# **€PA**

Targeted high throughput screening for nuclear receptor activation, cell stress, and immunosuppressive bioactivities with 147 perfluoroalkyl substances

**Presentation to FLUOROS** 

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## Outline of this talk

| Concept  | Publication(s)               | DOI  |
|--|------------------------------|--|
| Chemical library of PFAS screened                                    | Patlewicz et al. 2019        | <u>10.1289/EHP4555</u>                             |
| Tiered in vitro screening strategy                                   | Thomas et al. 2019 (general) | <u>10.1093/toxsci/kfz058</u>                       |
|  | EPA's PFAS Action Plan       | https://www.epa.gov/pfas/epas-<br>pfas-action-plan |
| Targeted screening for nuclear receptor activation and cell stress   | Houck et al. (2021)          | <u>10.1016/j.tox.2021.152789</u>                   |
| Targeted screening for<br>immunosuppressive bioactivity<br>signature | Houck et al. (in prep)       | Short preview today of work in progress            |



# **Chemical library of PFAS screened**

## How to select PFAS for tiered screening?

#### Brief Communication

A Section SDE-conformant HTML version of this article is available at https://doi.org/10.1269/EHP4555

U.S. EPA, generating such data to inform agency and partner deci-

sion making regarding potential human health hazard and risk

across the broad landscape of PFASs represents a real-world

challenge that HTT coupled with cheminformatic approaches is

substances were selected for HTT screening and tiered toxicity

testing, along with mention of the toxicity and toxicokinetic

Since there are no specific chemical catalogs for PFASs, an initial

scoping for potentially procurable PFAS substances relied on the

use of candidate PFAS structure lists generated from the U.S. EPA's Distributed Structure-Searchable Toxicity (DSSTox) chemical

database. DSSTox currently exceeds 760,000 substances, each of

which has undergone some level of chemical structure curation prior

to registration (Williams et al. 2017). The largest registered list of

PFAS chemicals available at the time this study was initiated was

the KEMI PFAS list in DSSTox (named PFASKEMI and available

for download at https://comptox.epa.gov/dashboard/chemical\_lists/

pfaskemi). Approximately 1,200 structures from this list were pro-

vided to the chemical contractor for scoping purposes, from which

able but likely to require on-demand synthesis and exceed standard costs. Based on this preliminary scoping, U.S. EPA funds were

secured for the purchase and processing of approximately 400 sub-

programs and regions and to include PFASs with associated tox-

final set of 31 PFASs recommended for further study by this

workgroup (list denoted as EPA PFAS WG 31) identified PFASs

whose review may support risk evaluation. Also included in the

icity data that would inform human health risk assessment. The

The first procurement phase considered the feasibility of pro-

roximately 600 substances were identified as potentially procur-

This article describes, in brief, the development of the PFAS screening library and the process by which a subset of 75 PFAS

uniquely designed to address

experiments currently underway.

**Development of the PFAS Screening Library** 

Discussion

#### A Chemical Category-Based Prioritization Approach for Selecting 75 Per- and Polyfluoroalkyl Substances (PFAS) for Tiered Toxicity and Toxicokinetic Testing

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SUMMARY: Per- and polyfluoroalkyl substances (PFASs) are a group of fluorinated substances of interest to researchers, regulators, and the public date to their widespread presence in the environment. A few PFASs have comparatively extensive amounts of human epidemiological, exposure, and experimental animal toxicity data (e.g., perfluceooctanoic acid), whereas little toxicity and exposure information exists for much of the broader set of PFASs. Given that traditional approaches to generate toxicity information are resource intensive, new approach methods, including in vitro highthroughput toxicity (HTT) testing, are being employed to inform PFAS hazard characterization and further (iv viw) testing. The U.S. Environm Protection Agency (EPA) and the National Toxicology Program (NTP) are collaborating to develop a risk-based approach for conducting PFAS toxic ity testing to facilitate PFAS human health assessments. This article describes the construction of a PFAS screening library and the process by which a targeted subset of 75 PFASs were selected. Multiple factors were considered, including interest to the U.S. EPA, compounds within targeted catego ries, structural diversity, exposure considerations, mocurability and testability, and availability of existing toxicity data. Generating targeted HTT data for PFASs represents a new frontier for informing priority setting. https://doi.org/10.1289/EHP4555

#### Introduction

Per- and polyfluoroalkyl substances (PFASs) are a group of fluorinated substances that have generated increased public attention due to their potential health hazard and widespread presence in the environment (Wang et al. 2017; Xiao 2017; Ross et al. 2018). The U.S. Environmental Protection Agency (EPA) Office of Research and Development (ORD) in partnership with the National Toxicology Program (NTP) are currently engaged in producing toxicity information to facilitate human health assessments for PFASs. A few PFASs have comparatively extensive amounts of toxicity data (e.g., perfluorooctanoic acid), but little toxicity information exists for much of the broader set of PFASs identified from preliminary exposure studies that capture potential occurrence in the environment. The hundreds of untested PFASs provide a scenario in which traditional one-by-one toxicity testing would consume tremendous resources and useful toxicity information would not be available for decades. The U.S. EPA's ToxCast program and the multi-federal agency Tox21 program (which includes the NTP and the U.S. EPA as major partners) have developed the capacity to screen hundreds to thousands of chemicals for bioactivity through in vitro highthroughput toxicity (HTT) testing. Data generated from these assays are already being used to inform hazard identification and prioritize chemicals for further in vivo testing (U.S. EPA 2012, 2014a, 2014b, 2015; Judson et al. 2010, 2015; Kleinstreuer et al. 2017). Within the

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stances to create a PFAS testing library. Received 9 October 2018; Revised 27 November 2018; Accepted 3 curing substances of interest to the U.S. EPA. A U.S. EPA work-December 2018; Published 11 January 2019. group was formed to identify PFASs of interest to U.S. EPA

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**Environmental Health Perspectives** 

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- A few PFAS (e.g., PFOA, PFOS) have extensive information whereas many PFAS have little to no information.
- Select the original 75, extended to select ~150 for screening activities.

