

# Unraveling the role of AHR gene duplication in PAH toxicity in zebrafish

ROBERT TANGUAY

OREGON STATE UNIVERSITY

DEPARTMENT OF ENVIRONMENTAL AND MOLECULAR TOXICOLOGY

SINNHUBER AQUATIC RESEARCH LABORATORY

ENVIRONMENTAL HEALTH SCIENCES CENTER

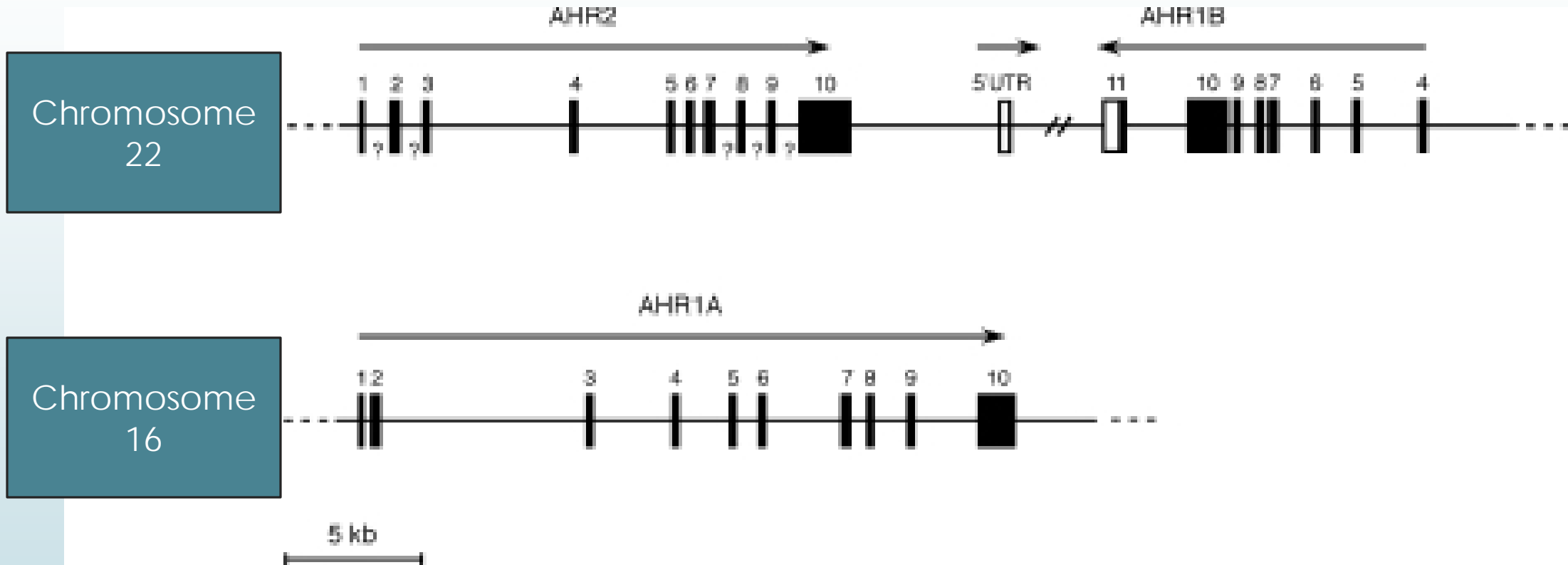


# Outline

- A little history on zebrafish AHR genes
- PAH toxicity
- Classifying PAHs
- AHR Downstream events
- AHR regulated non coding RNAs

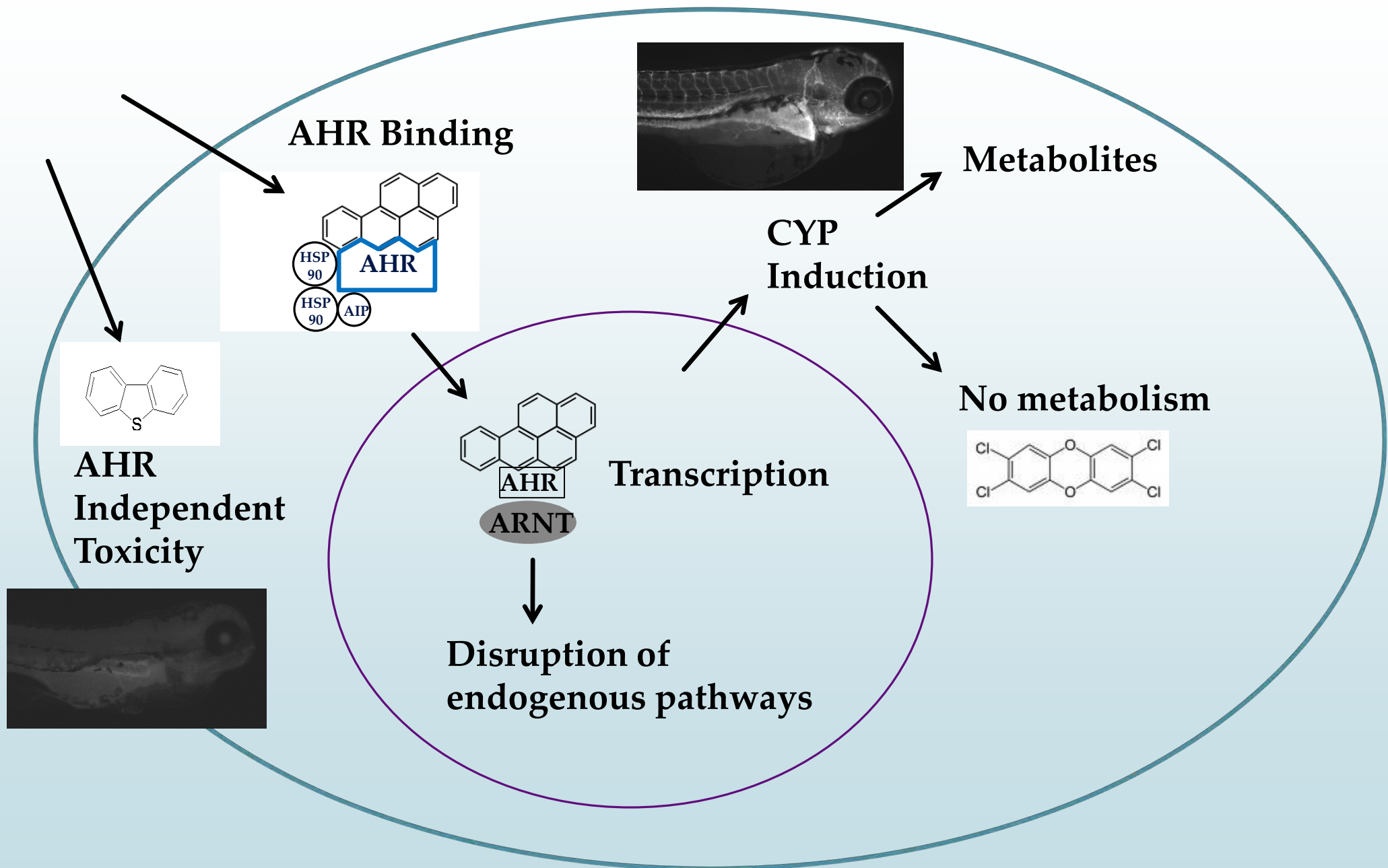


# AHR Gene Duplication



AHR1B, a new functional aryl hydrocarbon receptor in zebrafish: tandem arrangement of *ahr1b* and *ahr2* genes  
Sibel I. Karchner, Diana G. Franks, Mark E. Hahn, *Biochem J.* 2005 Nov 15; 392(Pt 1): 153–161.

# The AHR and PAH pathways of toxicity



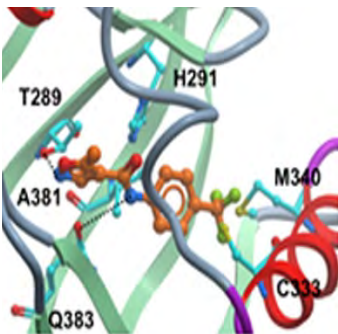
# Predicting AHR Ligands

# Modeling a “Target” Zebrafish AHRs



## Zebrafish three AHRs

- AHR2 primary mediator of toxicity
- AHR1A deficient in TCDD binding and transactivation activity
- AHR1B functional but no known toxicological roles

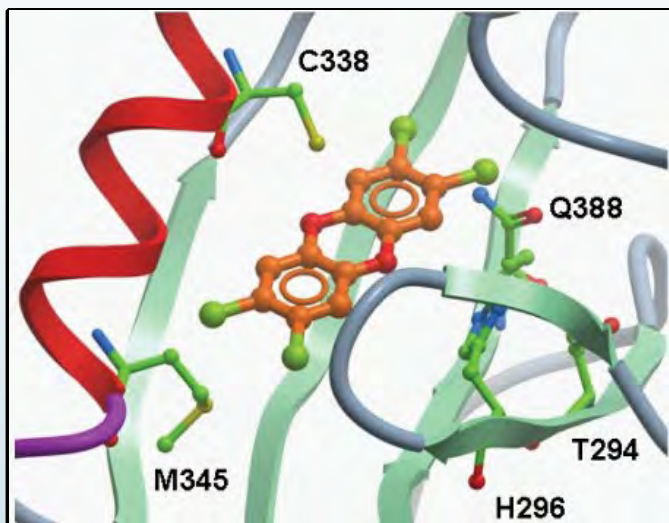


## AHR Homology Model

- AHR ligand binding domain models built using NMR structure of HIF2 $\alpha$  (PAS domain)
- Mouse, rat, human, zebrafish
- Performed molecular docking of putative AHR ligands

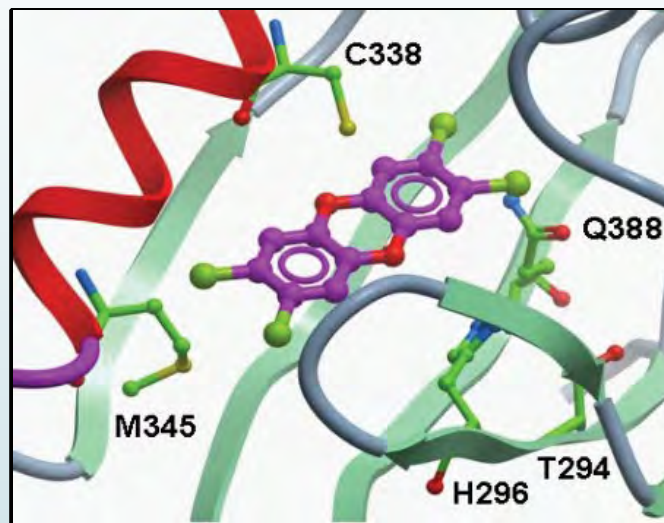
# TCDD Molecular Docking with the Zebrafish AHRs

AHR2



-3.97

AHR1B



-4.86

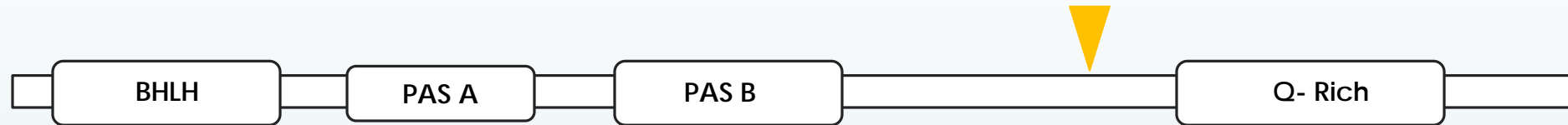
AHR1A

Unable to dock

Predicted binding energy  
(kcal/mole)

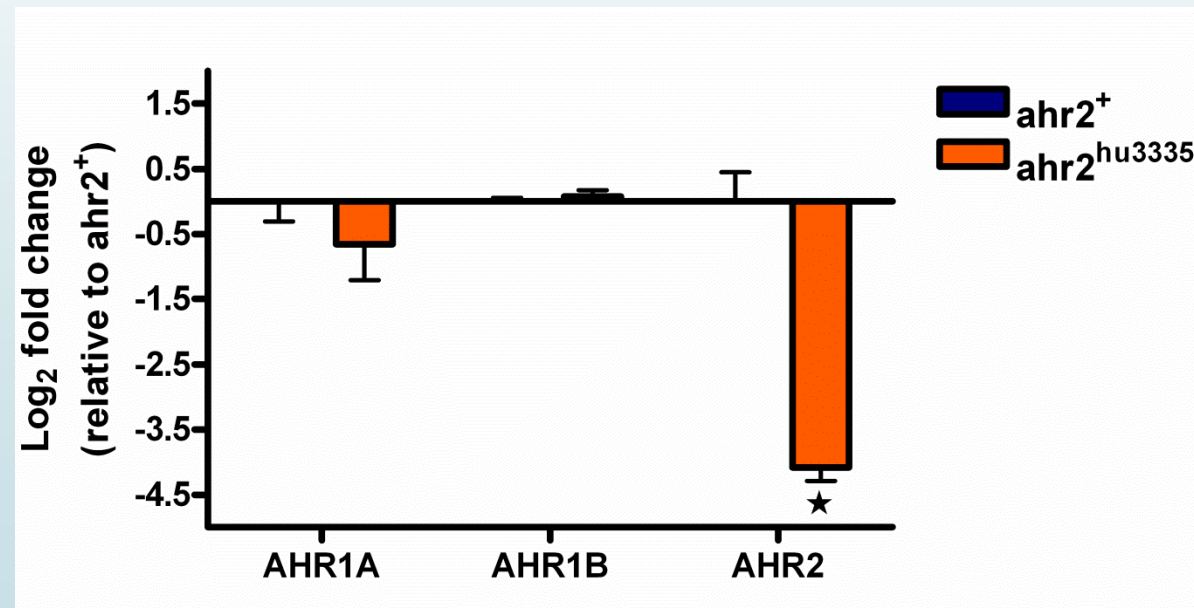
# The *ahr2*<sup>hu3335</sup> Zebrafish Line

