

Hepatic Transcriptome Comparative Analysis Reveals Similar Pathways and Targets Altered by Legacy and Alternative Per- and Polyfluoroalkyl Substances

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**Center for Computational Toxicology and Exposure
Office of Research and Development
Environmental Protection Agency**



Toxics Webinar | Issue Cover Authors of 2023

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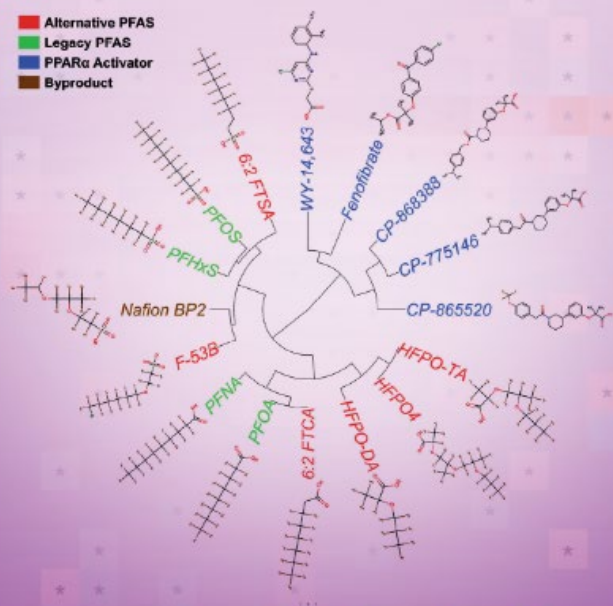


IMPACT
FACTOR
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PubMed

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Alternative PFAS
Legacy PFAS
PPARα Activator
Byproduct






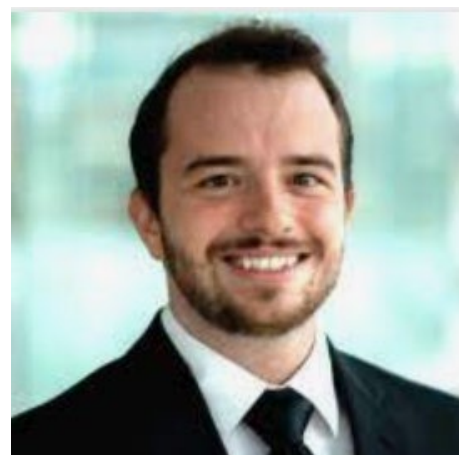
Comparative Analysis of Hepatic Transcriptome Altered by Eleven PFAS in Mice

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Article

Hepatic Transcriptome Comparative In Silico Analysis Reveals Similar Pathways and Targets Altered by Legacy and Alternative Per- and Polyfluoroalkyl Substances in Mice

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Pfizer Pharmaceutical Company**

**2021 Society of Toxicology
Colgate-Palmolive Awardee for
Student Research Training in
Alternative Methods**

Chris Lau – EPA

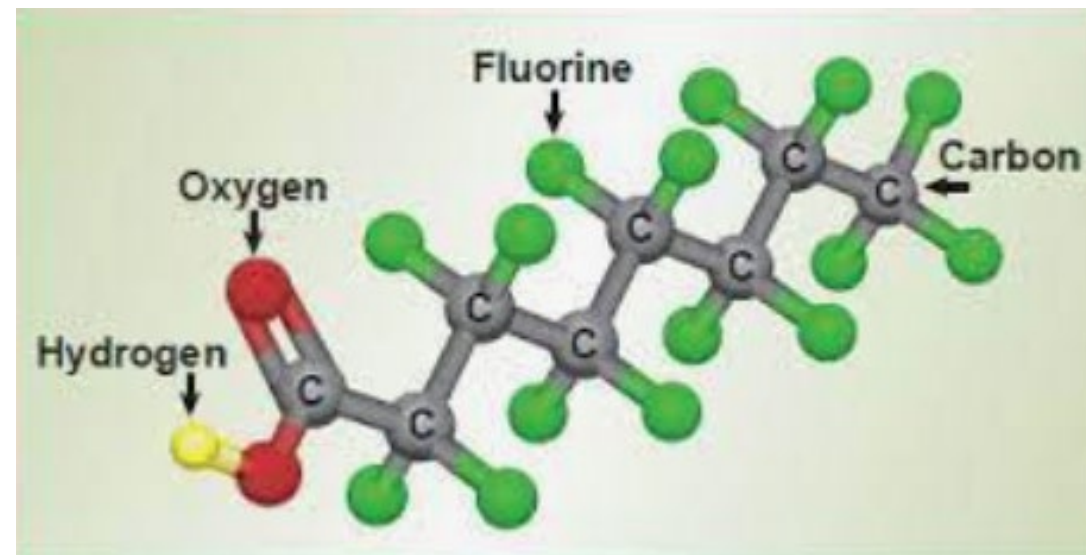
Jiayin Dai – Shanghai Jiao Tong University, China

Udayan Apte – University of Kansas Medical Center

What will be covered

- **Background on PFAS**
- **Summary of work in our Center related to PFAS**
- **The Robarts et al. study**
 - **Why we did it**
 - **What we found**

PFAS = Per- and poly-fluoroalkyl substances



<https://www.xdd-llc.com/remediation/pfas-introduction-and-chemical-properties/>

Why do we care about PFAS?

- They are everywhere and environmentally persistent
 - Globally distributed, detected in water, air, house dust, soil, sediment, sludge from wastewater treatment plants, biosolids
 - Non-biodegradable in environment
 - Found in consumer products



Why do we care about PFAS?