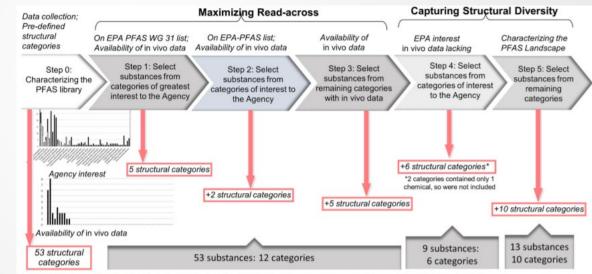
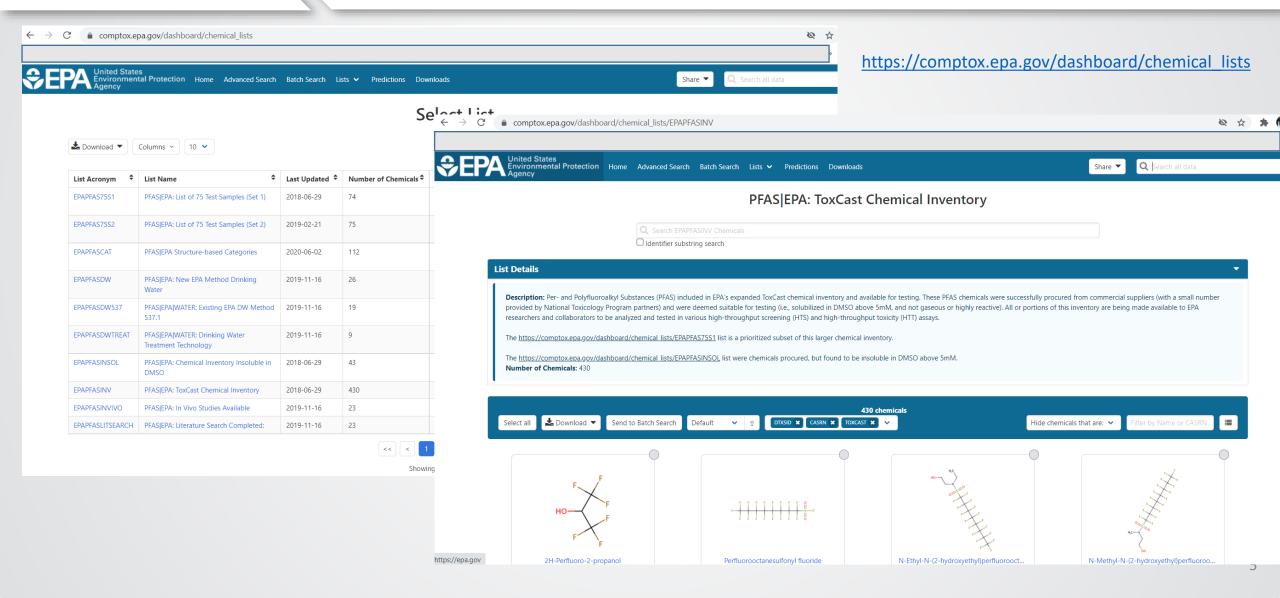


Figure 1. Workflow for selection of structural categories to identify the subset of 75 per- and polyfluoroalkyl substances (PFAS).

### https://comptox.epa.gov/dashboard/chemical lists/?search=PFAS

### Patlewicz et al. 2019

## You can search our lists of chemicals on our public CompTox Chemicals Dashboard, and from there link to data resources



**S**EPA



# Tiered in vitro screening strategy



Society of Toxicology

SOT

## Executing the Next Generation CompTox Blueprint to inform putative chemical hazard



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

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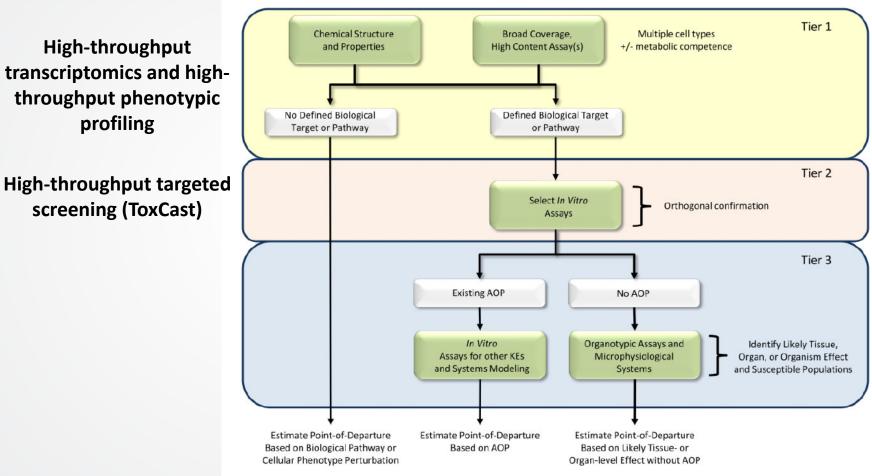


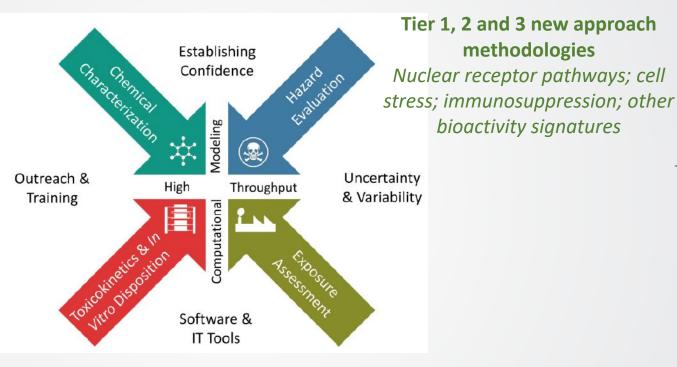
Figure 2. Tiered testing framework for hazard characterization. Tier 1 uses both chemical structure and broad coverage, high content assays across multiple cell types for comprehensively evaluating the potential effects of chemicals and grouping them based on similarity in potential hazards. For chemicals 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.



## Major elements of the EPA CompTox Blueprint can be applied to PFAS screening

methodologies

Analytical quality control (QC)



**Toxicokinetic screening data:** Hepatocellular clearance; renal transport; fraction unbound in plasma

**Public and** soon to be public Tier 2 data are the focus of today's talk



# This talk includes some of the publicly available data from research on PFAS

## EPA's PFAS Action Plan: A Summary of Key Actions

**\$EPA** 

RESEARCH

EPA is rapidly expanding the scientific foundation for understanding and managing risk from PFAS.

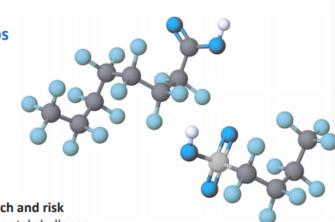
Improved detection and measurement methods, additional information about PFAS presence in the environment and drinking water, better understanding of effective treatment and remediation methods, and more information about the potential toxicity of a broader set of PFAS will help EPA, states, and others better manage PFAS risks.

EPA's PFAS Action Plan outlines concrete steps the agency is taking to address PFAS and to protect public health.