T → A mutation in residue 534 resulting in a premature stop

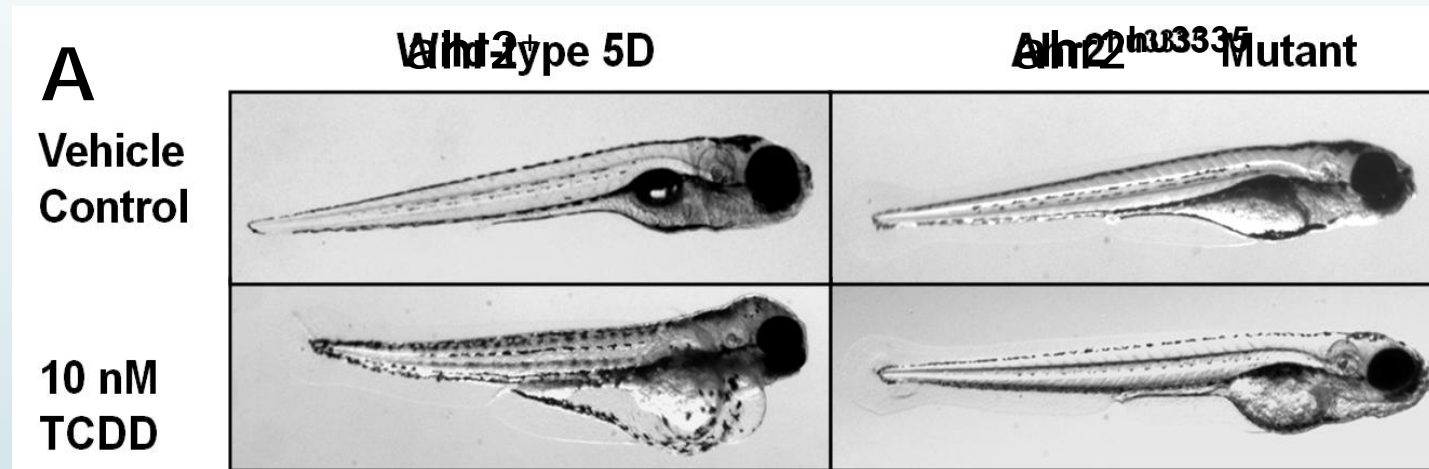


- Truncated protein is predicted to be non-functional

- Basal mRNA expression suggests mutant *ahr2*<sup>hu3335</sup> transcript is degraded



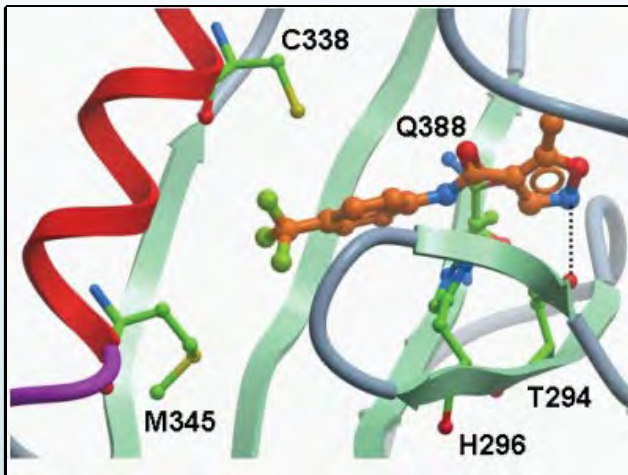
# *Ahr2*<sup>hu3335</sup> Mutants Are Resistant to TCDD-Induced Developmental Toxicity





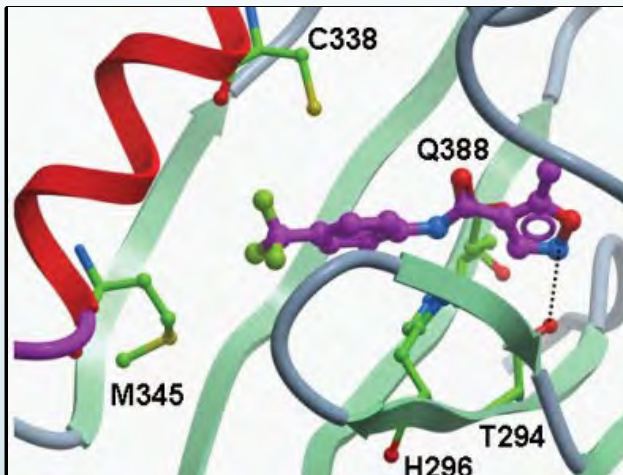
# Leflunomide Molecular Docking

AHR2



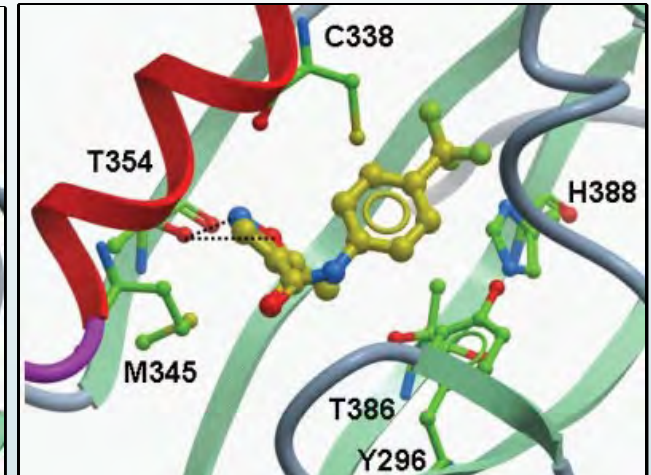
-2.13

AHR1B



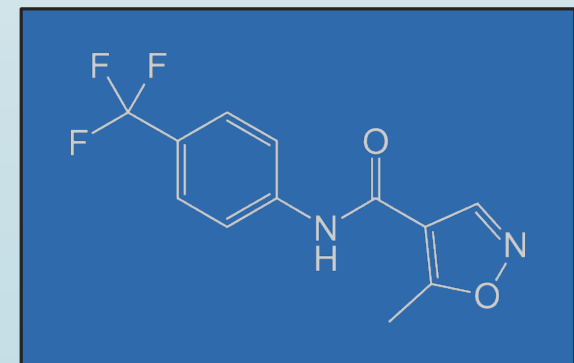
-1.97

AHR1A



-2.19

Predicted binding energy  
(kcal/mole)





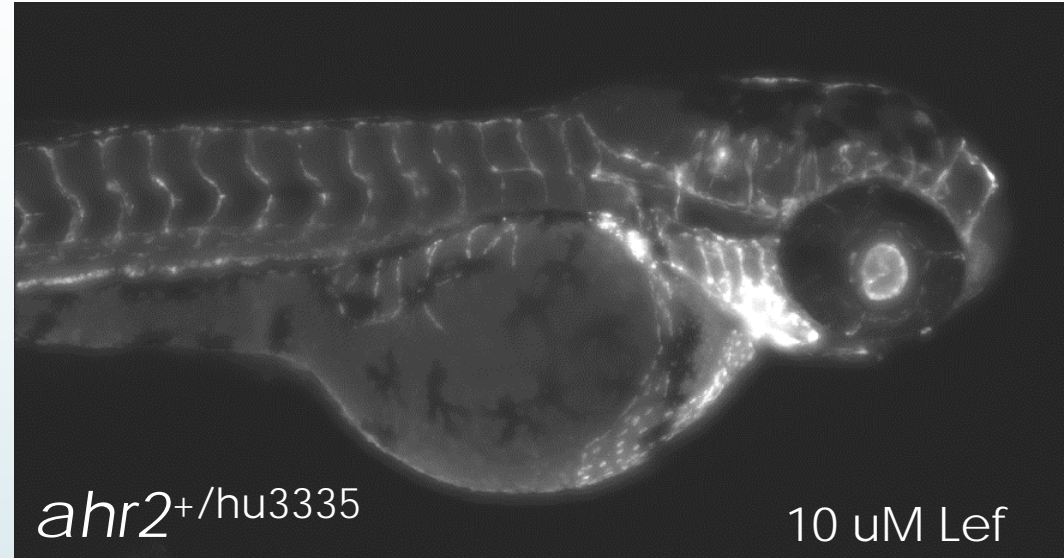
# Leflunomide-induced CYP1A expression is partially AHR2 dependent