- They are present in the blood of humans and wildlife

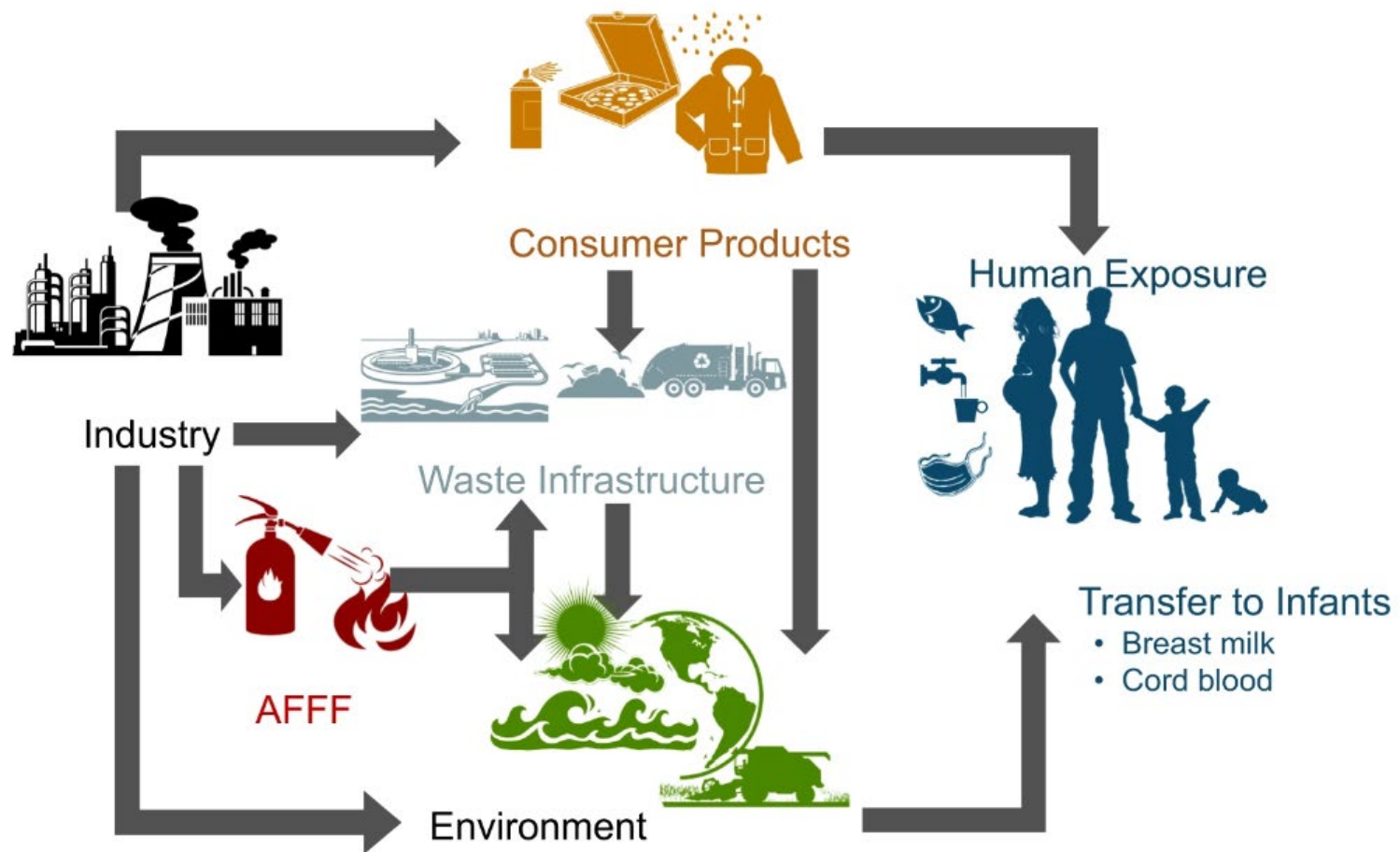
<i>Serum Levels (ppb)</i>	PFOS (C8)	PFOA (C8)	PFHxS (C6)	PFNA (C9)	PFDA (C10)
Production workers	1,500-2,000	500-1,000	~500	N/A	N/A
NHANES '99-'00	30.4	5.2	2.1	0.6	N/A
NHANES '01-'02	20.8	3.7	2.8	0.6	N/A
NHANES '03-'04	20.7	3.9	1.9	1.0	N/A
NHANES '05-'06	17.1	3.9	1.7	1.1	0.36
NHANES '07-'08	13.2	4.1	2.0	1.2	0.29
NHANES '09-'10	9.3	3.1	1.7	1.3	0.28
NHANES '11-'12	6.3	2.1	1.3	0.9	0.20
NHANES '13-'14	5.0	1.9	1.4	0.7	0.19
NHANES '15-'16	4.7	1.6	1.2	0.6	0.15
NHANES '17-'18	4.3	1.4	1.1	0.4	0.19
Arnsburg, Germany '06	5.8-10.5	23.4-25.3	1.1-2.5	N/A	N/A
Little Hocking, WV '07	19.2	32.9	3.3	1.4	0.4
Lake trout	121	4.4	0.6	2.9	N/A
Polar bear	~1,200	~10	--	~100	N/A



<https://thehill.com/>

Why do we care about PFAS?

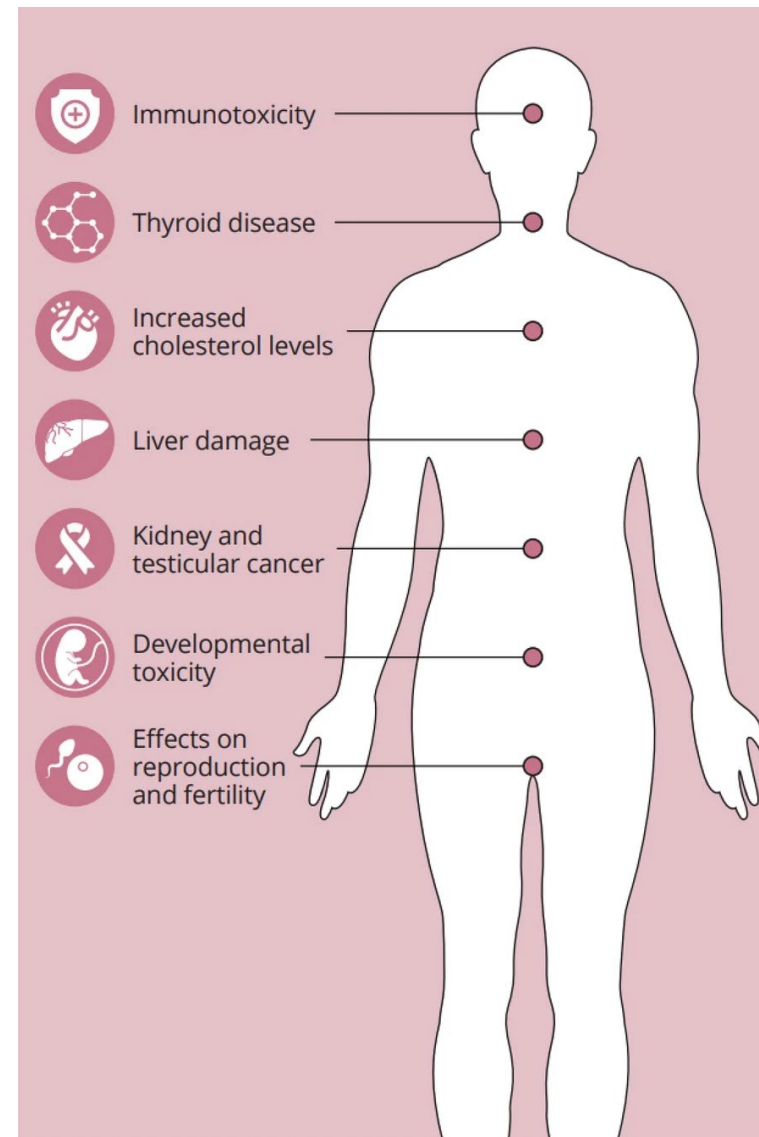
- People can be exposed to PFAS through multiple routes
 - food (fish, some edible plants), migration from food packaging
 - drinking water (contamination sites)
 - house dust, air
 - Consumer products



<https://www.nature.com/articles/s41370-018-0094-1/figures/1>

Why do we care about PFAS?

- Toxic effects have been identified in animal studies
 - Hepatotoxicity, reproductive and developmental toxicity, immunotoxicity, tumor induction, endocrine disruption, neurotoxicity
- Epidemiology studies indicate adverse effects in humans
- Human health risks assessed by regulatory bodies internationally
 - Legally enforceable Maximum Contaminant Levels (MCLs) for 5 PFAS in drinking water were issued by the US EPA on 4/10/2024:
 - PFOA (4 ppt),
 - PFOS (4 ppt)
 - PFHxS (10 ppt),
 - PFNA (10 ppt),
 - HFPO-DA (GenX, 10 ppt)

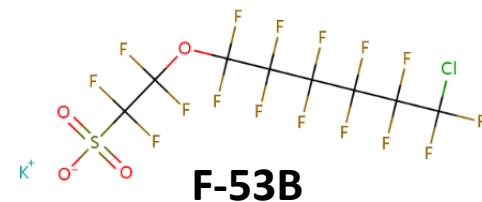


<https://enveurope.springeropen.com/articles/10.1186/s12302-023-00721-8/figures/2>

Slide modified from Dr. Chris Lau

Why do we care about PFAS?