EPA's Per- and Polyfluoroalkyl Substances (PFAS) Action Plan:

- Demonstrates the agency's critical national leadership by providing both short-term solutions and long-term strategies to address this important issue.
- Provides a multi-media, multi-program, national research and risk communication plan to address this emerging environmental challenge.
- Responds to the extensive public input the agency has received over the past year during the PFAS National Leadership Summit, multiple community engagements, and through the public docket.

EPA is taking a proactive, cross-agency approach to addressing PFAS. The key actions EPA is taking to help provide the necessary tools to assist states, tribes, and communities in addressing PFAS are summarized below.





# Targeted screening for nuclear receptor activation and cell stress



Bioactivity profiling of per- and polyfluoroalkyl substances (PFAS) identifies potential toxicity pathways related to molecular structure

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#### ARTICLE INFO ABSTRACT

Keywords: Perfluoroalkyl substances PFAS Transcription factors Chemical safety Nuclear receptors Per- and polyfluoroaliyl substances (FFAS) are a broad class of hundreds of fluorinated chemicals with environmental health concerns due to their widespread presence and pessitence in the environment. Several of these chemicals have been comprehensively studied for experimental toticity, environmental fate and exposure, and human epidemiology; however, most chemicals have limited or no data available. To inform methods for prioritizing these data-poor chemicals for detailed toxicity studies, we evaluated 142 FFAS using an in viro screening platform consisting of two multiplezed transactivation assays encompassing 81 diverse transcription factor activities and tested in concentration-response format ranging from 137 nM to 500 µM. Results showed activity for various machaer receptors, including three known FFAS targets-specifically estrogen receptor alpha and peroxisome proliferator receptors alpha and gamma. We also report activity against the refinal X receptor experiment of the transactivity of experimentitive FFAS against several of these targets. Finally, we identified they feast we confirmed cativity of ergresentitive FFAS against several of these targets. Finally, we identified torue in prioritizing chemicals for risk assessment and in the design of new structures devoid of biological activity.

1. Introduction

Per- and polyfinoroalkyl substances (PFAS) are a class of man-made chemicals that have been in use since the 1940s and are found in a broad array of industrial and consumer products (Glüge et al., 2020). Their common usage as non-stick surface repellants, in fire-fighting foams, in fluoropolymer manufacturing, and in other applications, couled with a tendency of some members of the class to bioaccumulate and be resistant to biodegradation, has led to a high level of concern for their contamination of the environment (Wang et al., 2017). There are well documented, widespread, human and wildlife exposure to some of these chemicals, the best known being perfluorooctanois acid (PFOS; DTXSID0031865) (Kelly et al., 2009; Photomg et al., 2020; Hannen

et al., 2002; Noorlander et al., 2011). These two chemicals are no longer manufactured in the U.S. and their international manufacturing has declined, but other PFAS chemicals have been developed to replace their commercial utility (REACH, 2014; OECD, 2015; Stockholm Convention, 2017; EPA, 2006; EPA, 2017). While the toxicities of PFOA and PFOS have been extensively studied by many researchers, numerous other PFAS have bitle to no toxicity or environmental faite information available. The lack of data and potential environmental after information available. The lack of data and potential environmental after of this class of chemicals led the U.S. Environmental Protection Agency (EPA) and the National Institute of Health's National Toxicology Program (NTP) to collaborate on conducting PFAS toxicity testing to facilitate PFAS have health assessments (Palewicz et al., 2019). A targeted selection of 430 PFAS (https://comptox.epa.gov/dashboard/chemic

Check for

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Houck et al. 2020

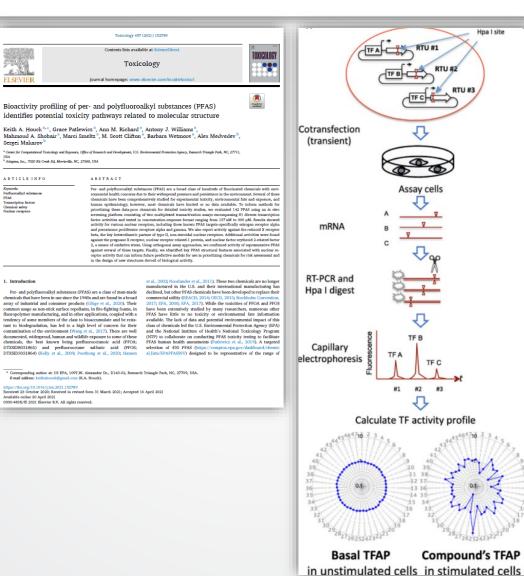
## Gathering information on nuclear receptor and cell stress pathways via transcription factor activity profiling (TFAP)

Hpa | site

>3800 ToxCast chemicals have been screened in concentration response in the Attagene transcription factor profiling system

SEPA

- HepG2 HG19 subclone for elevated xenobiotic metabolic capacity
- "CIS" assays: endogenous transcription factors that regulated transfected reporters (nuclear receptor promoter elements, cell stress)
- "TRANS" assays: exogenous receptorreporter system is transfected in (xenobiotic nuclear receptors)
- Used for environmental mixtures and single chemical screening