AHR Status

Expressed

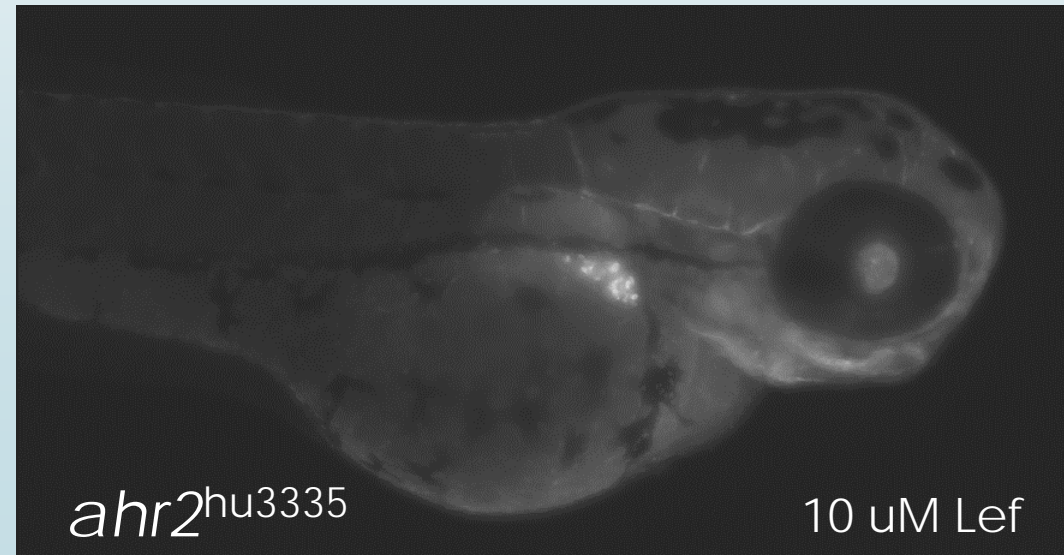


Not Expressed



*ahr2*<sup>+/hu3335</sup>

10 uM Lef

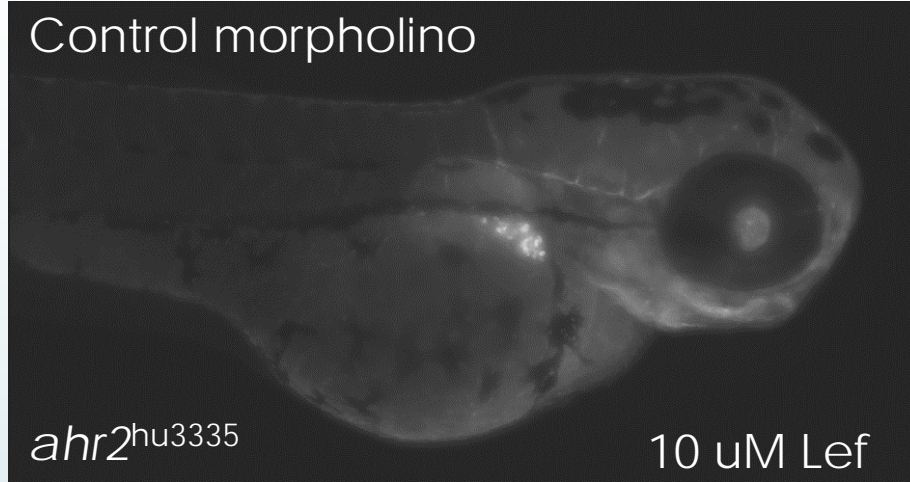


*ahr2*<sup>hu3335</sup>

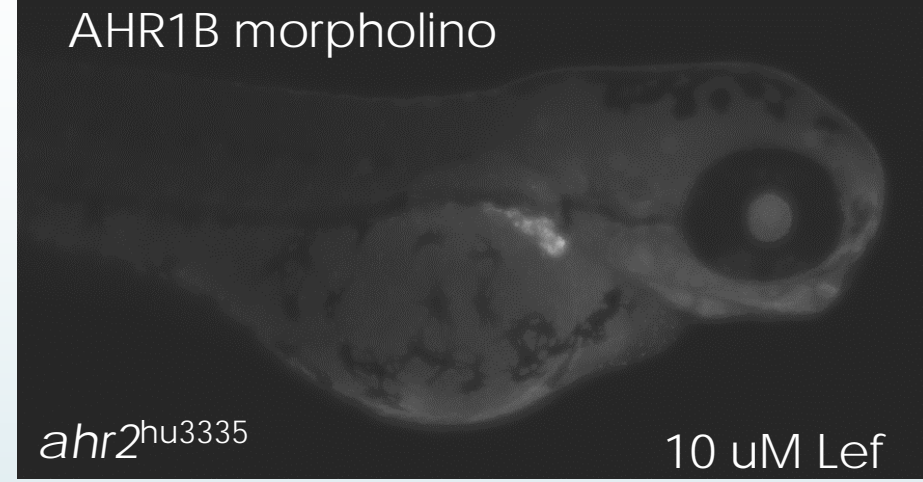
10 uM Lef

# AHR1A Dependent CYP1A Expression

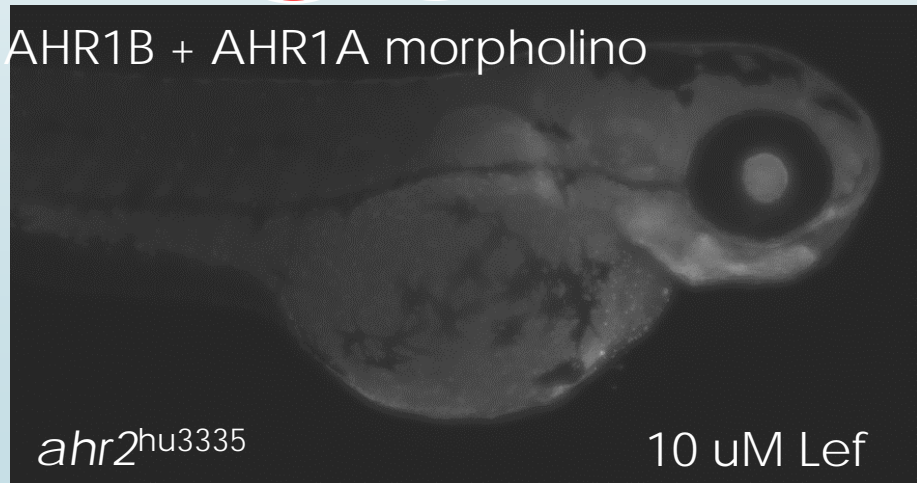
1a 1b 2



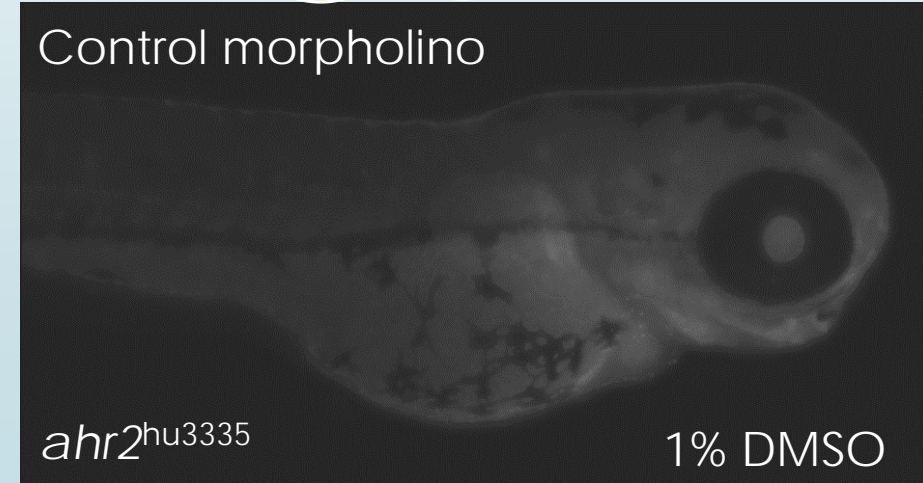
1a 1b 2



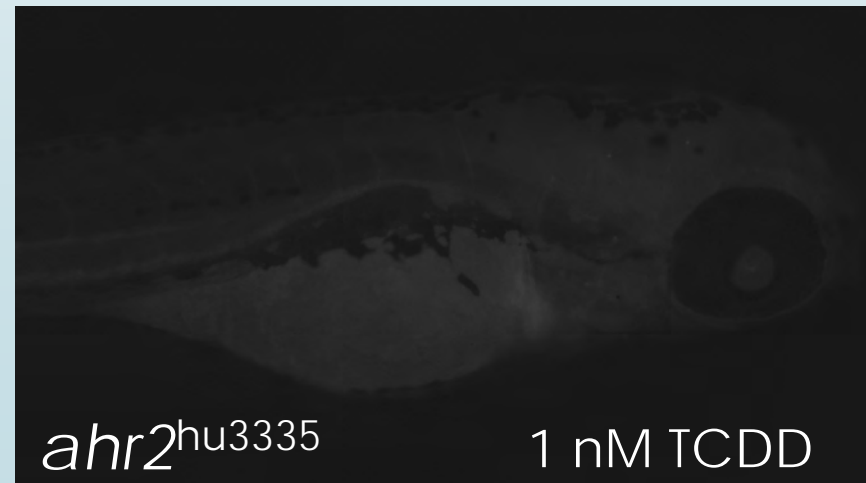
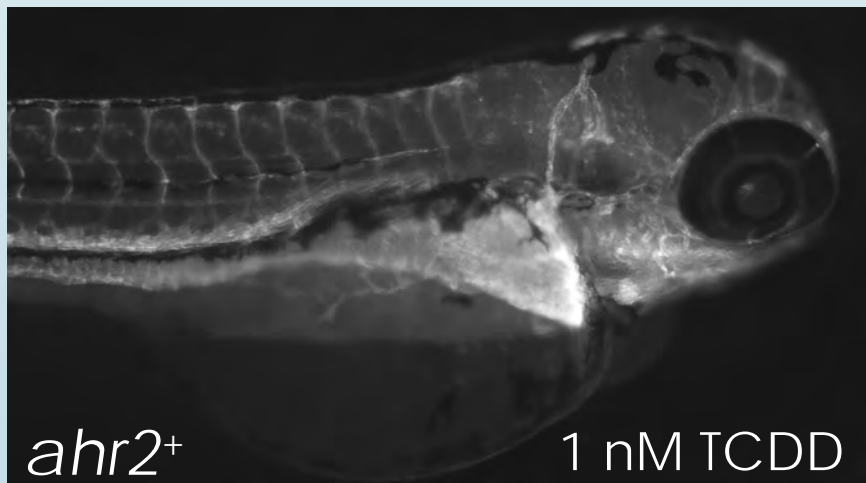
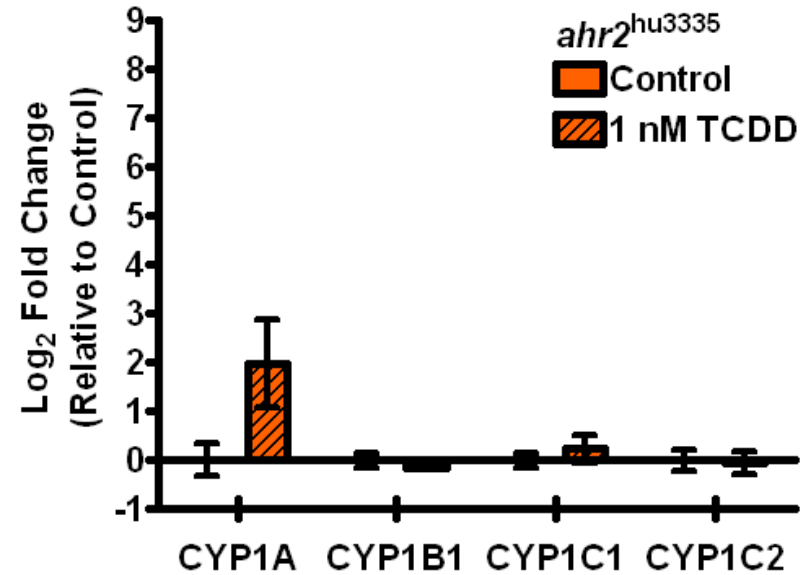
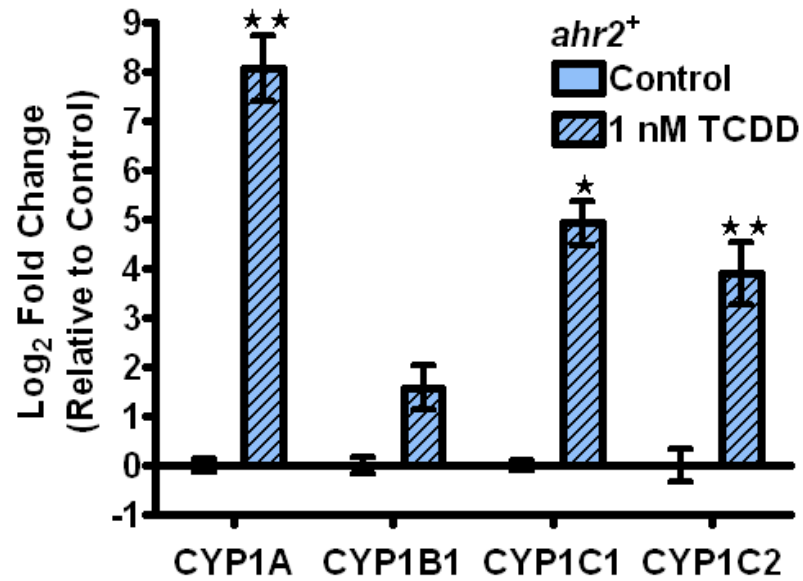
1a 1b 2



1a 1b 2



# *ahr2* Mutants Are Resistant to TCDD-induced CYP Expression Changes





# AHR2 importance confirmed in CRISPR/Cas9 line

A

-11 bp deletion in exon 1

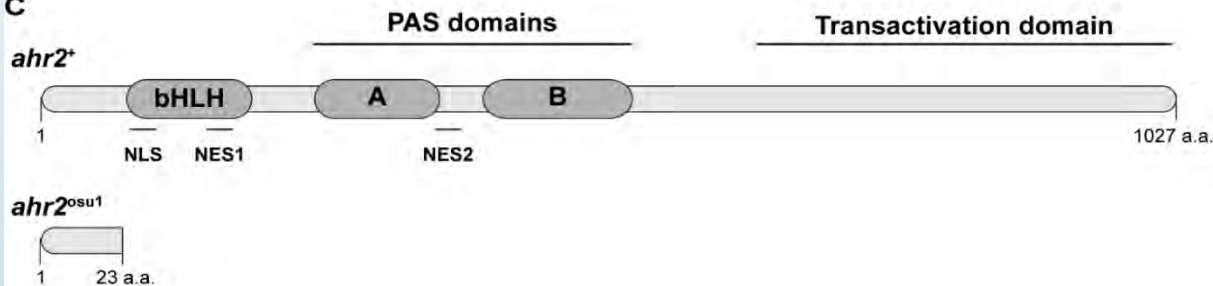
<i>ahr2</i> <sup>+</sup>	ATG TCG GCG GGT ATC GGT ACA TAT GCG GTC AAG AAA CGG AAG
<i>ahr2</i> <sup>osu1</sup>	ATG TCG GCG GGT ATC GGT.....C AAG AAA CGG AAG (-11 bp)

B

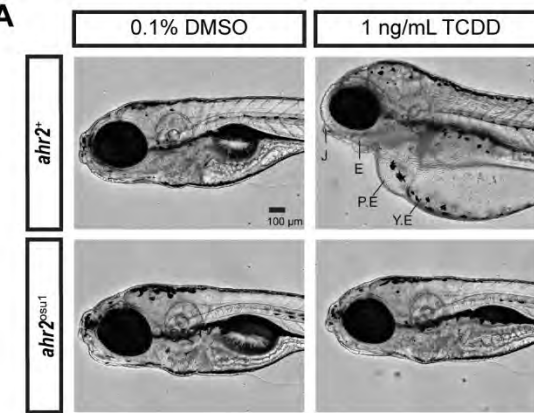
Translation of mutant sequence predicts premature stop codon

1	ATG TCG GCG GGT ATC GGT CAA GAA ACG GAA GAA GCC CGT TCA GAA	45
1	Met Ser Ala Gly Ile Gly Gln Glu Thr Glu Glu Ala Arg Ser Glu	15
46	AAT ACC CAA ACC ACC ACC CCC TGA	69
16	Asn Thr Gln Thr Thr Thr Pro End	23

C



A



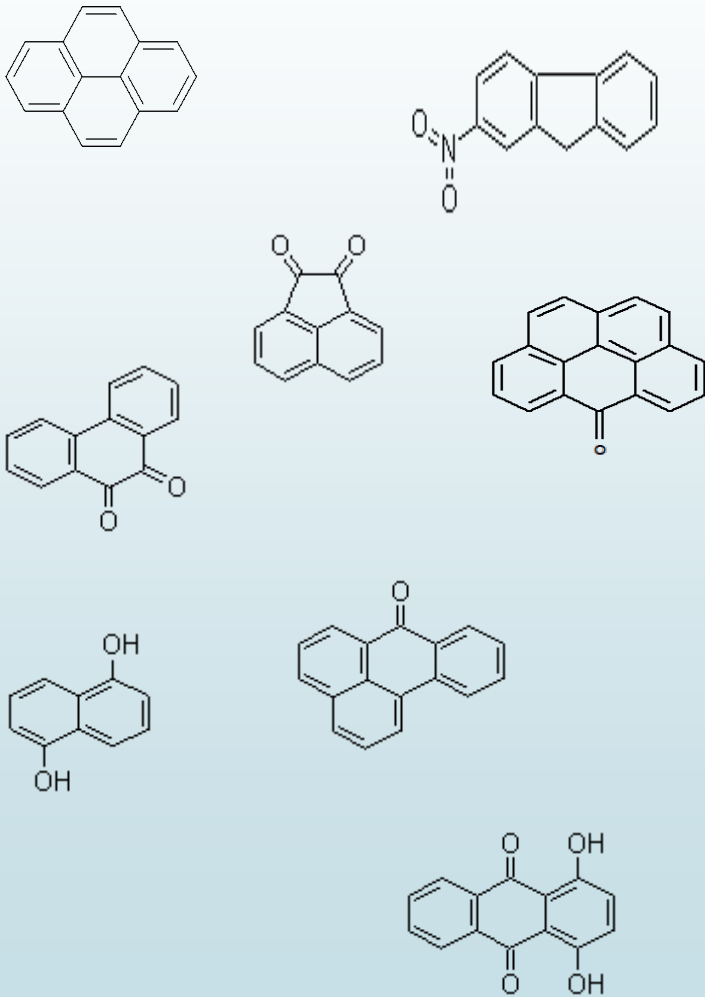
B



Garcia GR, Bugel SM, Truong L, Spagnoli S, Tanguay RL. AHR2 required for normal behavioral responses and proper development of the skeletal and reproductive systems in zebrafish. *PLoS one*. 2018;13(3)