- Some PFAS are very persistent in the blood of humans

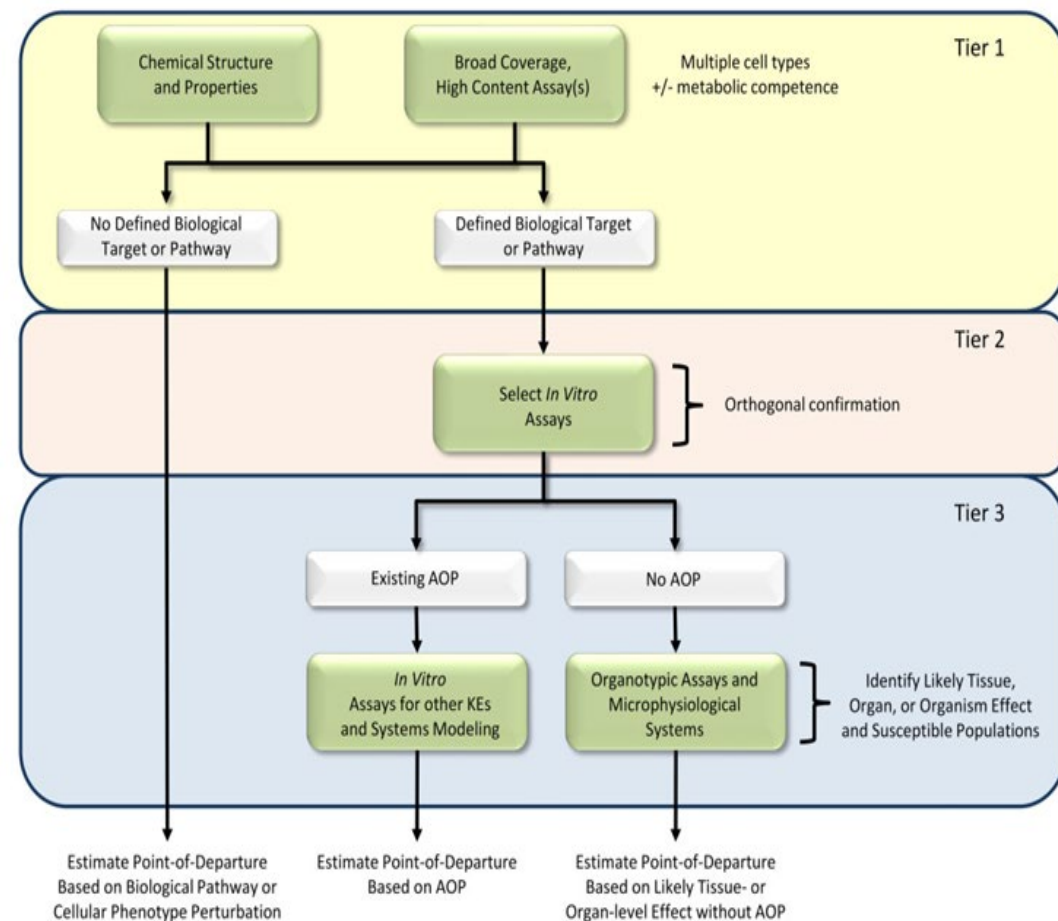


Serum half-life estimates of some per- and polyfluoroalkyl substances

	PFBS (C4)		PFHxS (C6)		PFOS (C8)		PFBA (C4)		PFHxA (C6)		PFHpA (C7)		PFOA (C8)		PFNA (C9)		PFDA (C10)		F-53B		GenX	
	F	M	F	M	F	M	F	M	F	M	F	M	F	M	F	M	F	M	F	M	F	M
<i>Rat</i>	4.0 hours	4.5 hours	1.4 days	26-27 days	28-43 days	34-36 days	1.8 hours	9.2 hours	0.5-7.3 hours	1.3-11 hours	1.2-2.1 hours	1.5-24 hours	1.7-4.8 hours	8.1-8.5 days	6.4 days	31-55 days	45-59 days	55-83 days			0.9-2.8 days	3.0-3.7 days
<i>Mouse</i>	4.5 hours	5.8 hours	25-27 days	28-30 days	31-38 days	36-43 days	6.2 hours	12 hours	~1.2 hours	~1.6 hours			16 days	22 days	26-68 days	34-69 days					1.0 day	1.5 days
<i>Monkey</i>	1.1 days	1.6 days	87 days	140 days	110 days	130 days	1.7 days		2.4 hours	5.3 hours			30 days	21 days							3.3 days	2.7 days
<i>Humans</i>	35 days	36 days	13 years	14 years	3.4 years	3.7 years	3 days		32 days		140 days	130 days	2.1-3.8 years		1.7 years	3.2 years			15.3 years		3.4 days	

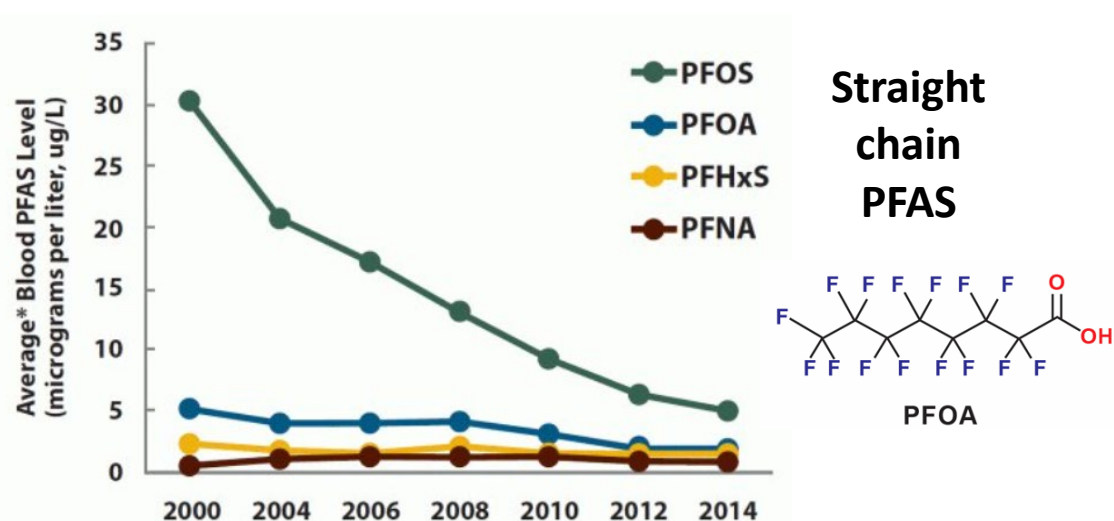
Center for Computational Toxicology and Exposure – Understanding the potential risks to PFAS exposure

- Considering the universe of PFAS (thousands)
- Tiered testing strategies
 - Tier 1 – high throughput testing examining cellular transcript or phenotypic changes in different cell lines; computational approaches for extrapolating test concentrations to exposure in humans; grouping by structure and biology allowing read across
 - Tier 2 – in vitro assays that are part of our new approach methodologies (NAMs)
 - Tier 3 – expose rats to different doses of individual PFAS to generate benchmark doses (EPA transcriptomic assessment product; ETAP)



Background and Goal of Study

The levels of legacy PFAS in human blood are decreasing

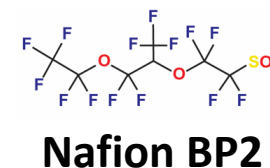


Brennan et al. (2021) Intl J Environ Res Pub Health 18:10900

The levels of alternative PFAS are increasingly detectable

- Increased blood levels of per- and polyfluoroalkyl ether acids (PFEAs)

- Kotlarz et al. (2024) EHP 132(2):27701; 132(2):27702
- Rosen et al. (2022) EHP 130(9):97002

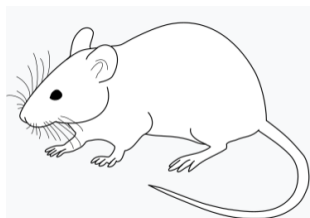


- Goal of study: Use available archived transcript profile data to identify molecular targets of legacy and alternative PFAS**

- Are there differences?
- How do alternative PFAS toxicities compare to toxicities of legacy PFAS?