| Number | Endpoint    | Go<br>Process                      | Number | Endpoint      | Go                               |
|--------|-------------|------------------------------------|--------|---------------|----------------------------------|
| 1      | GAL4 TRANS  |                                    | 41     | AP_1_CIS      |                                  |
| 2      | M_06_CIS    |                                    | 42     | HIF1a_CIS     |                                  |
| 3      | M 06 TRANS  |                                    | 43     | HSE CIS       | ess                              |
| 4      | M_19_CIS    |                                    | 44     | MRE_CIS       | str                              |
| 5      | M_19_TRANS  |                                    | 45     | NRF1 CIS      | eto                              |
| 6      | M_32_CIS    | 2                                  | 46     | NRF2 ARE CIS  | Suc                              |
| 7      | M 32 TRANS  | control                            | 47     | Oct_MLP_CIS   | response to stress               |
| 8      | M_61_CIS    | 0                                  | 48     | p53_CIS       | 2                                |
| 9      | M 61 TRANS  |                                    | 49     | Xbp1_CIS      |                                  |
| 10     | TA CIS      |                                    | 50     | CRE_CIS       |                                  |
| 11     | TAL_CIS     |                                    | 51     | ERRa_TRANS    | ietic<br>SS                      |
| 12     | CMV_CIS     |                                    | 52     | ERRg_TRANS    | osynthet                         |
| 13     | E Box CIS   |                                    | 53     | GR_TRANS      | biosynthetic<br>process          |
| 14     | E2F_CIS     | ion                                | 54     | GRE_CIS       | ā                                |
| 15     | EGR_CIS     | cell<br>proliferation              | 55     | DR5_CIS       |                                  |
| 16     | Ets_CIS     | olif                               | 56     | RARa_TRANS    |                                  |
| 17     | Pax6_CIS    | ā                                  | 57     | RARb_TRANS    |                                  |
| 18     | AR TRANS    |                                    | 58     | RARg_TRANS    | ion.                             |
| 19     | ERa TRANS   | e l                                | 59     | RXRa_TRANS    | tiat                             |
| 20     | ERE CIS     | reproduction                       | 60     | RXRb_TRANS    | ren                              |
| 21     | THRa1 TRANS |                                    | 61     | NURR1_TRANS   | cell differentiation             |
| 22     | VDR TRANS   |                                    | 62     | RORb_TRANS    | alld                             |
| 23     | VDRE_CIS    | -                                  | 63     | RORg_TRANS    | 0                                |
| 24     | ISRE_CIS    | è e s                              | 64     | RORE_CIS      |                                  |
| 24     | ISRE_CIS    | mmune<br>system<br>process         | 65     | Sox_CIS       |                                  |
| 25     | NF_kB_CIS   | in sy ad                           | 66     | AP_2_CIS      |                                  |
| 26     | IR1_CIS     |                                    | 67     | BRE_CIS       |                                  |
| 27     | FXR_TRANS   | 50                                 | 68     | C_EBP_CIS     |                                  |
| 28     | DR4_LXR_CIS | Seces                              | 69     | FoxA2_CIS     | ent                              |
| 29     | LXRa_TRANS  | bro                                | 70     | FoxO_CIS      | mq                               |
| 30     | LXRb_TRANS  | olic                               | 71     | GATA_CIS      | velo                             |
| 31     | PPARa_TRANS | tab                                | 72     | GLI_CIS       | de                               |
| 32     | PPARd_TRANS | lipid metabolic process            | 73     | HNF4a_TRANS   | snatomical structure development |
| 33     | PPARg_TRANS | bid                                | 74     | HNF6_CIS      | DD D                             |
| 34     | PPRE_CIS    |                                    | 75     | Myb_CIS       | str                              |
| 35     | SREBP_CIS   |                                    | 76     | Myc_CIS       | Ca.                              |
| 36     | Ahr_CIS     | 0.0                                | 77     | NFI_CIS       | tom                              |
| 37     | CAR_TRANS   | xenobiotic<br>metabolic<br>process | 78     | Sp1_CIS       | nat                              |
| 38     | PBREM_CIS   | kenobiotik<br>metabolic<br>process | 79     | STAT3_CIS     | 10                               |
| 39     | PXR_TRANS   | pre                                | 80     | TCF_b_cat_CIS |                                  |
| 40     | PXRE_CIS    |                                    | 81     | TGFb_CIS      |                                  |



# There are differences in assay sensitivity by mode and receptor, based on expression and design differences.

#### Table 1

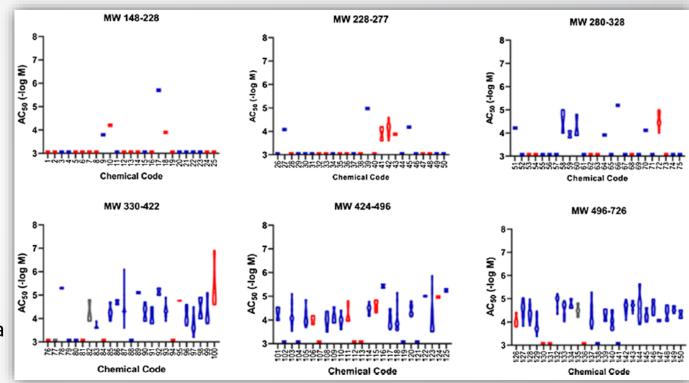
#### Nuclear receptors included in FACTORIAL-TRANS assay.

| #  | Abbreviation | Receptor Name                                    | Nomenclature | Reference Agonist (Fold-<br>Increase) | <i>cis</i> -Factorial Assay<br>(Fold-Increase) | Receptor Expression in $HepG2^1$                                  |
|----|--------------|--|--------------|---------------------------------------|--|---|
| 1  | FXR          | Farnesoid X receptor                             | NR1H4        | Lithocholic acid (3.5)                | IR1 (1.9)                                      | Moderate  |
| 2  | AR           | Androgen receptor                                | NR3C4        | Testosterone propionate<br>(44.1)     | NA   | Very low  |
| 3  | RARγ         | Retinoic acid receptor-γ                         | NR1B3        | All-trans retinoic acid (3.9)         | DR5 (20.2)                                     | Moderate (RAR subfamily) <sup>2</sup>                             |
| 4  | GAL4         | Yeast GAL4, negative control                     | GAL4         | NA                                    | NA   | NA  |
| 5  | RXRα         | Retinoid X receptor-α                            | NR2B1        | Bexarotene (18.5)                     | DR5 (8.3)                                      | Moderate (RXR subfamily) <sup>2</sup>                             |
| 6  | GR           | Glucocorticoid receptor                          | NR3C1        | Betamethasone (29.1)                  | GRE (4.6)                                      | Moderate  |
| 7  | RARβ         | Retinoic acid receptor-β                         | NR1B2        | All-trans retinoic acid (1.6)         | DR5 (20.2)                                     | Moderate (RAR subfamily) <sup>2</sup>                             |
| 8  | RARα         | Retinoic acid receptor-α                         | NR1B1        | All-trans retinoic acid (5.5)         | DR5 (20.2)                                     | Moderate (RAR subfamily) <sup>2</sup>                             |
| 9  | PPARγ        | Peroxisome proliferator-<br>activated receptor-γ | NR1C2        | Rosiglitazone maleate (44.8)          | PPRE (3.8)                                     | High  |
| 10 | ERRγ         | Estrogen-related receptor-γ                      | NR3B3        | 4-Nonylphenol, branched<br>(2.7)      | NA   | NA  |
| 11 | RORβ         | RAR-related orphan receptor-β                    | NR1F1        | SSR69071 (7.8)                        | RORE (5.9)                                     | NA  |
| 12 | ERα          | Estrogen receptor-α                              | NR3A1        | 17β-Estradiol (22.6)                  | ERE (19.1)                                     | Very low; full-length human ERα co-<br>expressed in FACTORIAL-CIS |
| 13 | LXRα         | Liver X receptor-α                               | NR1H3        | Lynestrenol (13.9)                    | DR4 (2.3)                                      | High (LXR subfamily) <sup>2</sup>                                 |
| 14 | ERRα         | Estrogen-related receptor- $\alpha$              | NR3B1        | 4-Nonylphenol, branched<br>(2.7)      | NA   | NA  |
| 15 | PXR          | Pregnane X receptor                              | NR1I2        | Rifampicin (3.8)                      | PXRE (9.1)                                     | Moderate; full-length human PXR co-<br>expressed in FACTORIAL-CIS |
| 16 | ΤRα          | Thyroid hormone receptor- $\alpha$               | NR1A1        | 3,5,3′-Triiodothyronine<br>(33.0)     | NA   | High  |
| 17 | LXRβ         | Liver X receptor-β                               | NR1H2        | Lynestrenol (8.7)                     | DR4 (2.3)                                      | High (LXR subfamily) <sup>2</sup>                                 |
| 18 | CAR          | Constitutive androstane receptor                 | NR1I3        | p,p'-DDT (3.5)                        | PBREM (1.0)                                    | Very low  |
| 19 | ΡΡΑΒα        | Peroxisome proliferator-<br>activated receptor-α | NR1C1        | Pirinixic acid (14.1)                 | PPRE (2.4)                                     | Moderate  |
| 20 | RORy         | RAR-related orphan receptor-y                    | NR1F3        | SSR69071 (14.2)                       | RORE (5.9)                                     | NA  |
| 21 | RXRβ         | Retinoid X receptor- $\beta$                     | NR2B2        | Bexarotene (15.2)                     | DR5 (8.3)                                      | Moderate (RXR subfamily) <sup>2</sup>                             |
| 22 | HNF4α        | Hepatocyte nuclear factor-4-α                    | NR2A1        | NA                                    | NA   | High  |
| 23 | NURR1        | Nuclear receptor related 1                       | NR4A2        | Bexarotene (24.6)                     | NA   | NA  |
| 24 | VDR          | Vitamin D receptor                               | NR1I1        | Ergocalciferol (32.6)                 | VDRE (1.2)                                     | Very low  |
| 25 | ΡΡΑRδ        | Peroxisome proliferator-<br>activated receptor-δ | NR1C3        | 12-Hydroxyoctadecanoic<br>acid (9.3)  | PPRE (2.9)                                     | NA  |