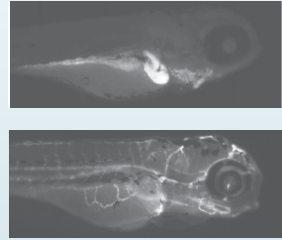
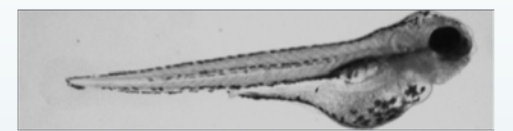
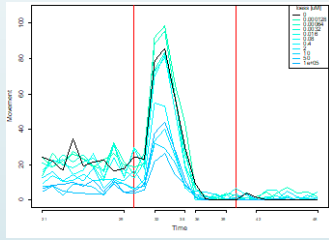
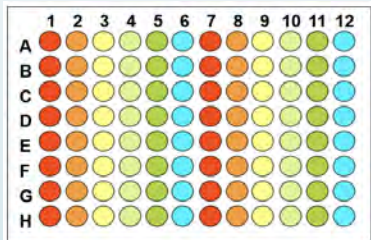
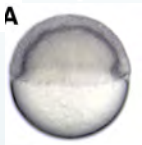
**OSU/PNNL SUPERFUND RESEARCH PROGRAM**  
**POLYCYCLIC AROMATIC HYDROCARBON MIXTURES: NEW**  
**TECHNOLOGIES AND EMERGING HEALTH RISKS**

# Toxicity Mechanisms for Most PAHs are Unknown



- Environmental samples can contain 100's PAHs
- Environmentally Dynamic
- Parent, substituted compounds
- Toxicity data is scarce for substituted PAHs
- PAHs induce AHR-dependent and AHR-independent developmental toxicity, dependent on structure
- We lack the structural basis for developmental and neurotoxicity

# High-Throughput Screening of PAHs

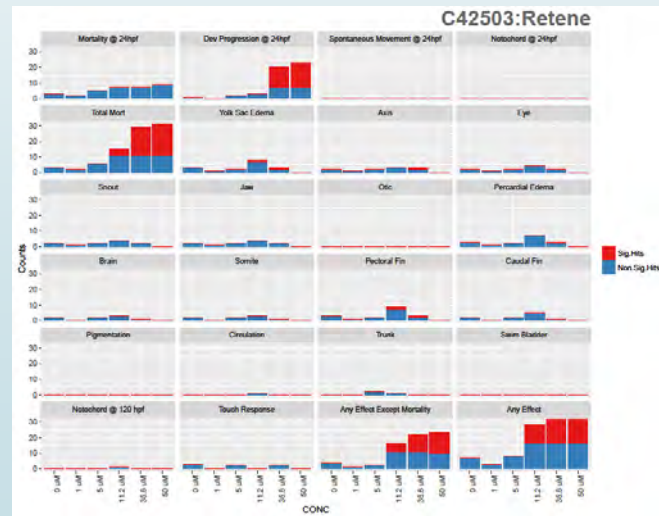


## Exposure

- Dechorionated
- 6 hpf to 120 hpf
- 5 concentrations
  - 50-1  $\mu\text{M}$
  - 5-0.1  $\mu\text{M}$
- N=32

## 24 hour evaluations

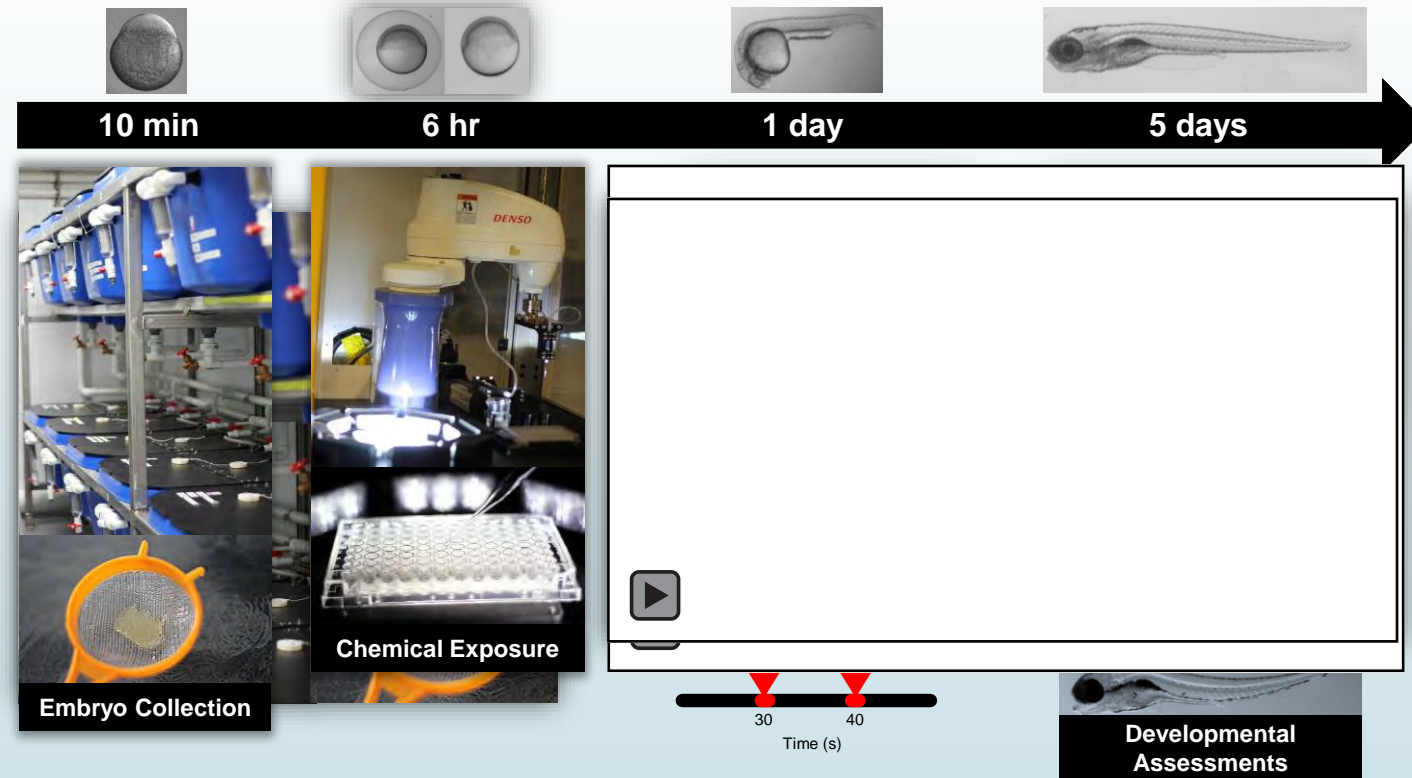
- Behavior
- Morphology



## 5 day evaluations

- Morphology
- Behavior
- CYP1A Localization

# HTS Platform for Chemical Screening



Truong et al. (2014) *Toxicol Sci* 137: 212-233.

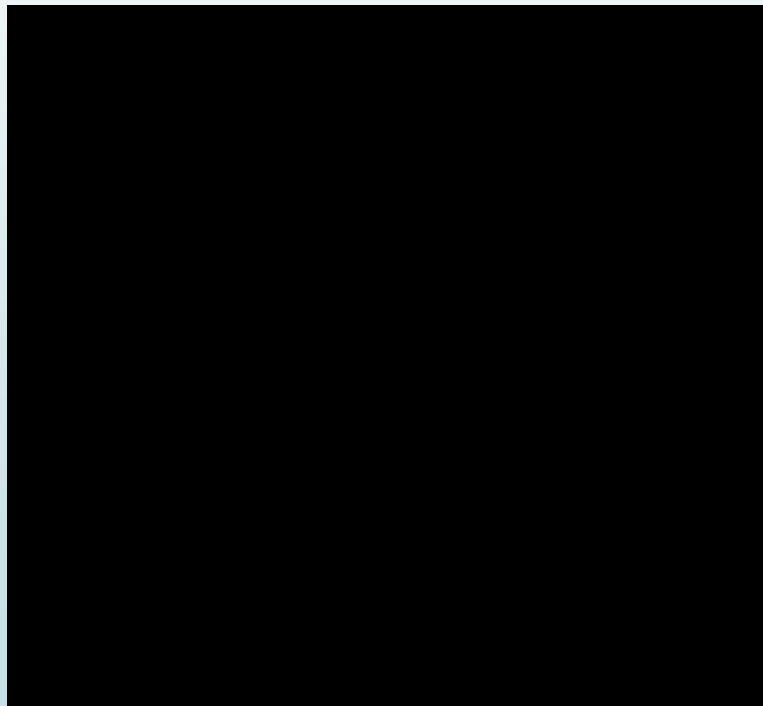
Mandrell, D., **Truong, L.**, et al. 2012. Automated zebrafish chorion removal and single embryo placement: Optimizing throughput of zebrafish developmental toxicity screens. *Journal of Laboratory Automation* 17 (1) 66-74.

All dependent on a custom made LIM system "ZAAP"

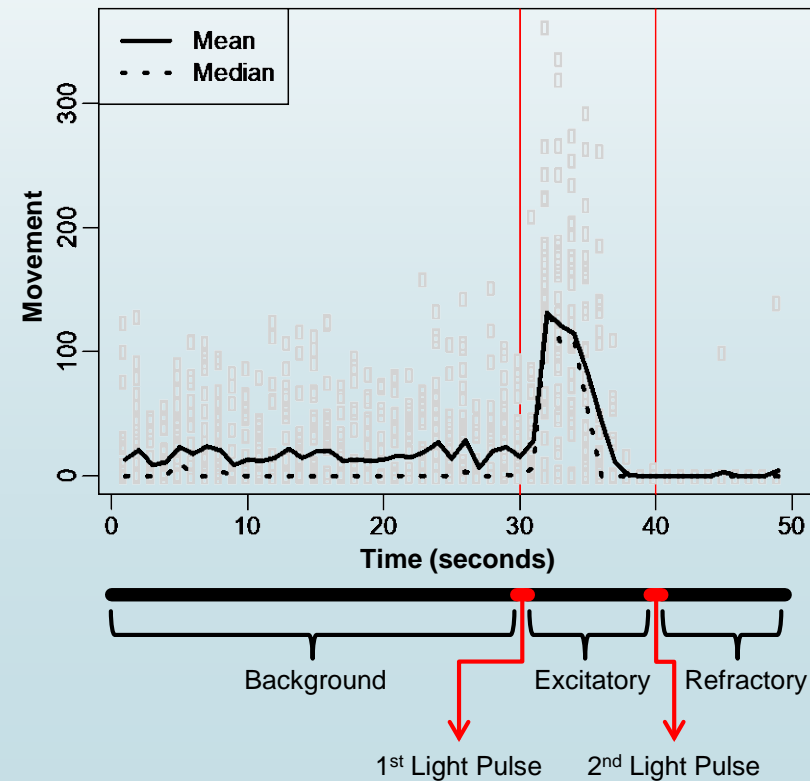


# What About The 24 hpf Photomotor Response (EPR)?

- Control fish (in the absence of chemical) will respond after the 1<sup>st</sup> light pulse (**Excitatory** interval) but not after the 2<sup>nd</sup> light pulse (**Refractory** interval).



EPR recording (50 seconds)

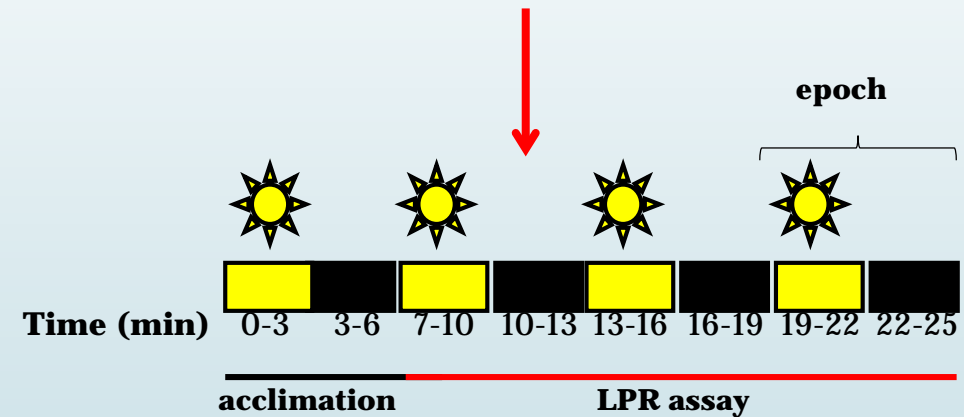


# Larval Photomotor Response (LPR) Behavioral Testing (5 day Larvae)

Distance Moved During Alternating Periods of Light and Dark



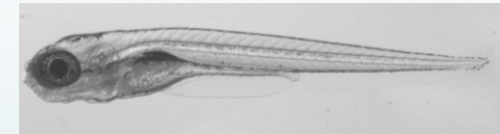
## Larval Photomotor Behavior



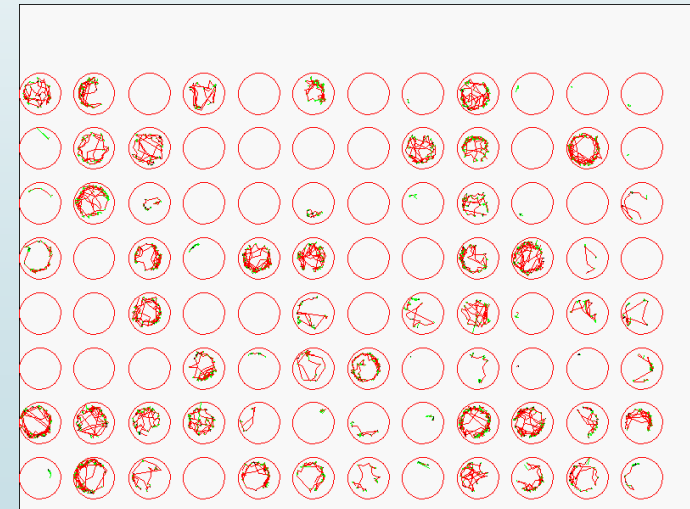
The LPR assay is 18 mins, with a 6 min acclimation period. Statistical significant was determined using a K-S test with a  $p < 0.01$ , and a minimum of 30% AUC change (relative to control).