Methods

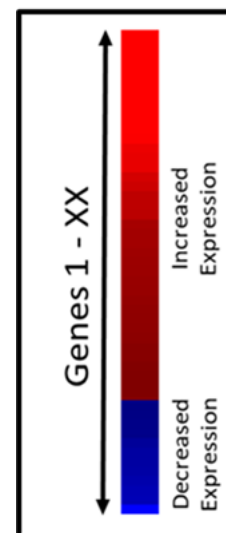
**Treat mice with 11
legacy or
alternative PFAS**



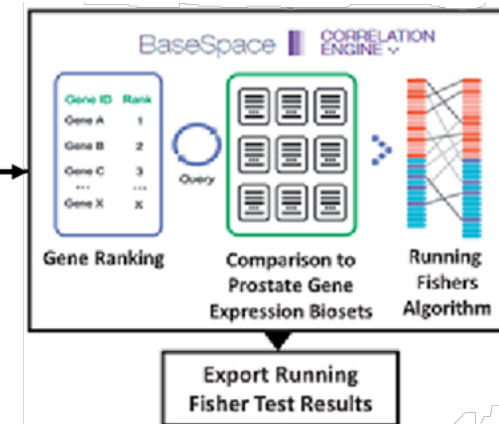
**Collect liver;
Isolate RNA;
Measure
transcript profile
changes using
microarrays or
RNA-Seq**



**Archived
transcript
profiles**

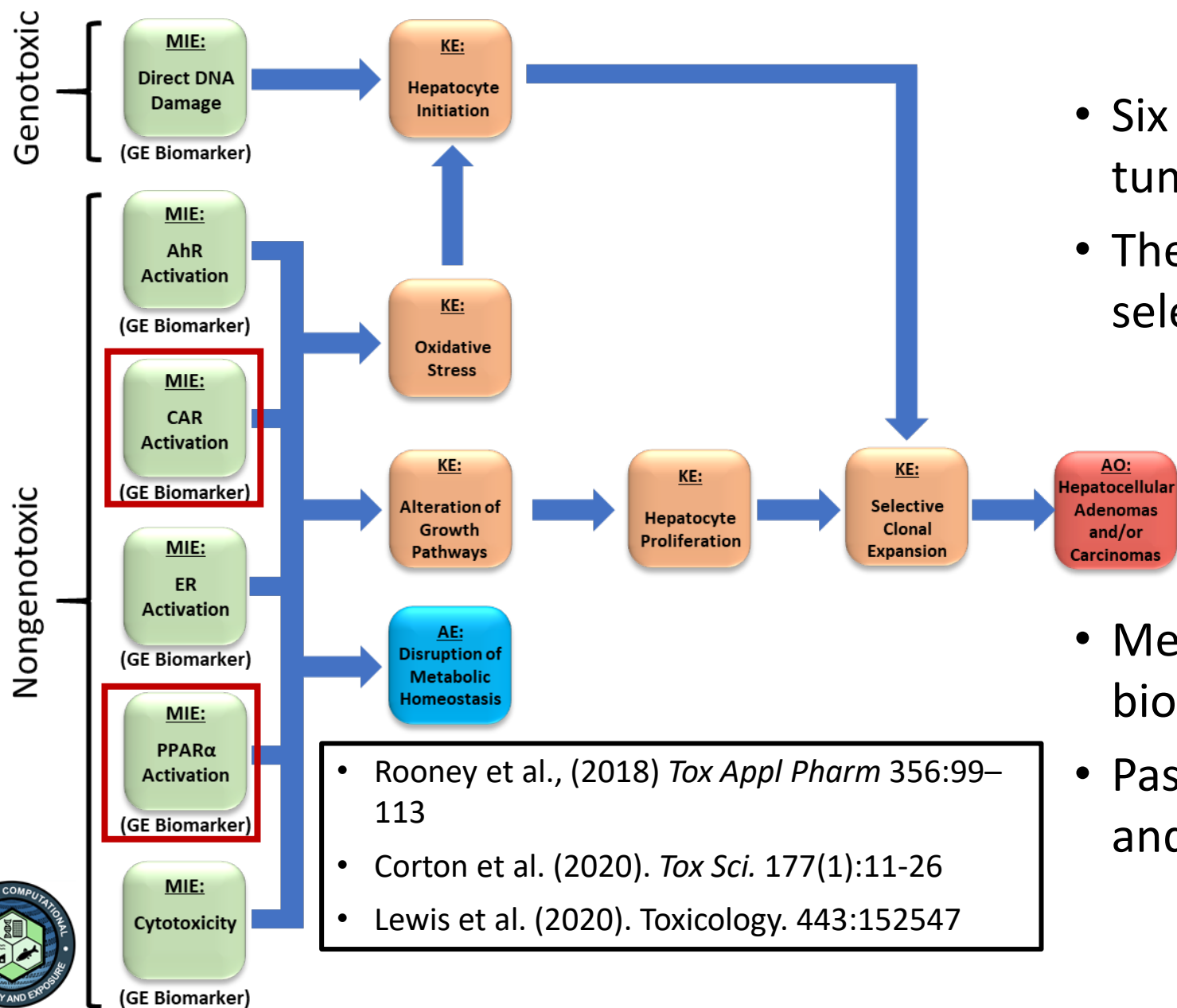


**Illumina's BaseSpace
Correlation Engine**



**Compare to gene lists
and biomarkers to
understand biological
effects**

Major Adverse Outcome Pathways That Lead to Rodent Liver Tumors



- Six major AOPs lead to rodent liver tumors
- The AOPs converge on the key event of selective clonal expansion

- Measure MIEs with gene expression biomarkers
- Past studies have revealed that PPAR α and CAR are targets of the legacy PFAS

- Rooney et al., (2018) *Tox Appl Pharm* 356:99–113
- Corton et al. (2020). *Tox Sci.* 177(1):11-26
- Lewis et al. (2020). *Toxicology.* 443:152547



