanistic Evidence  
in Hazard identification and Risk  
Assessment: from KCs to AOPs

Jingbo Pi



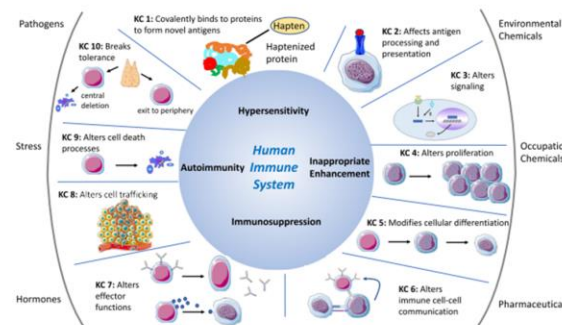
## Outline

- CMU-Unilever Workshop on NAMs-based NGRA
- Understanding from IARC Monographs
- Key characteristics (KCs) vs. AOPs
- A vision for future: KCs-structured NAMs-based NGRA?

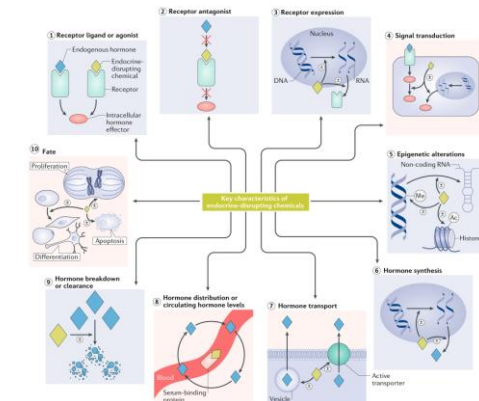


Guyton KZ, et al. Chem Res Toxicol. 2018; 31: 1290–1292.

### KCs associated with immunotoxins



### KCs associated with endocrine disruptors



Nature Reviews Endocrinology (Nat Rev Endocrinol) ISSN 1759-5037 (online) ISSN 1759-5029 (print)

Germolec D.R., et al. Environmental Health Perspectives 130(10) October 2022

# A Decade of Progress for TATT with NAMs/NGRA/TT21C





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