# Larval Photomotor Response (LPR) Behavioral Testing (5 day Larvae)

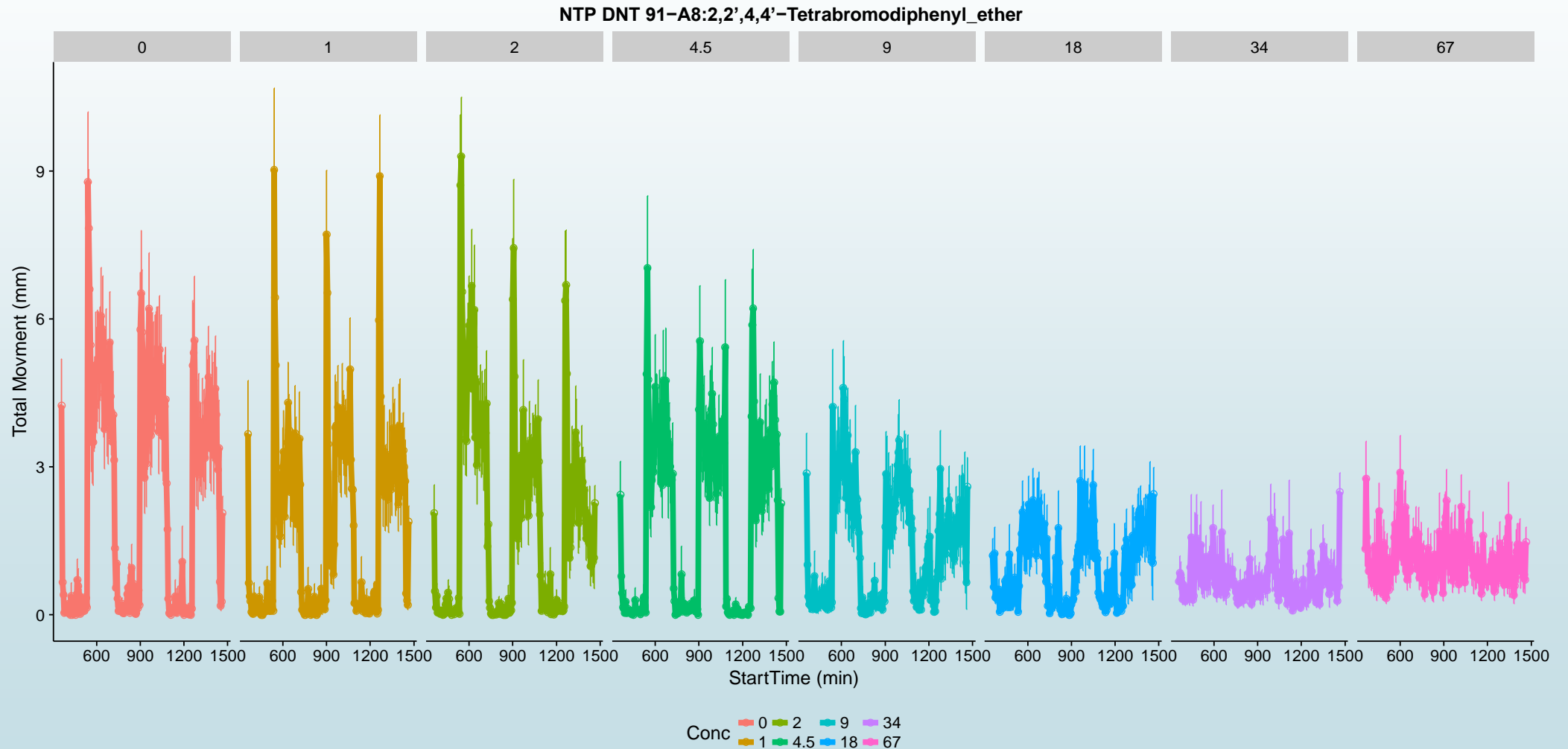
96-well plate



Distance moved over time



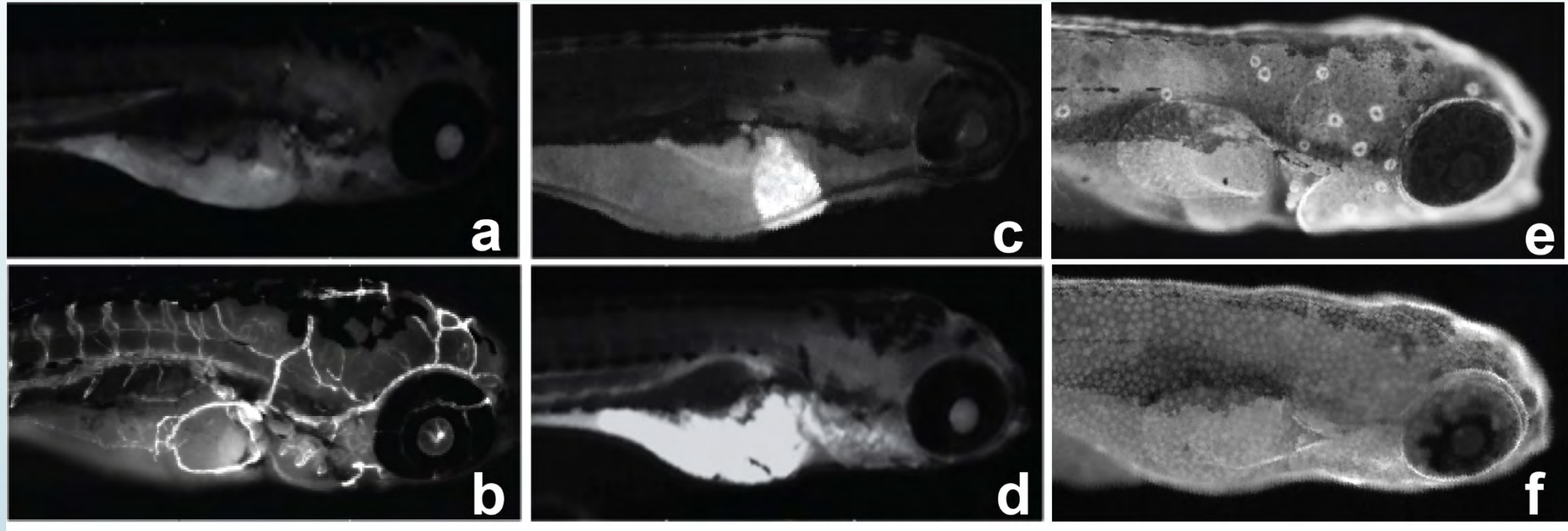
# Larval Photomotor Response (LPR) Example Output (5 day Larvae)