Compiled Studies Examined

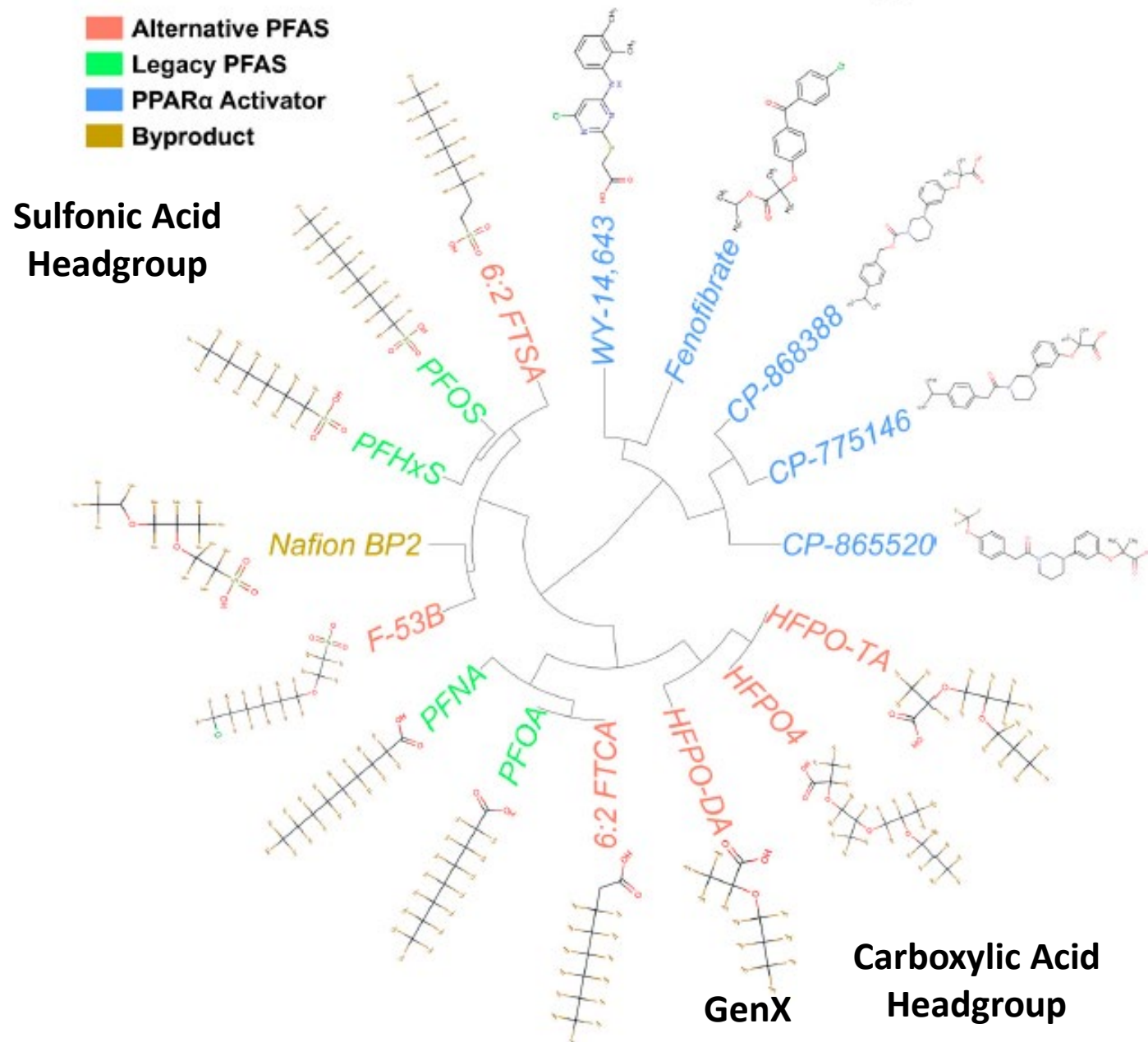
- Studies in the public domain
- Two main profiling platforms (Affymetrix microarrays, RNA-Seq)
- All profiles from male mouse liver
- Exposures were 7-28d
- One to four doses
- Examined 11 PFAS and 5 PPAR α activators

Chemical Name	Abbreviation	DTXSID	PMID	Dose	Timepoint
Perfluorooctane sulfonate	PFOS	DTXSID3031864	20936131	Daily 3 mg/kg or 10 mg/kg	7 days
Perfluorooctanoic acid	PFOA	DTXSID8031865	18281256	Daily 3 mg/kg	7 days
Perfluorononanoic acid	PFNA	DTXSID8031863	28558994	Daily 1 mg/kg or 3 mg/kg	7 days
perfluorohexane sulfonate	PFHxS	DTXSID7040150	28558994	Daily 3 mg/kg or 10 mg/kg	7 days
Perfluoro-2-[(perfluoro-3-(perfluoroethoxy)-2-propanyl)oxy]ethanesulfonic acid	Nafion BP2	DTXSID10892352	#	Daily 0.03, 0.3, 3, or 6 mg/kg	7 days
Ammonium perfluoro-2-methyl-3-oxahexanoate	HFPO-DA (HFPO2) (GenX)	DTXSID40108559	27553808 32138627	Daily 1 mg/kg Daily 0.1, 0.5, or 5 mg/kg	28 days 90 days
Perfluoro-(2,5,8-trimethyl-3,6,9-trioxadodecanoic)acid	HFPO4	DTXSID70276659	27553808	Daily 1 mg/kg	28 days
Perfluoro-2,5-dimethyl-3,6-dioxanonanoic acid	HFPO-TA	DTXSID00892442	29927593	Daily 0.02, 0.1, 0.5 mg/kg	28 days
Potassium 9-chlorohexadecafluoro-3-oxanonane-1-sulfonate	F-53B	DTXSID60881236	#	Daily 5 mg/kg	28 days
6:2 Fluorotelomer sulfonic acid	6:2 FTSA	DTXSID6067331	28032147	Daily 5 mg/kg	28 days
2-Perfluorohexyl ethanoic acid	6:2 FTCA	DTXSID50472556	28032147	Daily 5 mg/kg	28 days
(S)-2-methyl-2-[3-(1-(2-(4-(trifluoromethoxy)phenyl)acetyl)piperidin-3-yl)phenoxy]propanoic acid sodium salt	CP-865520	DTXSID4044032	18971326	Daily 1 mg/kg	5 days
(S)-2-[3-(1-(2-(4-isopropylphenyl)acetyl)piperidin-3-yl)phenoxy]-2-methylpropanoic acid sodium salt	CP-775146	DTXSID9044033	18971326	Daily 1 mg/kg	5 days
(S)-2-[3-(1-((4-isopropylbenzyloxy)carbonyl)piperidin-3-yl)phenoxy]-2-methylpropanoic acid sodium salt	CP-868388	DTXSID4044034	18971326	Daily 1 mg/kg	5 days
Propan-2-yl 2-[4-(4-chlorobenzoyl)phenoxy]-2-methylpropanoate	Fenofibrate	DTXSID2029874	18301758	Single 4 mg/mL	6 h
[[4-Chloro-6-(2,3-dimethylanilino)pyrimidin-2-yl]sulfonyl]acetic acid	WY-14,643	DTXSID4020290	26215100	Single 250 mg/kg	8 h

Structural similarity of the compounds examined in the study

- Used ToxPrint to determine structural relatedness
- PPAR α agonists clustered together
- PFAS cluster based on the head groups