- Low- to negligible-expression in HepG2 cells of ERα and PXR was overcome by cotransfection of fulllength receptors in the TRANS assay
- CAR and VDR have very low sensitivity to ligands due to reliance only on endogenous receptor expression in the host cell.

## As with other assay platforms screened, lower MW often corresponded to more limited bioactivity, but there may be more than one reason.

- PFAS with molecular weight less than 330 g/mol appeared less likely to be active in the Attagene assays and more likely to "fail" analytical QC (defined as parent structure not detected).
- Activity was not detected for 76 distinct structures, of which 55 % failed analytical QC.
- 67% of the "failed" samples had predicted vapor pressures in excess of 100 mmHg, suggesting that chemical volatilization may have played a role in limited bioactivity of some of these samples.
- The specific acid form of PFAS may also be important, as the free acid form of the chemical known as "GenX" (perfluoro-2-methyl-3oxahexanoic acid (DTXSID70880215) did not have a high vapor pressure (was unlikely to have volatilized), but the ammonium salt form of this chemical (DTXSID40108559) showed activity as a PPARα agonist when solubilized in water (rather than DMSO).

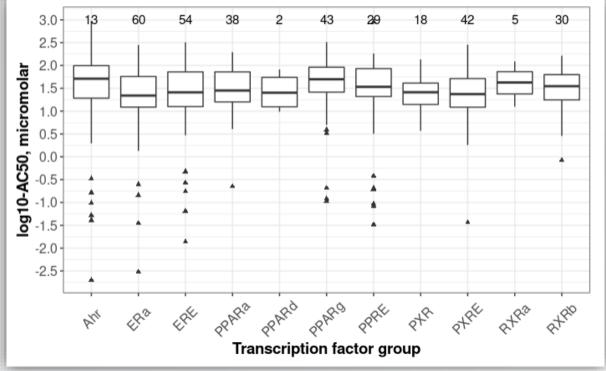


Houck et al. 2020, Fig1B.



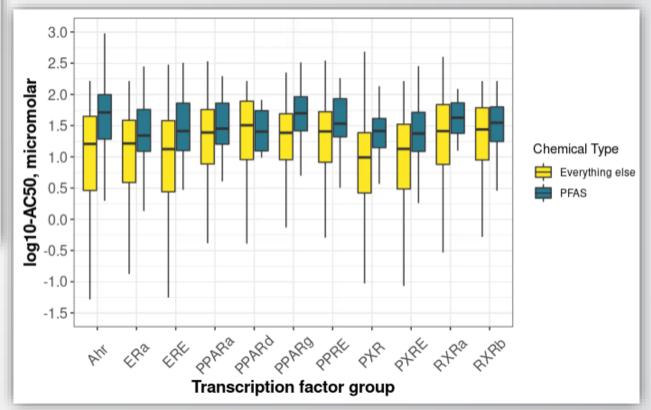
Potency for the PFAS that were positive at key transcription factor targets tended to be somewhat left-shifted from the rest of the ToxCast library

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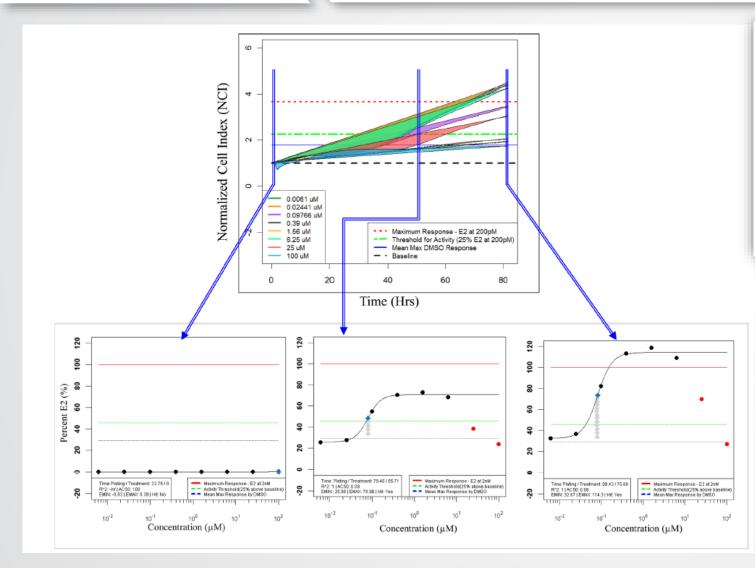
- Many PFAS were negative in the transcription factor activity screening
- Aryl hydrocarbon receptor (AhR), estrogen receptor alpha (ERa), PPAR alpha, delta, and gamma (PPARa,d,g), the pregnane X receptor (PXR), and RXR alpha and beta (RXRa,b) emerged as targets.