# Comparative PAH Screening Effort

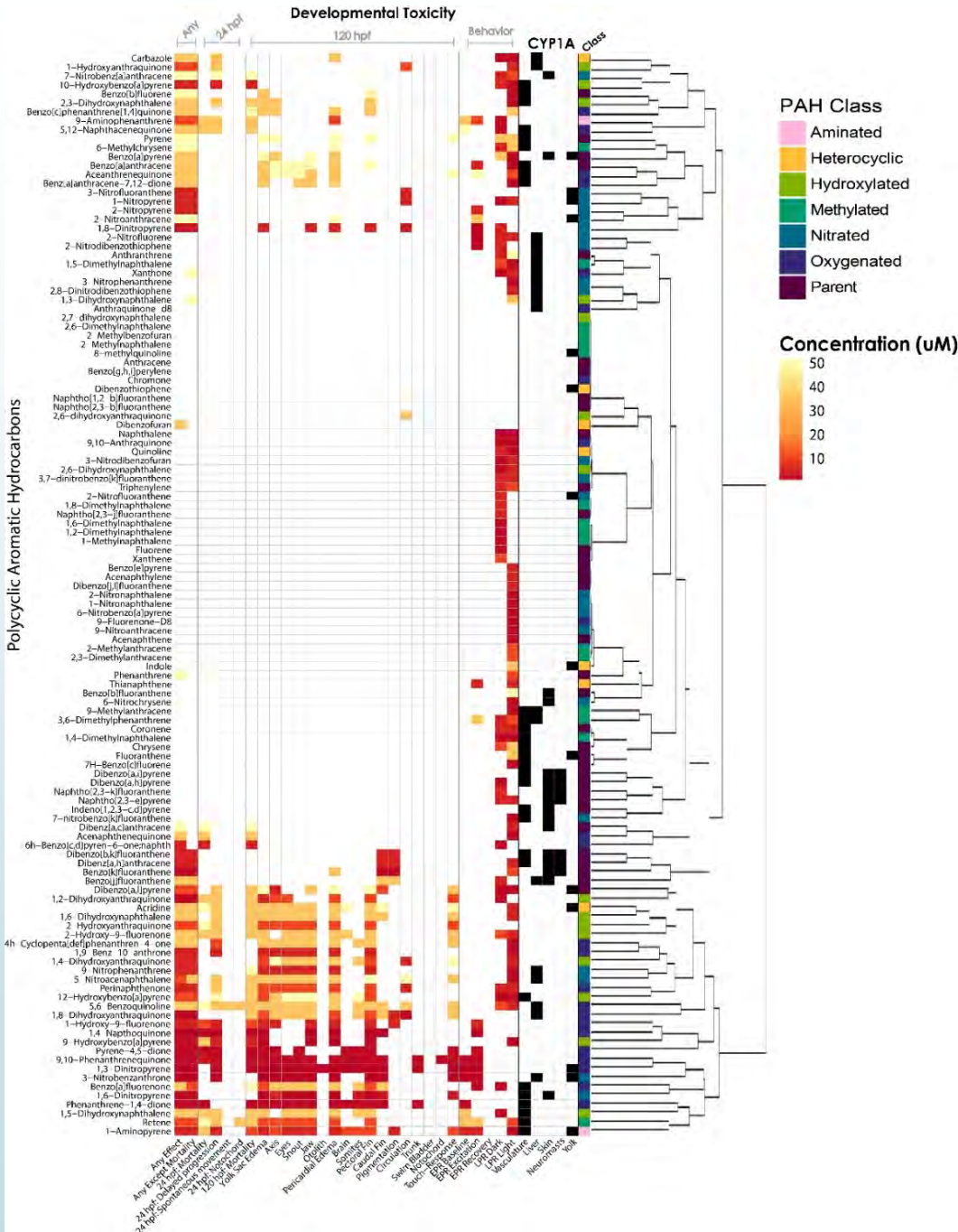


# CYP1A Expression Patterns as a Biomarker of AHR Activation



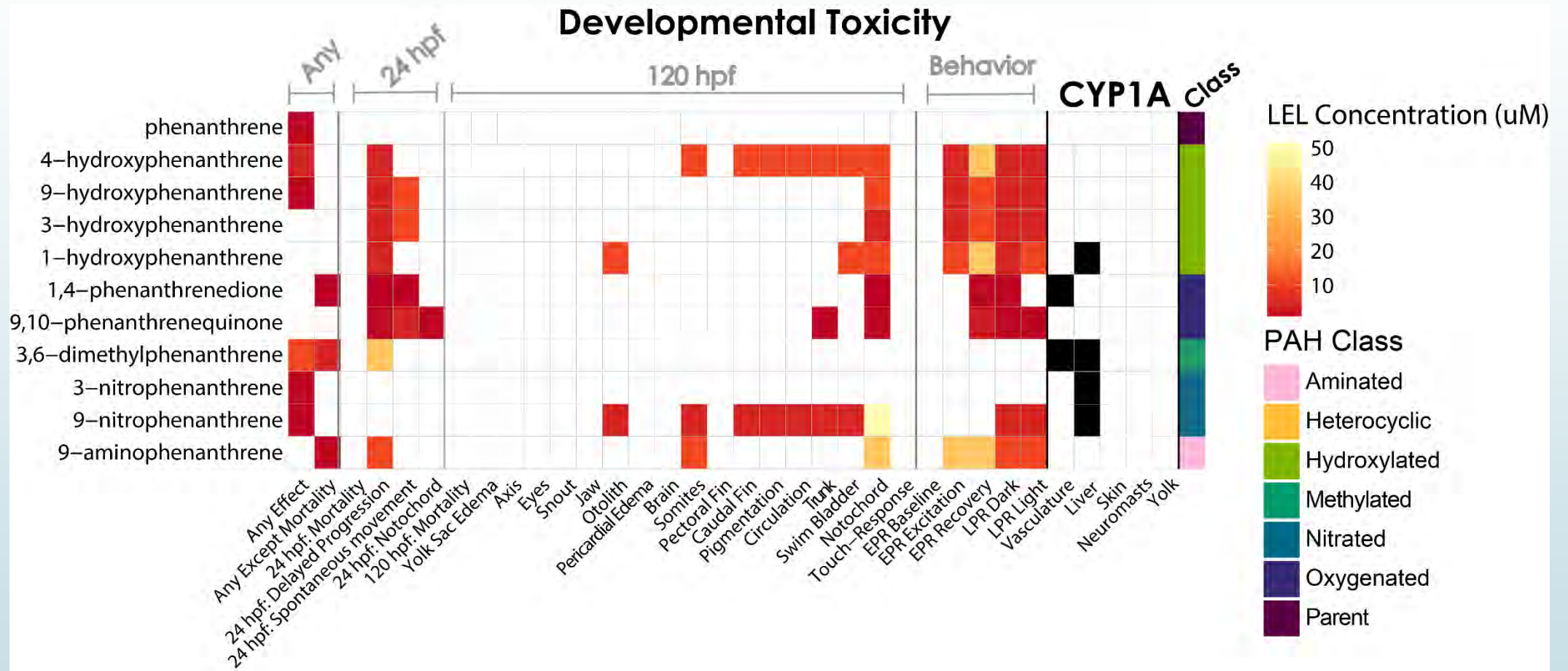
120 hpf

# Summary of Results

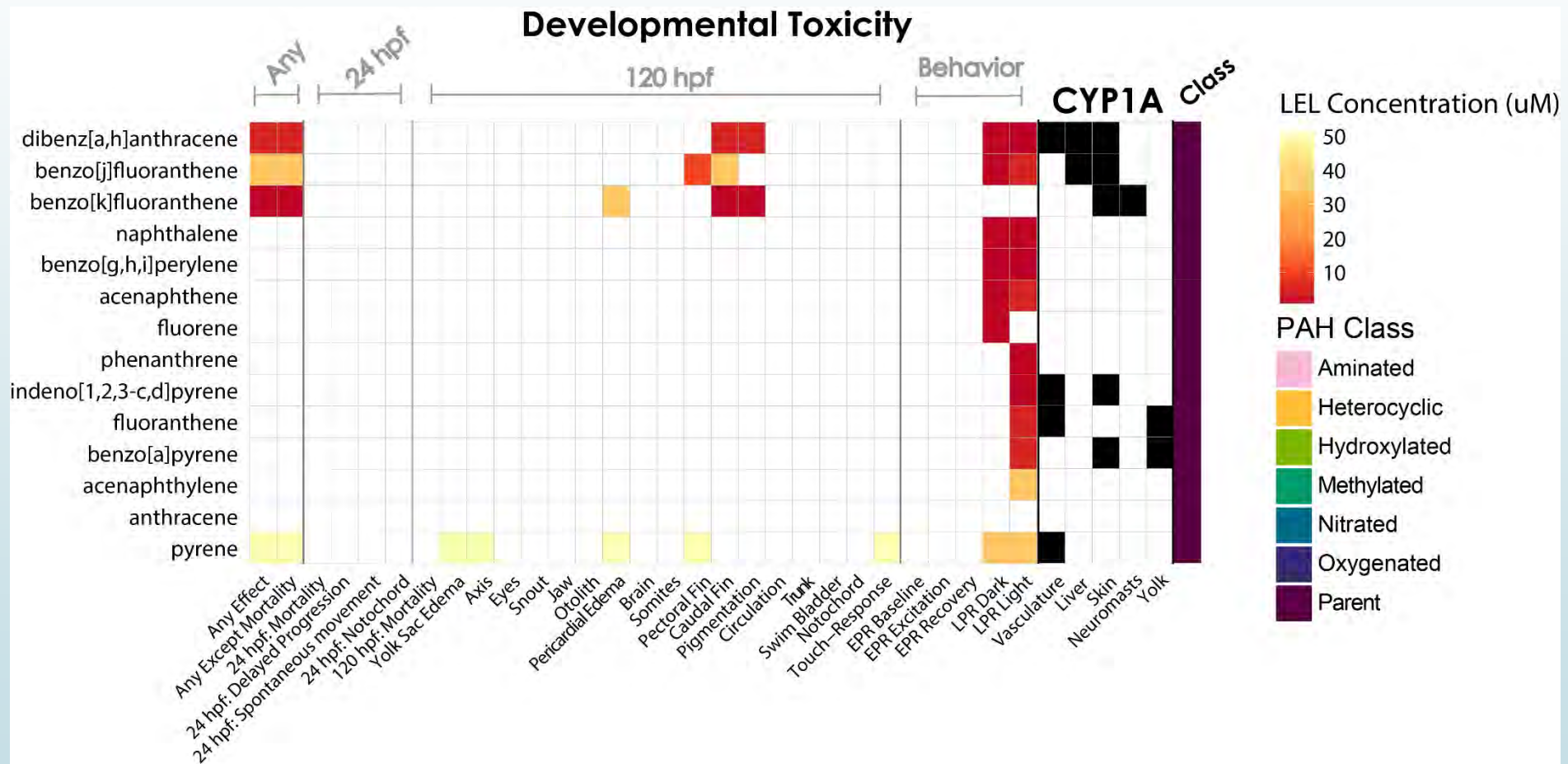




# Example Parent and Derivatives



# 16 EPA Priority PAHs Do Not Reflect Full Range of Effects



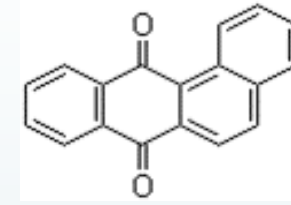




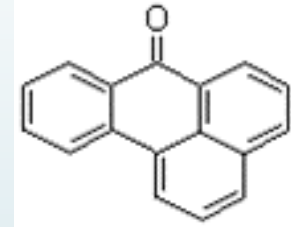


# Case Study – two OPAHs

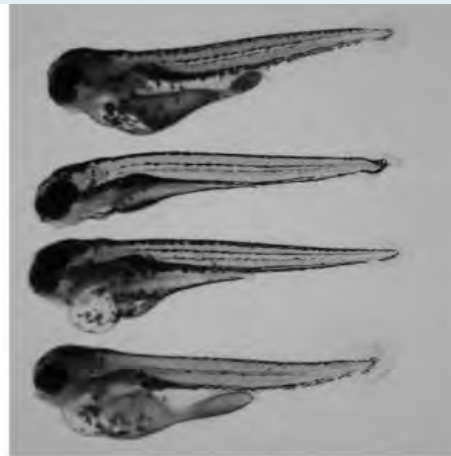
Benz[a]anthracene-7,12-dione  
(7,12-Ba[A]Q)



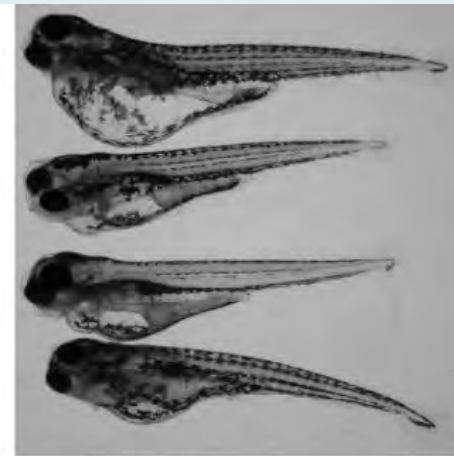
1,9-Benz-10-anthrone (BEZO)



Control

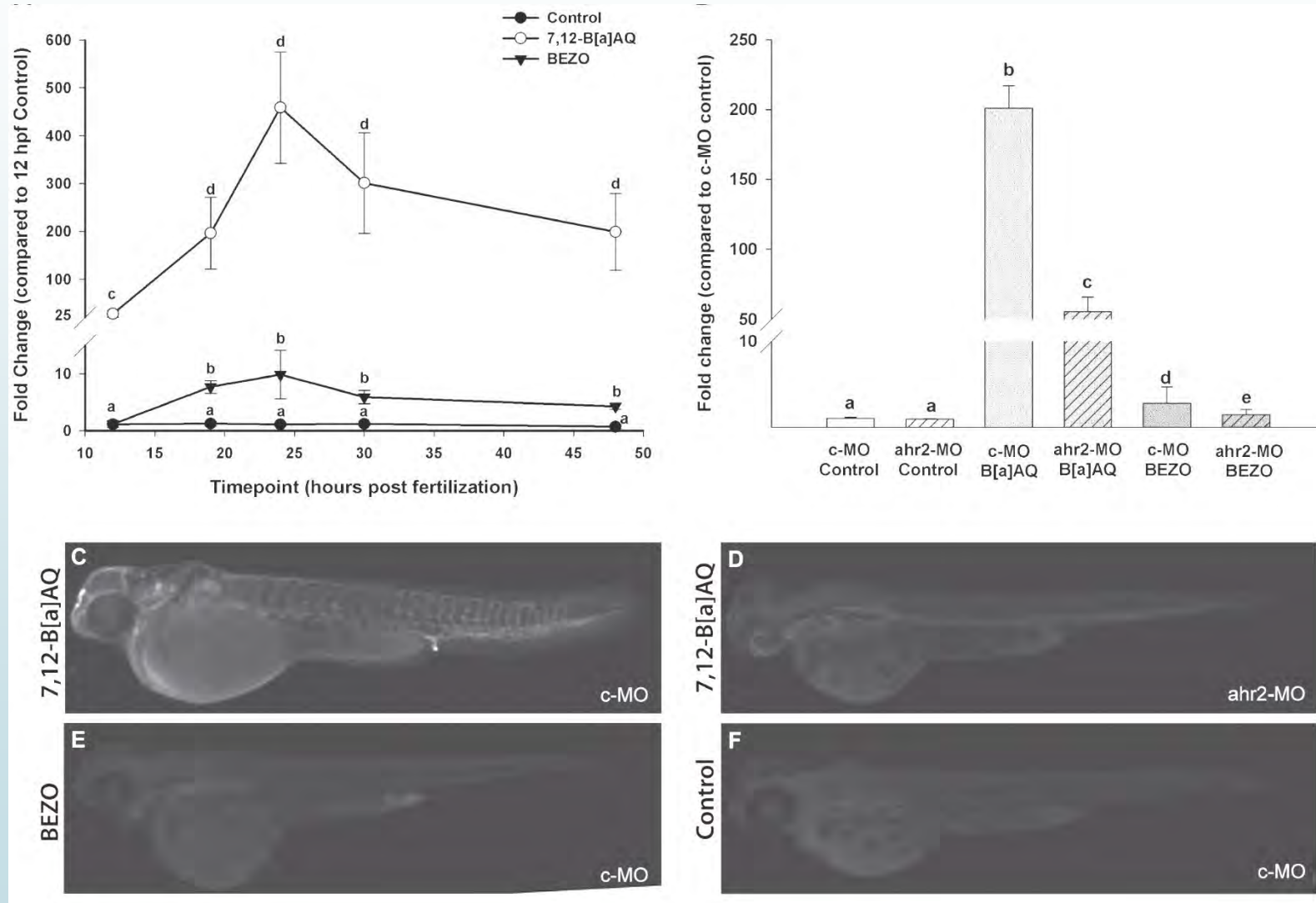


7,12-B[a]AQ

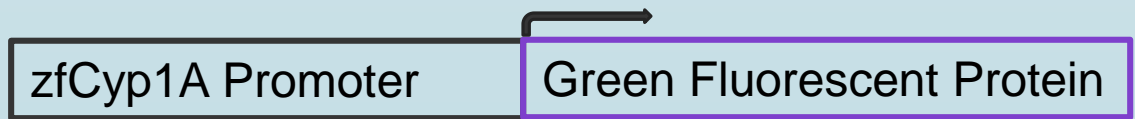
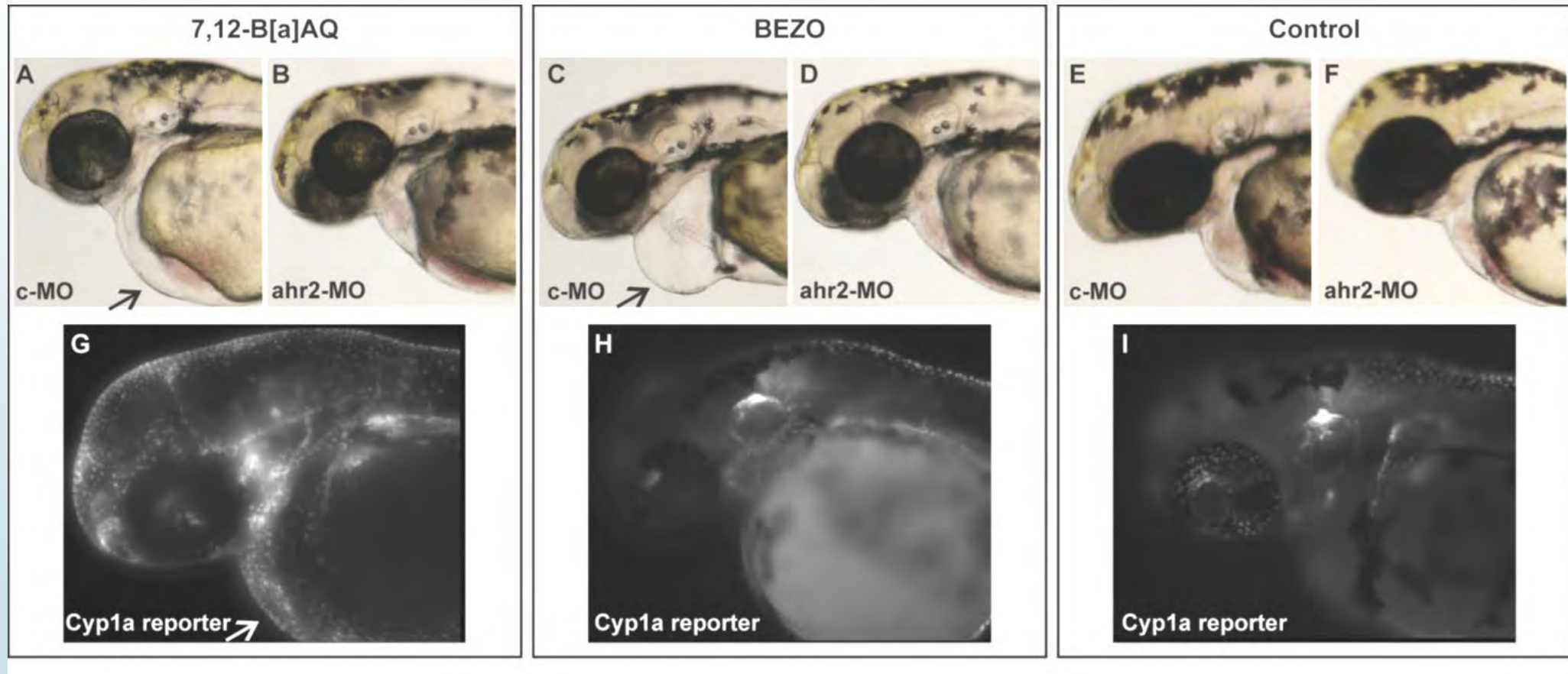