Structure Similarity

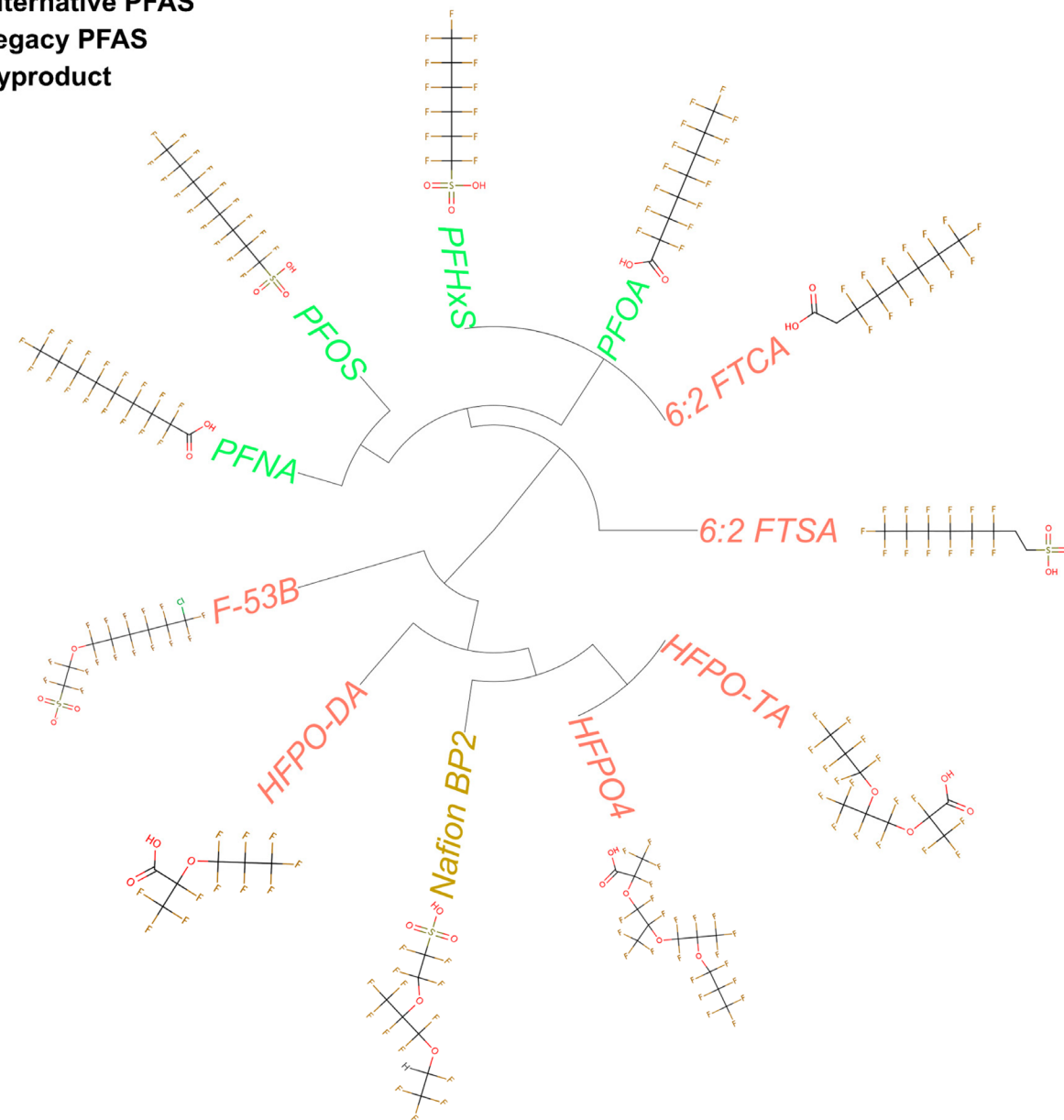


Structural similarity of the compounds examined in the study

- Used ToxPrint to determine structural relatedness
- Removed the head groups
- Clustered based on whether straight chain or contains ether linkages

Structure Similarity

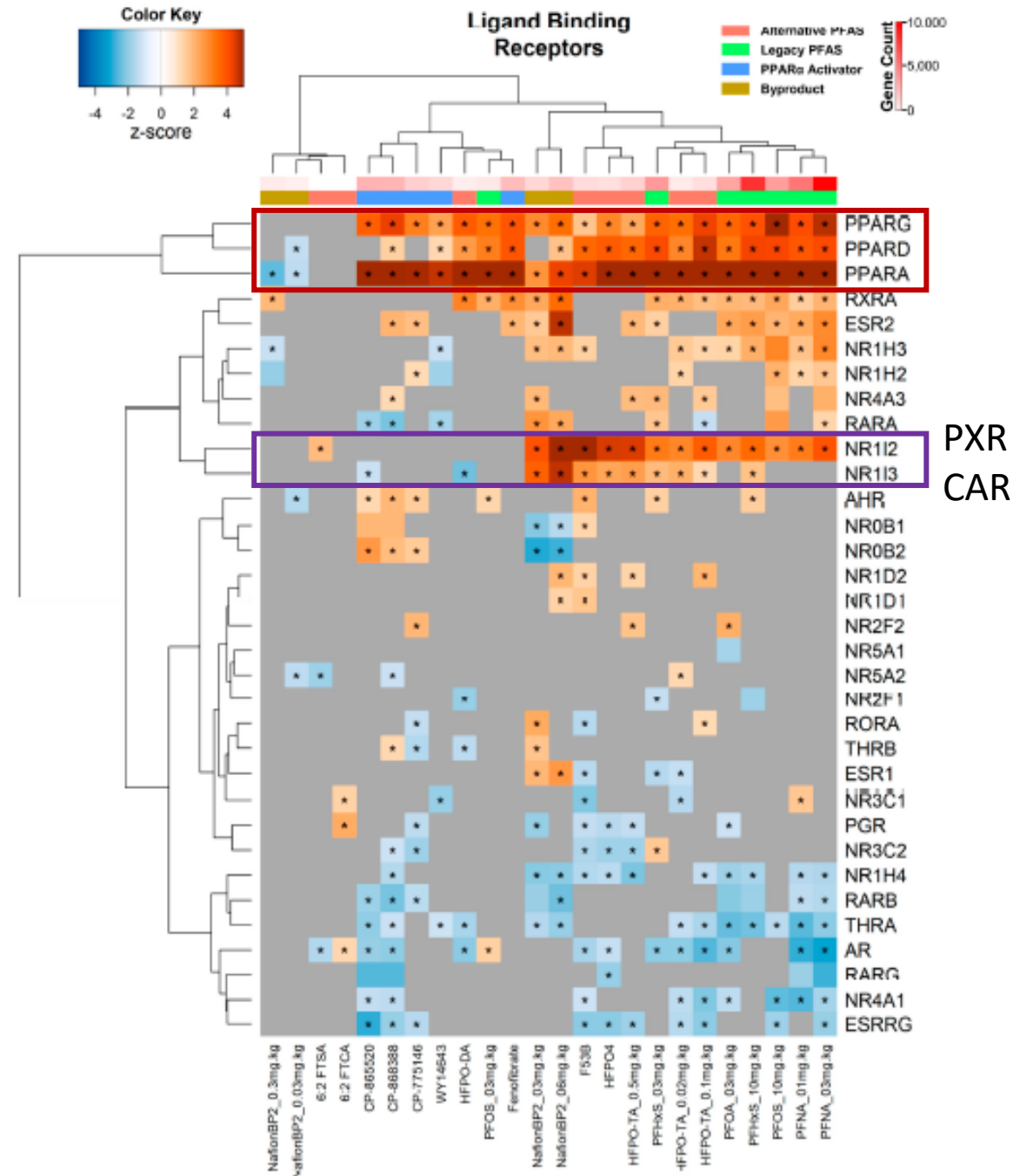
Alternative PFAS
Legacy PFAS
Byproduct



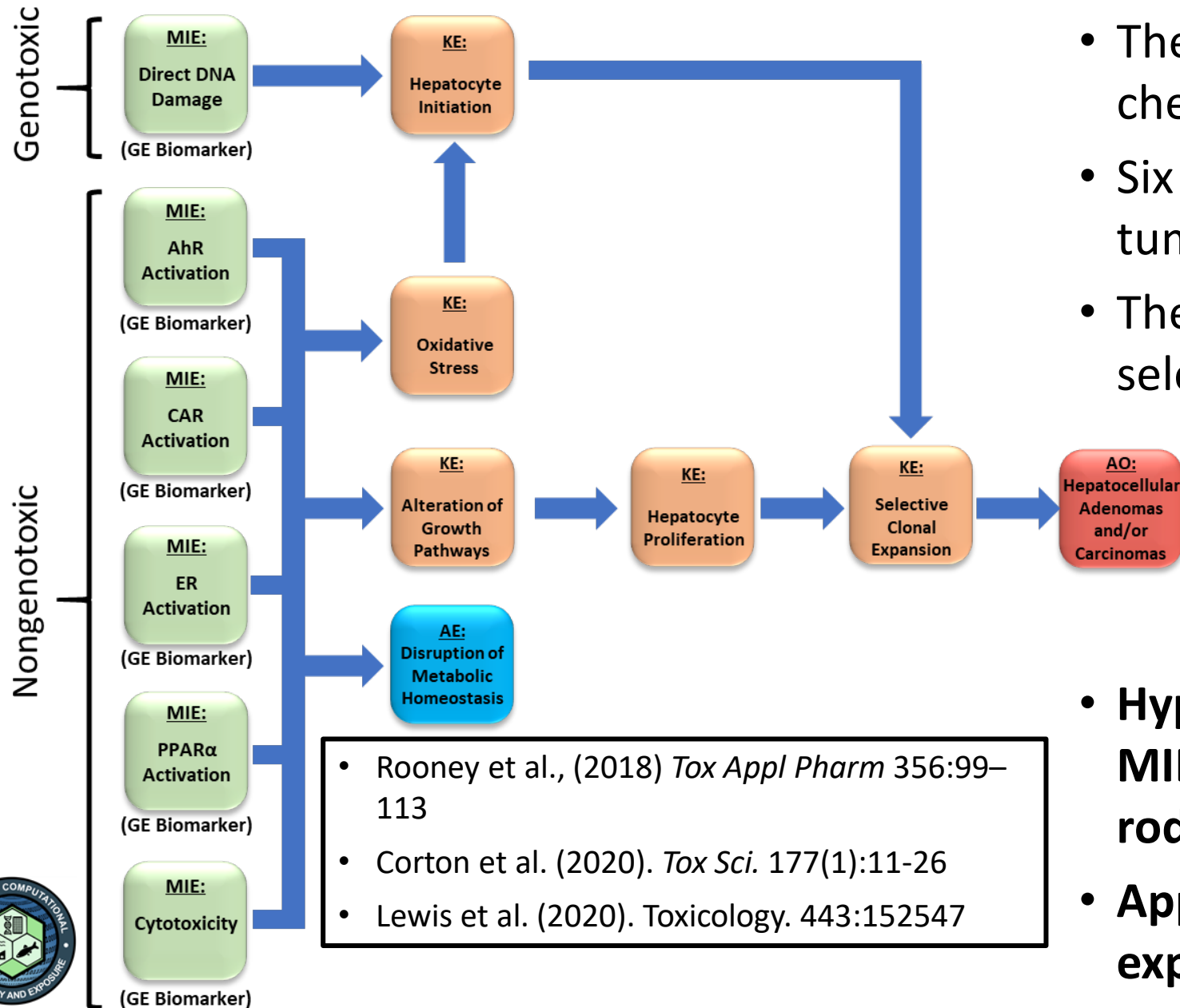
(B)