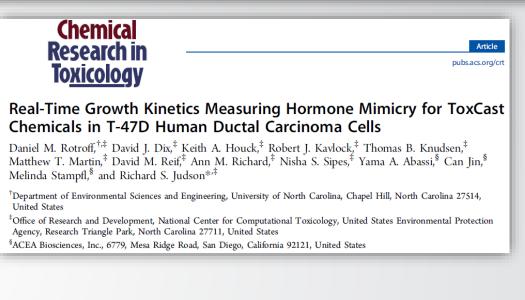
The number of chemicals that simply hit one or more relevant assays for a particular transcription factor group can be examined in more depth for confirmation.





Estrogen receptor activity can be confirmed with orthogonal assays including ACEA: Real Time Cell Analysis Based on Electrical Impedance

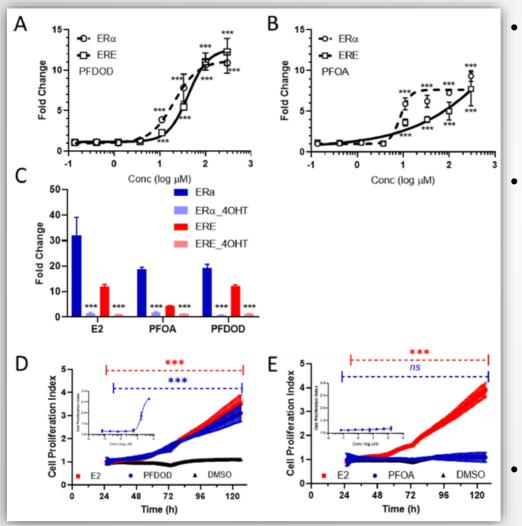




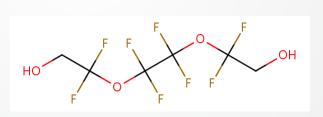
- Can measure cell proliferation or cytotoxicity depending on the direction
- Electrical impedance measured over 80 hr
- ACEA ER assay uses T-47D breast cancer cells

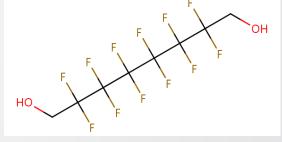
# **\$EPA**

# Confirmation of transcriptional responses with functional activity is an important strategy for ER bioactivity



- 40-60 PFAS demonstrated some activity in the ATG ERa TRANS or ERE CIS assays; viewing these assays as orthogonal reduces the set to <10.</li>
  - All of these were less potent than  $17\beta$ -estradiol.
  - Acrylates and N-akyl perfluoroalkyl (linear) sulfonamide structural categories were significantly associated with ER activity.
- Adding in ACEA as another orthogonal assay to confirm specificity leads indicates few PFAS with transcription factor *and* functional ERdependent cell proliferation.





1H,1H,8H,8H-Perfluoro-3,6-dioxaoctane-1,8-diol

• PFOA activated ATG\_ERa\_TRANS and ERE\_CIS but failed to produce functional ER-dependent cell proliferation in ACEA.

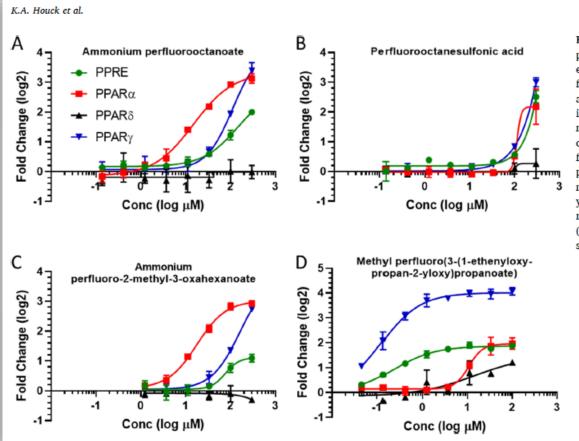
Houck et al. 2020, Fig5.

<sup>1</sup>H,1H,8H,8H-Perfluorooctane-1,8-diol

**Set EPA**

# As expected PPAR activity was observed for a subset of PFAS.

- The TRANS assay contained endpoints for all three human PPARs (α,δ,γ) whereas the CIS assay contained a reporter gene controlled by a PPARresponse element that responds to all three PPARs endogenously expressed in the HepG2 host cells.
- Functional groups enriched within the actives were mostly carboxylates along with sulfonates, sulfonamides and a thenoylketone, which all have a negative ionic charge at physiological pH, consistent with known critical components for ligand-binding.
- Not much activity at PPARδ (smaller binding pocket?).

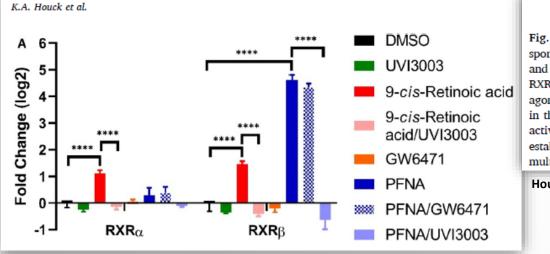


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Fig. 6. Transactivation of the peroxisome proliferator-activated receptors (PPARs) by example PFASs. Concentration-response data for PPAR- $\alpha$ , - $\delta$ , and - $\gamma$  in the FACTORIAL-TRANS assays and the PPAR response element (PPRE) in the FACTORIAL-CIS assay following treatment for 20-24 h with increasing concentrations of ammonium perfluorooctanoate (A), perfluorooctanesulfonic acid (B), ammonium perfluoro-2-methyl-3-oxahexanoate (C), and methyl perfluoro(3-(1-ethenvloxypropan-2yloxy)propanoate) (D). Values are the mean reporter gene activity expressed as fold-change (log2) normalized by solvent control (dimethyl sulfoxide) values.

Houck et al. 2020, Fig6.