BEZO

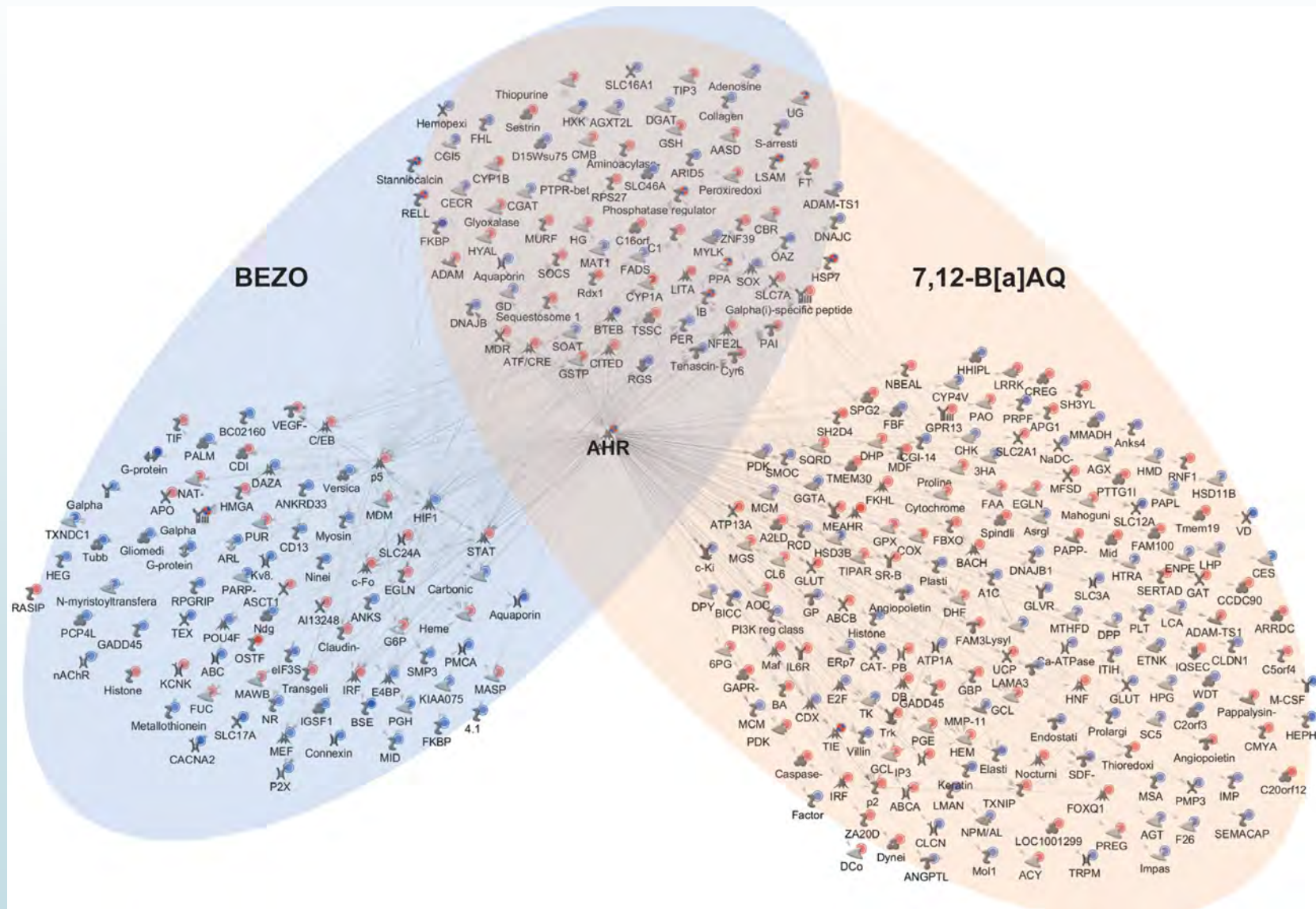
# Differential Downstream AHR Activation (Measured CYP1A Expression)



# Overt Toxicity for both – AHR2 Dependent



# Complexity of Defining Molecular Responses – Even if MIE is known





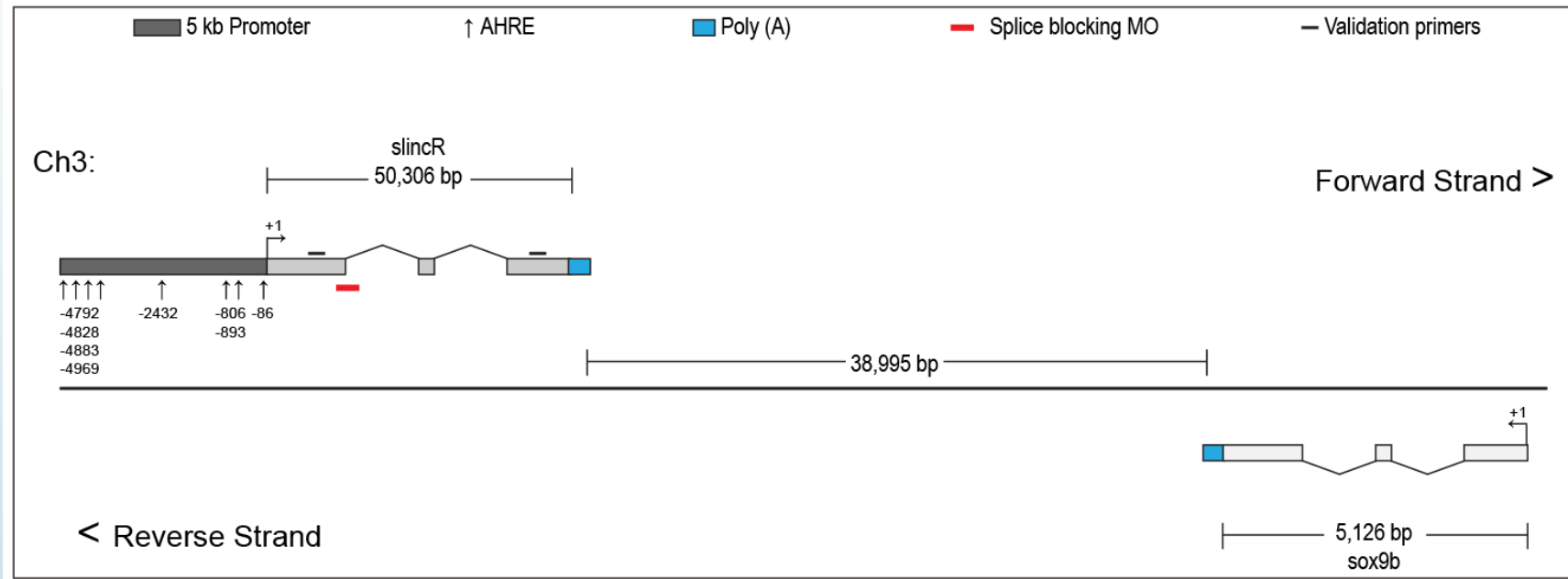
# Looking Downstream of AHR2 activation

# Novel Transcript identified as long non-coding RNA (lncRNA) and mapped adjacent to Sox9b gene

Top significantly elevated transcripts identified from RNA-Seq in whole 48 hpf embryo exposure to 10  $\mu$ M B[a]AQ

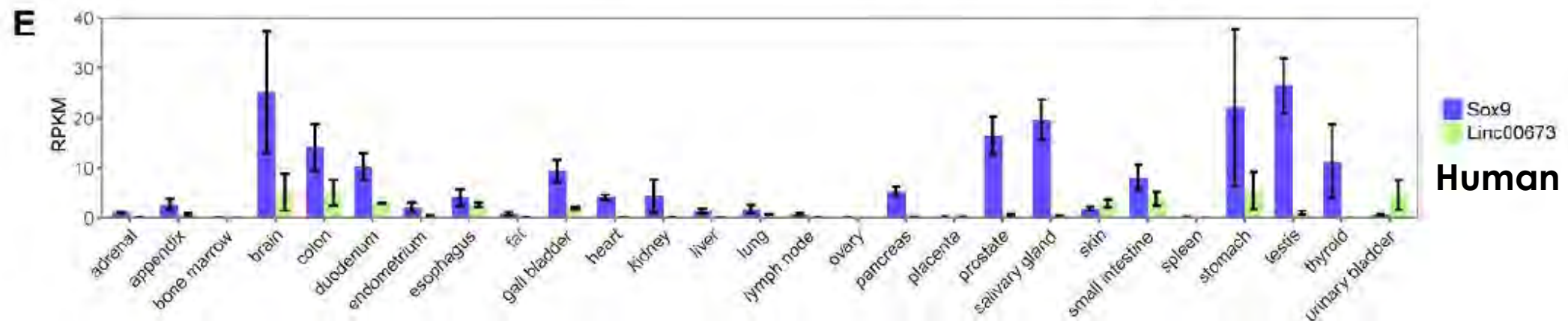
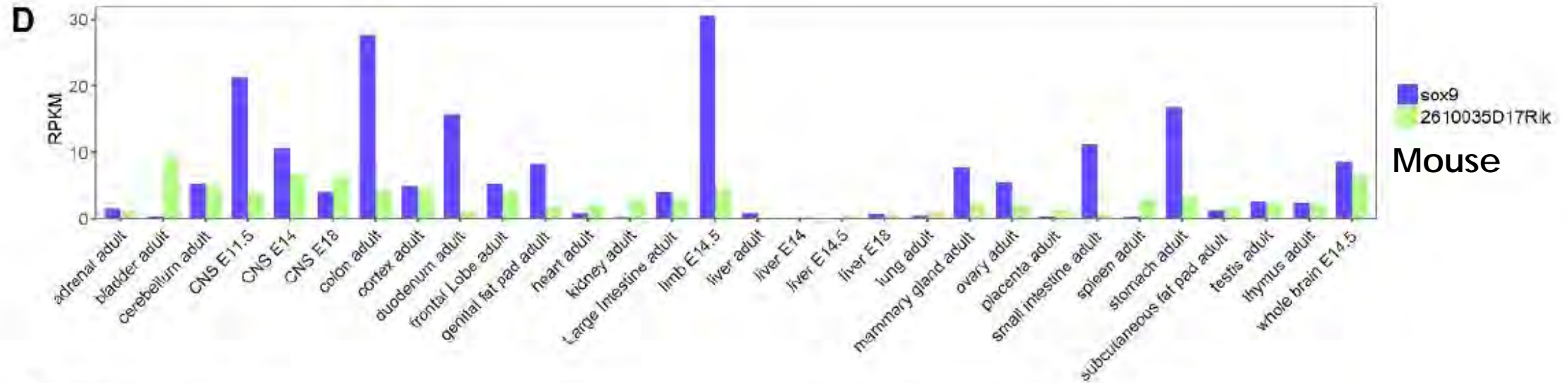
**sox9b long intergenic non-coding RNA (slincR)**

Gene	log2 (fold change)
cyp1a	7.9
cyp1c1	4.8
cyp1c2	4.6
cyp1b1	3.6
<b>novel transcript</b>	<b>3.2</b>



- Spatial arrangement conserved
- Adjacency of AHREs in promoter region of lncRNA conserved
- Expect lncRNA-target regulation to be similar between fish and mammals

# Sox9b and "SlincR" Expression in Mammalian Tissues



RNA-seq expression data from NCBI BioProjects PRJNA66167 (Mouse) and PRJEB4337 (Human)

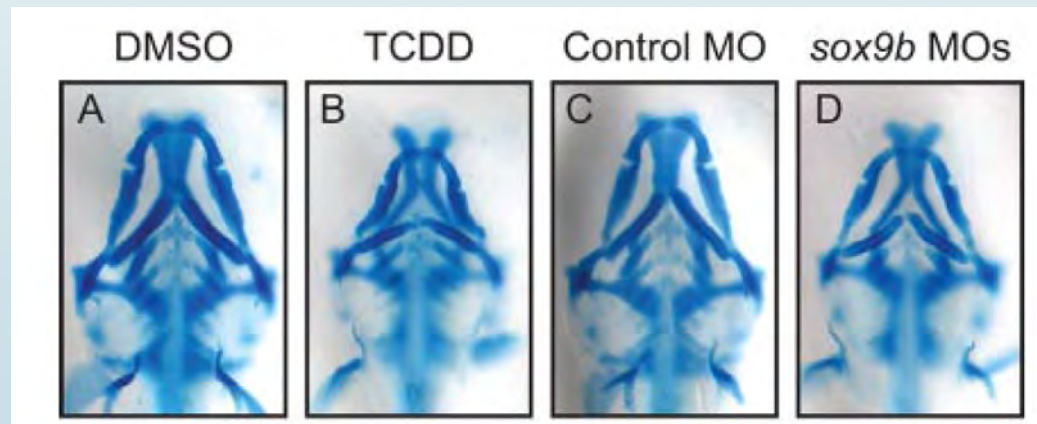
# Sox9 Importance for Toxicology and Human Health