Examination of the role of nuclear receptors

- Use the upstream activator analysis function of Ingenuity Pathways Analysis (IPA)
- Most PFAS activate PPAR family members
- A subset of PFAS activate CAR and/or PXR
- No major distinctions between legacy and alternative PFAS
- Qualifier: the gene lists in IPA have not been characterized for prediction – they are hypothesis generating tools



Major Adverse Outcome Pathways That Lead to Rodent Liver Tumors



- The liver is the most frequent target of chemical tumorigens
- Six major AOPs lead to rodent liver tumors
- The AOPs converge on the key event of selective clonal expansion

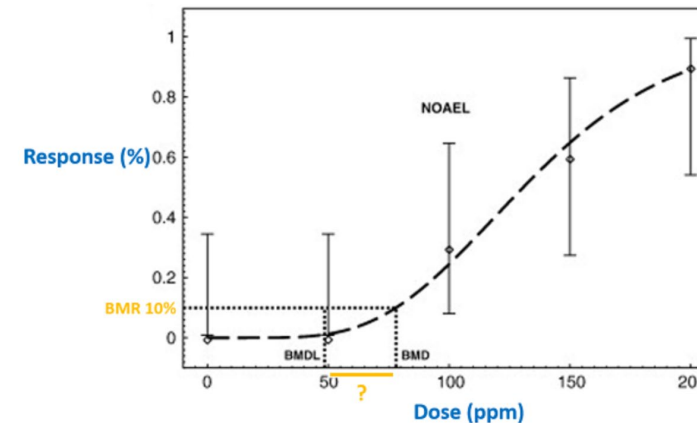
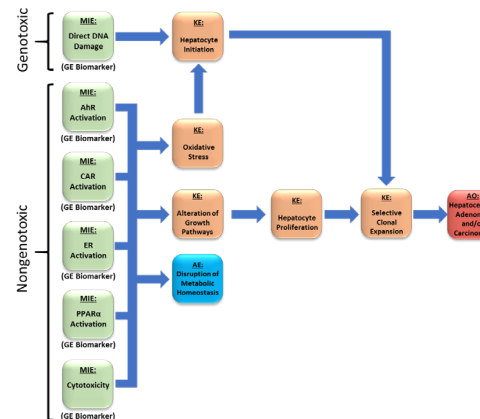
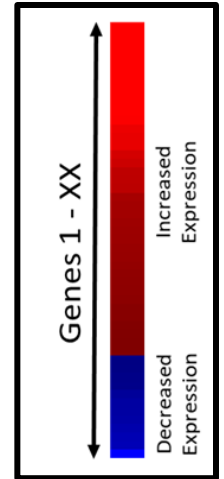
- **Hypothesis: measurement of the six MIEs will be sufficient to predict rodent liver tumors**
- **Approach: measure MIEs with gene expression biomarkers**

- Rooney et al., (2018) *Tox Appl Pharm* 356:99–113
- Corton et al. (2020). *Tox Sci.* 177(1):11-26
- Lewis et al. (2020). *Toxicology.* 443:152547



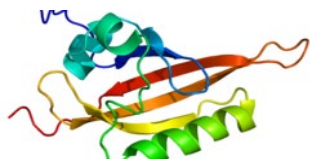
Gene expression biomarkers

- Biomarker is defined as “a characteristic that is objectively measured and evaluated as an indicator of normal biological processes, pathogenic processes, or pharmacologic responses to a therapeutic intervention.” (1998, the National Institutes of Health Biomarkers Definitions Working Group)
- A gene expression biomarker is a short list of genes and associated fold-change values or ranks used to predict the activity of a factor important in mediating effects of chemicals or toxicity
- Can be used to
 - Identify mode of action
 - Predict tumorigenic potential
 - Determine a benchmark dose
- Very few examples of well characterized gene expression biomarkers with known accuracies
 - Gene lists as signature/pathway analysis often used as hypothesis generators