# ~17 PFAS activated RXR $\beta$ , with two of these active at RXR $\alpha$



S FPA

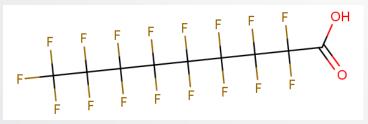
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Fig. 8. PFAS activity for retinoid X receptors (RXR). A) Responses of RXR $\alpha$  and RXR $\beta$  to perfluorononanoic acid (PFNA) and effects of pharmacological agents UVI3003 (5  $\mu$ M), a pan-RXR antagonist; 9-cis retinoic acid (0.02  $\mu$ M), a pan-RXR agonist; and GW6471 (5  $\mu$ M), a PPAR $\alpha$ -selective antagonist; in the presence and absence of PFNA (66  $\mu$ M). No significant activation of RXR $\alpha$  by PFNA was observed. Significance was established with an ordinary one-way ANOVA and Tukey's multiple comparisons test. (\*\*\*\* = P < .0001). B) Radioligand

### Houck et al. 2020, Fig8A.

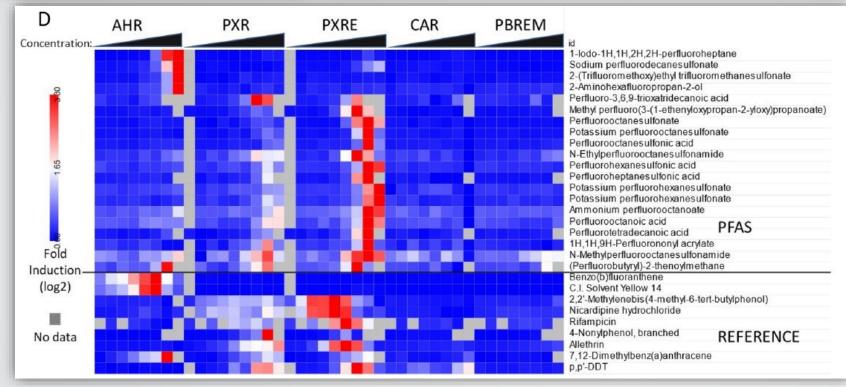
- Seventeen of the PFAS, mostly linear, fluorinated carboxylic acids, showed a novel finding of activation of RXRβ.
- Most also activated PPARα, PPARγ and NRF2, with varying levels of selectivity. Only two activated RXRα; however, NURR1 was activated, presumably through agonist effects on RXRβ.
- All are structurally related perfluorinated carboxylic acids and meet defined ligand structural requirements for RXR.

PFNA appears to work through RXR specifically: an RXR-selective antagonist, UVI3003 (DTXSID501024375), completely blocked PFNA activation of RXR, whereas the PPARα antagonist GW6471 was ineffective.



# **\$EPA**

# Xenobiotic nuclear receptor responses associated with hepatic metabolism may also be important targets to screen for PFAS bioactivity.



Houck et al. 2020, Fig3B.

- Many of the PFAS modulated the xenobiotic response, particularly PXR.
- Responses were generally modest with respect to potency and efficacy relative to prototypical PXR inducers.
- None of the PFAS were determined to be CAR activators, recognizing limitations in the FACTORIAL-CIS assay for CAR, likely due to negligible expression of CAR in HepG2 cells.
- Several PFAS structures activated the AhR, somewhat surprising in that all were linear fluoroalkyl molecules while the protypical activator is a polycyclic aromatic hydrocarbon. Except for sodium perfluorodecanesulfonate and 1-lodo-1H,1H,2H,2H-perfluoroheptane, the responses were very weak with unknown *in vivo* relevance.



# Immunosuppressive activity of the PFAS150 in an *in vitro* assay suite

Houck KA, Paul Friedman K, Feshuk M, Patlewicz G, Smeltz M, Clifton MS, Wetmore BA, Velichko S, Berenyi A, Berg EL. (*In internal review*). Evaluation of 147 Perfluoroalkyl Substances for Immunosuppressive and Other Activities through Phenotypic Screening of Human Primary Cells.



## PFOA and PFOS are suspected of being immunosuppressive

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## Exposure to per- and polyfluoroalkyl substances leads to immunotoxicity: Epidemiological and toxicological evidence

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### Abstract

In this perspective, we evaluate key and emerging epidemiological and toxicological data concerning immunotoxicity of perfluorooctanoic acid (PFOA) and perfluorooctane sulfonate (PFOS) and seek to reconcile conflicting conclusions from two reviews published in 2016. We summarize ways that immunosuppression and immunoenhancement are defined and explain how specific outcomes are used to evaluate immunotoxicity in humans and experimental animals. We observe that different approaches to defining immunotoxicological outcomes, particularly those that do not produce clinical disease, may lead to different conclusions from epidemiological and toxicological studies. The fundamental point that we make is that aspects of epidemiological studies support epidemiological findings. Taken together, we find that results of epidemiological studies, supported by findings from toxicological studies, provide strong evidence that humans exposed to PFOA and PFOS are at risk for immunosuppression.

- A 2016 NTP review of PFOA/PFOS had concluded that suppression of antibody responses in animals had the most evidence.
- Chang et al. 2016 had concluded that the available evidence was insufficient to make a conclusion about causality; could cause immunosuppression in animals, but inconsistencies were present across species and strain.
- DeWitt et al. 2019 attempts to reconcile these opinions and concludes what the NTP concluded, that PFOA and PFOS may be associated with immunosuppression based on available data.
- The limited data landscape, focused on PFOA and PFOS, is complex, with differences by species, strain, sex, endpoints measured, and doses used.



# BioMAP is a suite of primary and primary co-culture models for assessing phenotypes, including some elements of immunosuppression

| System | lcon | Cell Type   | Disease Relevance  | Biomarker Readouts   | Description  |
|--------|------|---|--|--|--|
| SAg    |      | Peripheral blood<br>mononuclear cells<br>+ Venular endothelial<br>cells | Autoimmune<br>Disease, Chronic<br>Inflammation                         | MCP-1, CD38, CD40, E-selectin,<br>CD69, IL-8, MIG, PBMC<br>Cytotoxicity, Proliferation, SRB        | The SAg system models chronic inflammation of the Th1 type and<br>T cell effector responses to TCR signaling with costimulation. This<br>system is relevant to inflammatory conditions where T cells play a<br>key role including organ transplantation, rheumatoid arthritis,<br>psoriasis, Crohn's disease and multiple sclerosis. |
| вт     |      | Peripheral blood<br>mononuclear cells<br>+ B cells                      | Asthma, Allergy,<br>Oncology,<br>Autoimmunity                          | B cell Proliferation, PBMC<br>Cytotoxicity, Secreted IgG, sIL-17A,<br>sIL-17F, sIL-2, sIL-6, sTNFα | The BT system models T cell dependent B cell activation and class<br>switching as would occur in a germinal center. This system is<br>relevant for diseases and conditions where B cell activation and<br>antibody production are relevant. These include autoimmune<br>disease, oncology, asthma and allergy.                       |
| Mphg   |      | Venular endothelial<br>cells<br>+ Macrophages                           | Cardiovascular<br>Inflammation,<br>Restenosis, Chronic<br>Inflammation | MCP-1, MIP-1α, VCAM-1, CD40,<br>E-selectin, CD69, IL-8, IL-1α, M-<br>CSF,<br>sIL-10, SRB, SRB-Mphg | The <i>M</i> phg System models chronic inflammation of the Th1 type<br>and macrophage activation responses. This system is relevant to<br>inflammatory conditions where monocytes play a key role including<br>atherosclerosis, restenosis, rheumatoid arthritis, and other chronic<br>inflammatory conditions.                      |

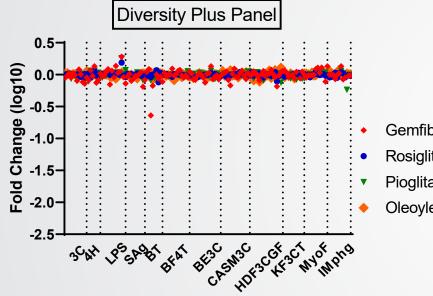
A subset of BioMAP includes biomarkers in relevant cell types with measures of specific immune cells and interleukins known to be immune-relevant.