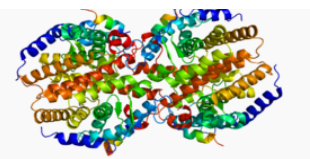


Biomarkers that predict key events in the livers of mice and rats

AhR



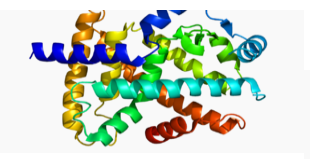
CAR



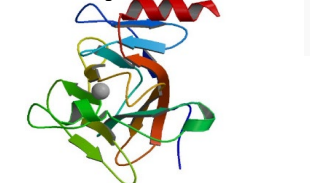
NRF2



PPAR α



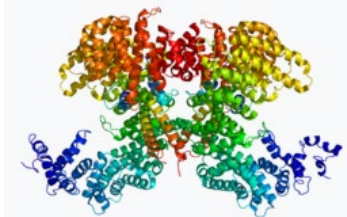
p53



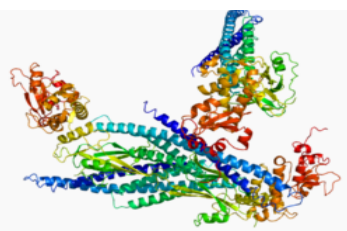
Estrogen
Receptor α



SREBP



STAT5b



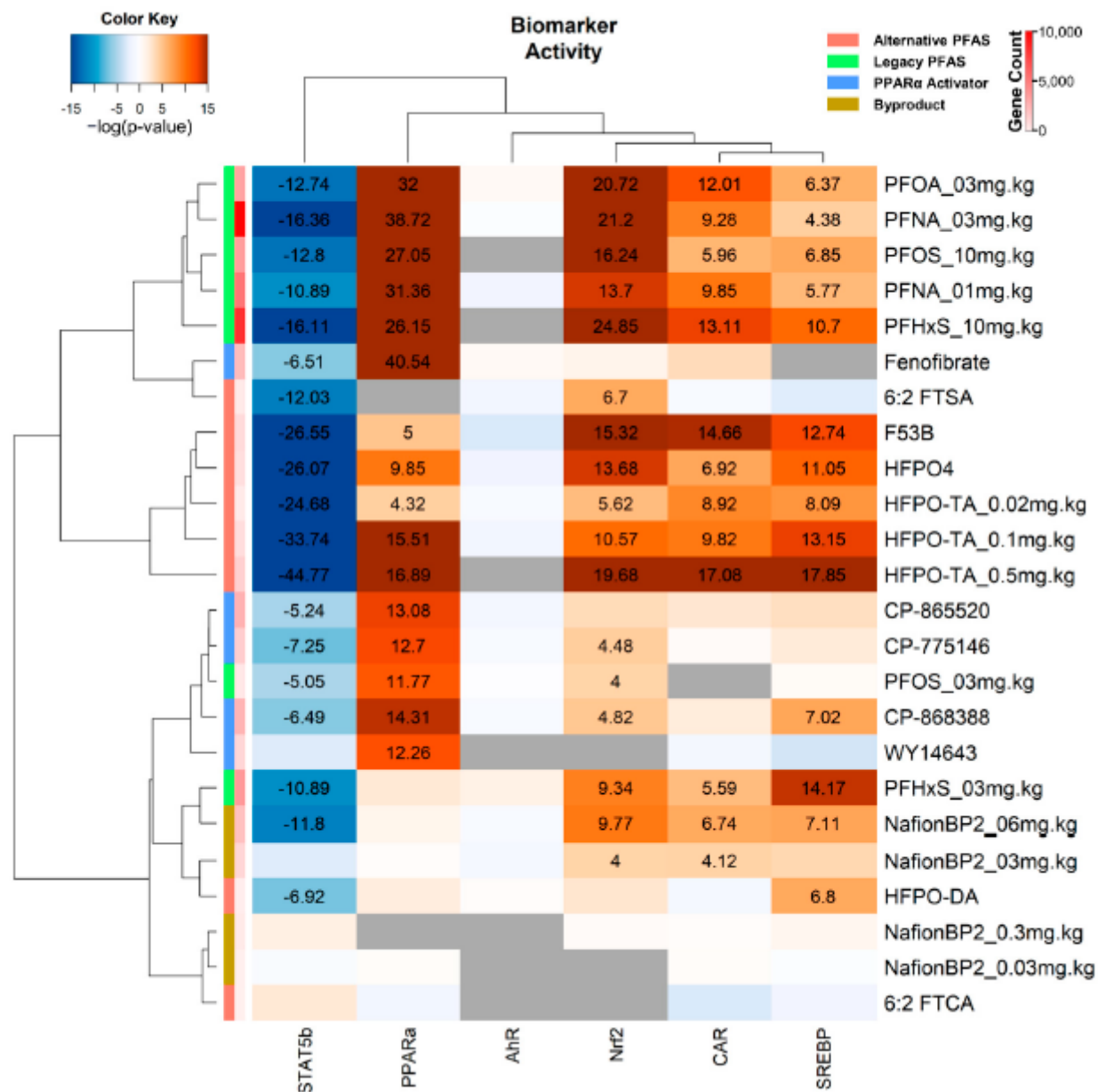
- Oshida et al. (2015). Identification of Modulators of the Nuclear Receptor Peroxisome Proliferator-Activated Receptor α (PPAR α) in a Mouse Liver Gene Expression Compendium. PLoS One. 10(2):e0112655.
- Oshida et al. (2015). Identification of Chemical Modulators of the Constitutive Activated Receptor (CAR) in a Mouse Liver Gene Expression Compendium. Nuclear Receptor Signaling. 13:e002.
- Oshida et al. (2015). Screening a Mouse Liver Gene Expression Compendium Identifies Effectors of the Aryl Hydrocarbon Receptor (AhR). Toxicology. 336:99-112.
- Oshida et al. (2015). Disruption of STAT5b-Regulated Sexual Dimorphism of the Liver Transcriptome by Diverse Factors Is a Common Event. PLoS One. 11(3):e0148308.
- Oshida et al. (2015). Chemical and Hormonal Effects on STAT5b-Dependent Sexual Dimorphism of the Liver Transcriptome. PLoS One. 2016 11(3):e0150284.
- Rosen et al. (2017). PPAR α -independent transcriptional targets of perfluoroalkyl acids revealed by transcript profiling. Toxicology. 387:95-107.
- Rooney et al. (2017). Genomic Effects of Androstenedione and Sex-Specific Liver Cancer Susceptibility in Mice. Toxicol Sci. 160(1):15-29.
- Rooney et al. (2018) Activation of Nrf2 in the liver is associated with stress resistance mediated by suppression of the growth hormone-regulated STAT5b transcription factor. PLoS One. 13(8):e0200004.
- Rooney et al. (2018). Activation of CAR leads to activation of the oxidant-induced Nrf2. Toxicol Sci. 167:172-189.
- Rooney et al. (2018). Adverse outcome pathway-driven identification of rat liver tumorigens in short-term assays. Toxicol Appl Pharmacol. 356:99-113.
- Corton (2019). Frequent Modulation of the Sterol Regulatory Element Binding Protein (SREBP) by Chemical Exposure in the Livers of Rats. Comput. Toxicol. 10:113-129.

The mouse biomarkers have excellent predictive accuracy

Mouse Biomarker	Number of Genes	Mutant mice used	Predictive Accuracy for Activation	Publication
PPARα	131	<i>Ppara</i>	98%	PLoS One. 2015 10(2):e0112655
CAR	83	<i>Nr1i3</i>	97%	Nucl Recept Signal. 2015 13:e002
AhR	63	<i>Ahr</i>	95%	Toxicology. 2015 336:99-112
Nrf2	Activated by oxidative stress	<i>Nfe2l2, Keap1</i>	96%	PLoS One 2018 13(8):e0200004
Stat5b	Regulates growth hormone responsive genes		97%	PLoS One 2016 11(3):e0150284
Srebp	Regulates genes involved in the synthesis of cholesterol and triglycerides	<i>Srebp1a, Srebp1b</i>	97%	Genes 10 (2019) 77

Biomarker analysis

- Most PFAS suppress Stat5b – indicative of suppression of growth hormone signalling
- Most chemicals activate PPAR α (not Nafion BP2, 6:2 FTSA and 6:2 FTCA) that regulates fatty acid and glucose metabolism
- Many activate CAR and Nrf2 – indicative of increases in xenobiotic metabolism and associated oxidative stress
- Many activate SREBP – linked to steatosis commonly observed in the livers of treated rodents
- No clear distinctions between the legacy and alternative PFAS
- 6:2 FTCA and 6:2 FTSA activate fewer factors - 6:2 FTCA is more quickly metabolized



Conclusions

- To understand the diversity of molecular targets of the PFAS in the mouse liver, we performed a comparative toxicogenomics analysis of the gene expression changes after exposure to 11 PFAS
- Using hierarchical clustering, pathway analysis, and predictive biomarkers, we found that most of the alternative PFAS modulate molecular targets that overlap with legacy PFAS
- Only three of the 11 PFAS tested did not appreciably activate PPAR α (Nafion BP2, 6:2 FTSA, and 6:2 FTCA)
- Predictive biomarkers showed that most PFAS (PFHxS, PFOA, PFOS, PFNA, HFPO-TA, F-53B, HFPO4, Nafion BP2) activated CAR
- PFNA, PFHxS, PFOA, PFOS, HFPO4, HFPO-TA, F-53B, Nafion BP2, and 6:2 FTSA activated NRF2
- A subset of PFAS activated SREBP that may underlie the steatosis observed
- The work highlights the similarities in molecular targets between the legacy and alternative PFAS
- We predict that the alternative PFAS (except 6:2 FTCA) would be no less toxic than the legacy PFAS

Thanks for listening!

Questions?