Does not cover every type of cell involved in immune suppression nor every marker

- A TDAR will measure T-cell dependent IgM antibody response, can also IgG if modified.
- SAg: markers of decreased T-cell proliferation or specifically cytotoxic to PBMCs.
- BT: Decreased IgG and B cell proliferation or specifically cytotoxic to PBMCs.
- Mphg: Decreased IL-10 (like dexamethasone)
- Reference chemicals used: cyclosporin A, azathioprine, methotrexate [these 3 strongly suppress IgG production in BT], and dexamethasone [affected IL-10 in Mphg]

# SEPA

## It's unlikely that PPAR is related to any immunosuppressive activity in the BioMAP assay suite as PPAR agonists have little activity

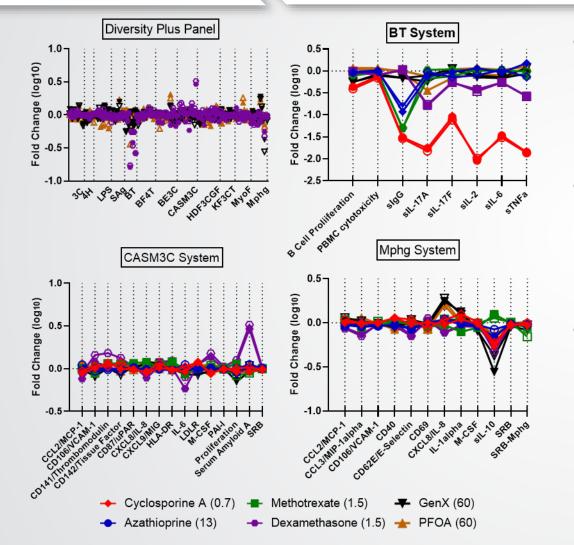


- Gemfibrozil Rosiglitazone Pioglitazone
- Oleoylethanolamide

**BioMAP Diversity Plus profiles of PPARa and PPARg reference** chemicals. Profiles for PPARg agonists rosiglitazone  $(3.7 \,\mu\text{M})$ , pioglitazone (10  $\mu$ M) and PPARa agonists gemfibrozil (200  $\mu$ M) and oleoylethanolamide  $(1.1 \,\mu\text{M})$  are shown for the 12 assay systems of the BioMAP Diversity Plus platform. Concentrations were selected from the database to exceed reported in vitro EC50's for the corresponding receptor targets by 5- to 40-fold.

## **Set EPA**

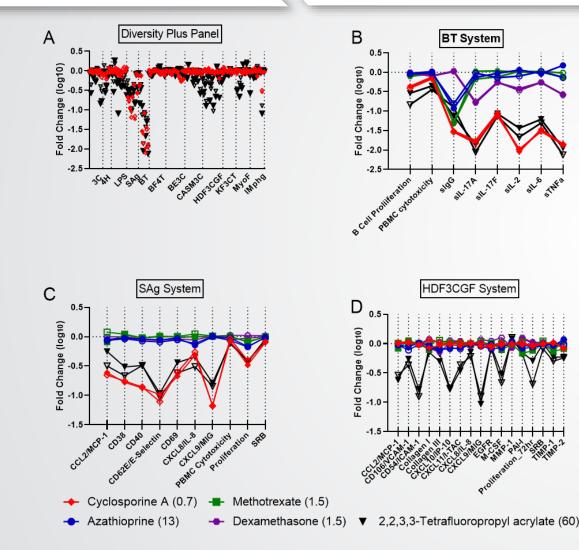
# PFOA (and PFOS) failed to provide support for suppression of IgG in BioMAP at the screened concentration range



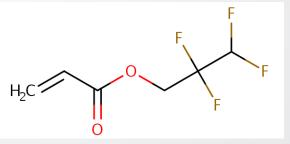
- Overall, neither PFOA nor PFOS seemed to show activity similar to the reference immunosuppressants except perhaps PFOA at its highest testing concentration of 60 μM where it clustered with dexamethasone.
- PFOA and PFOS decreased IL-10 in a co-culture system (Mphg assay) that detects macrophage responses. IL-10 is a cytokine that promotes B cell IgG production, but human relevance of this in vitro finding is unknown.
- Several other PFAS, *i.e.* 3-Bis(trifluoromethyl)-2-propenoic acid, 3H-perfluoro-2,2,4,4-tetrahydroxypentane and perfluoropinacol, have activities similar to the reference immunosuppressants in some of the cell systems, including suppression of IgG secretion.

# **\$EPA**

# A single PFAS at its highest screened concentration associated closely with cyclosporin



- Suppression of multiple endpoints in the BT system and the SAg system is similar for both chemicals with strong reduction in secreted IgG and the cytokines IL-17A, IL-2, IL-6 and TNFα in the BT assay.
- Notably, while cyclosporine A was very selectively active for these two assay systems, 2,2,3,3-tetrafluoropropyl acrylate was also active in others, in particular the wound healing and inflammation model (HDF3CGF).



2,2,3,3-Tetrafluoropropyl acrylate



# Overall conclusions of these high-throughput screening data for a PFAS library

- PFAS with MW >330 g/mol tend to be more active in vitro in the current aqueous media, cell-based assays.
- Analytical quality control is exceedingly important particularly for PFAS which may include substances that volatilize. Additionally the specific salt form of a PFAS greatly impacts its bioactivity.
- In general, the PFAS tend to be similarly or less potent than the rest of the ToxCast chemical library for any of the targets screened to date.
- Subsets of PFAS have activity for various nuclear receptor targets, and the use of orthogonal assays *in vitro* can further inform interpretation of these transcription factor targets. Screening for AhR, ER, PPAR, PXR, and RXR may be important.
- Work in progress on research-based screening models of immunosuppression fails to support PFOA and/or PFOS induced IgG suppression.

# 

## Acknowledgements

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Barbara Wetmore

Antony Williams

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|-----------------|--------------------|---------------------------|
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| Sergei Markarov | Sergei Markarov    | Sharlene Velichko         |

Antal Berenyi



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