Sox9 in humans:

- Dysregulation of Sox9 has been implicated in skeletal deformities, fibrosis, and cancer (Pritchett 2011)

Upon developmental exposure to TCDD, zebrafish Sox9b:

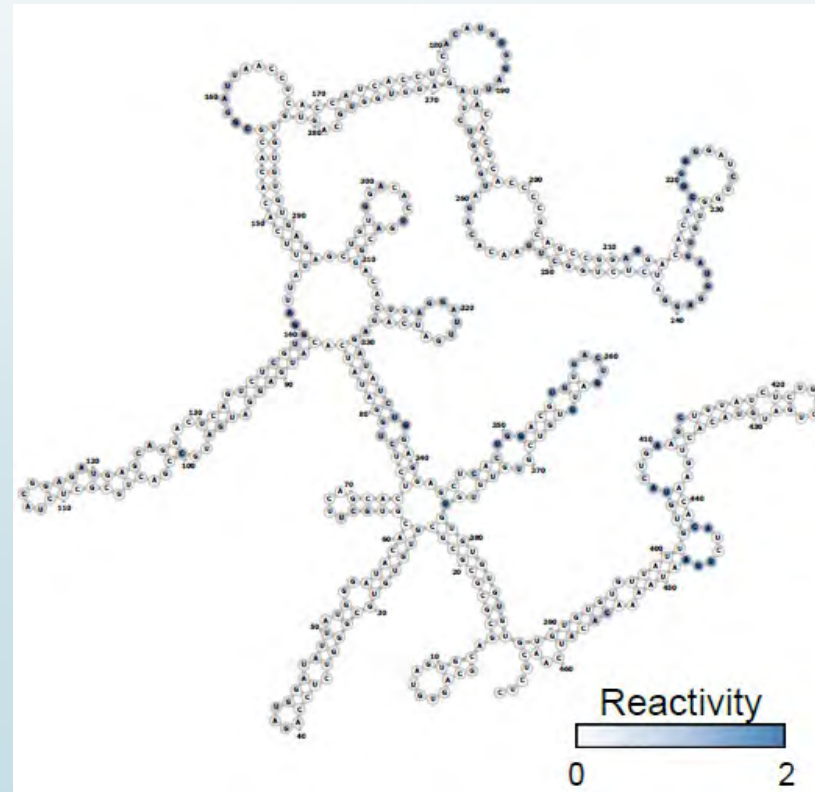
- Is one of the most reduced transcripts in several target organ toxicities
- Is hypothesized to have a **causal** role in TCDD toxicity



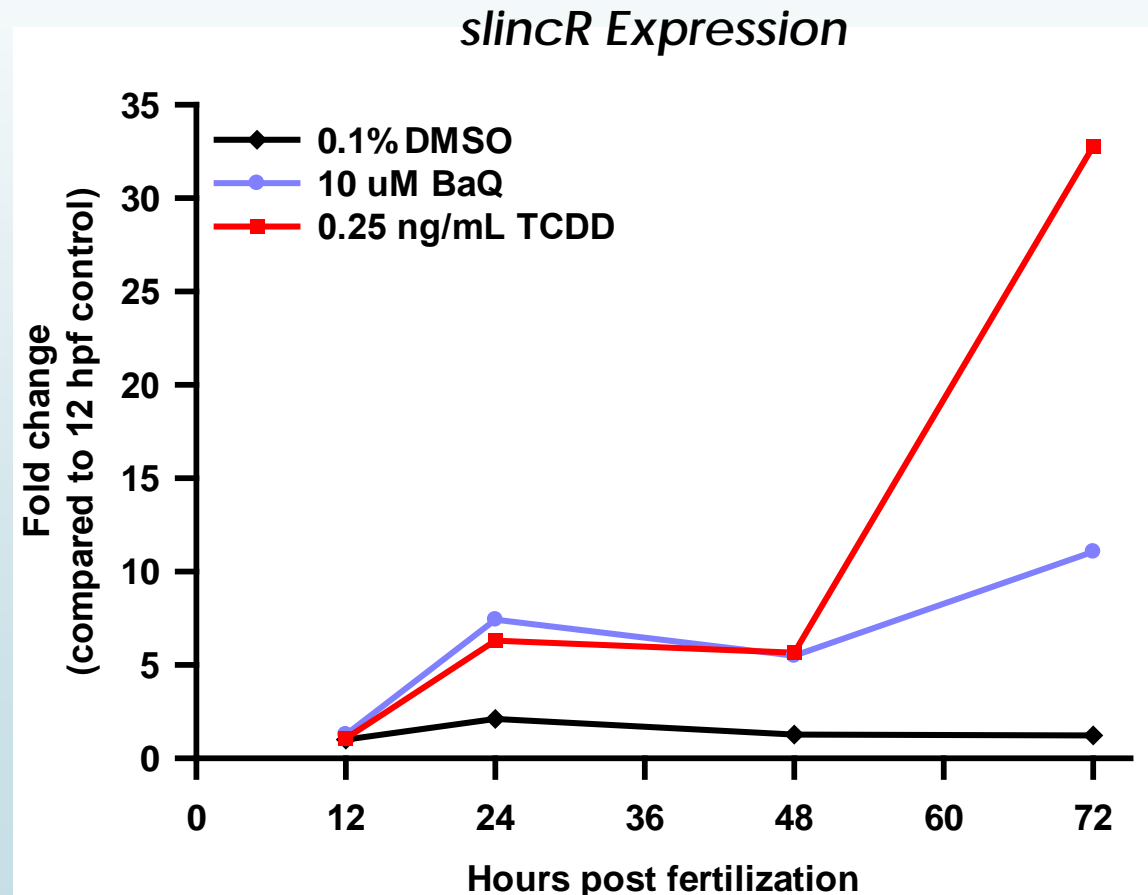
- The mechanism of Sox9b repression was unknown

# Predicted slincR Structure

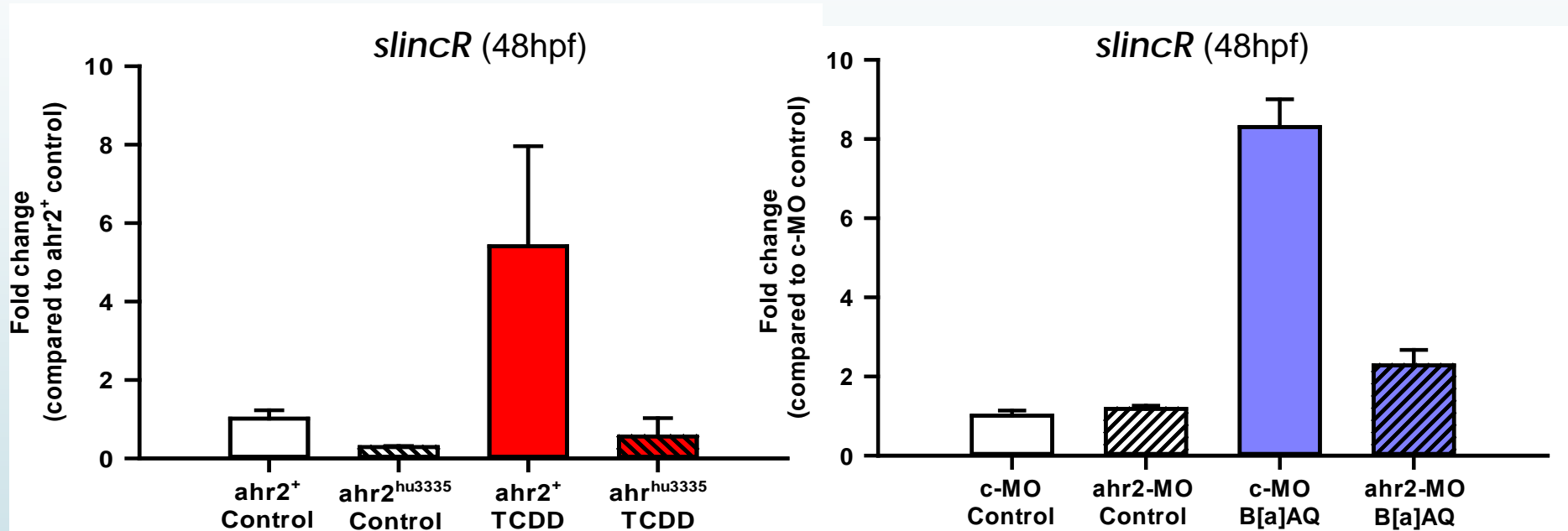
*Selective 2'-Hydroxyl Acylation Analyzed  
by Primer Extension (SHAPE)*



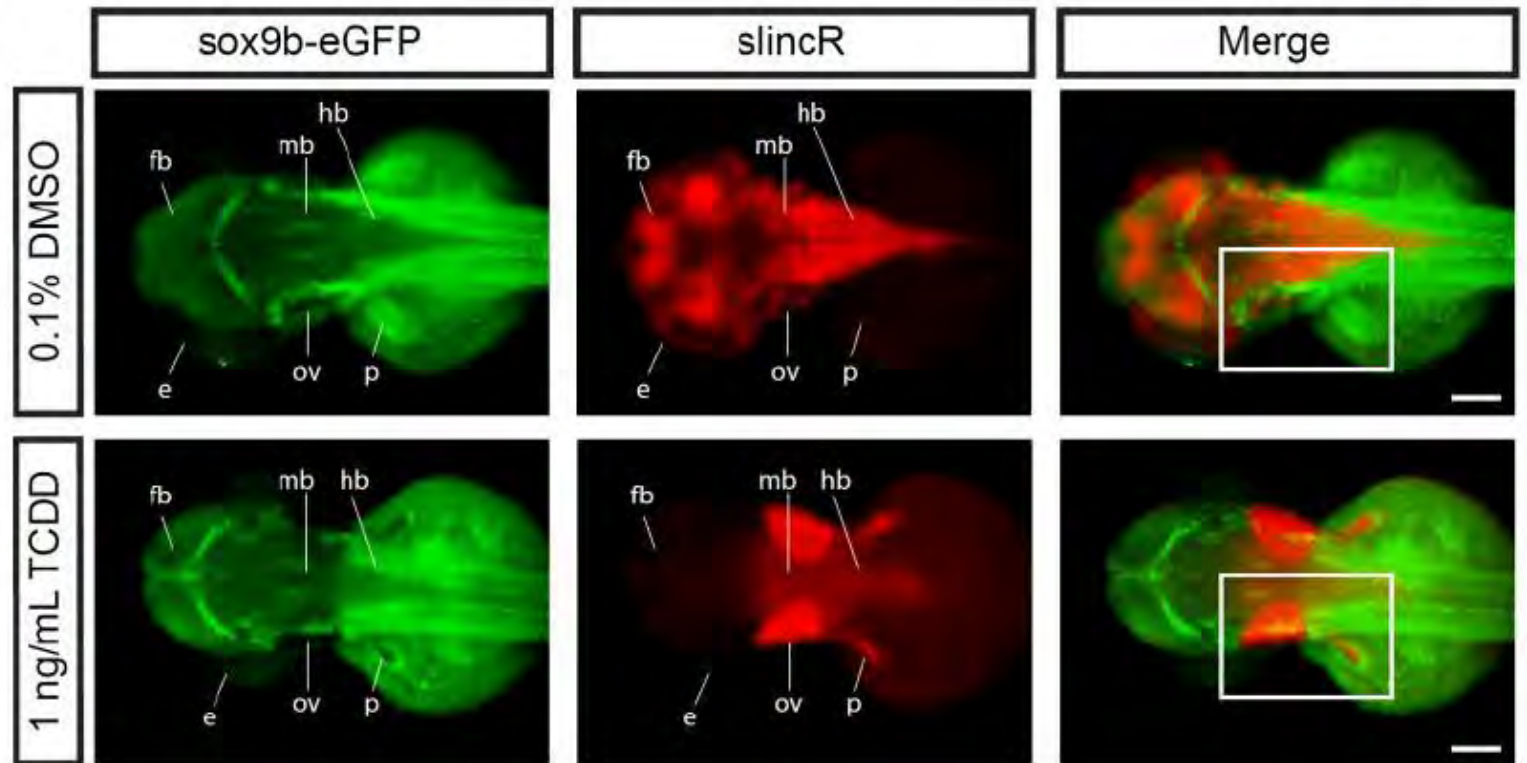
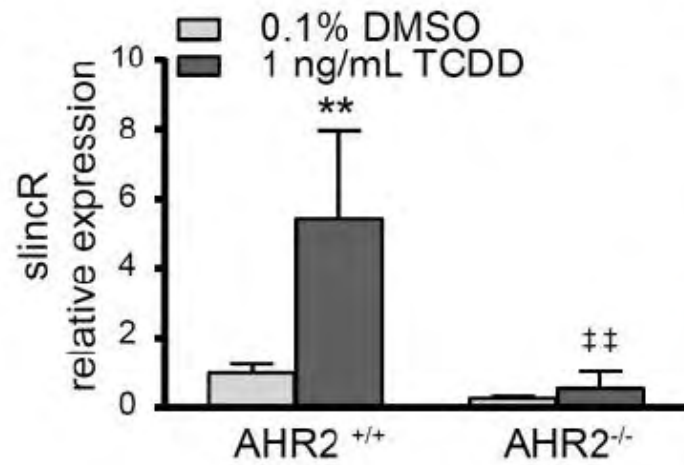
# *slincR* Expression is Elevated by Other AHR Ligands



# Induction of *slincR* Expression Requires AHR2



# Expression of Sox9b and SlicnR Expression

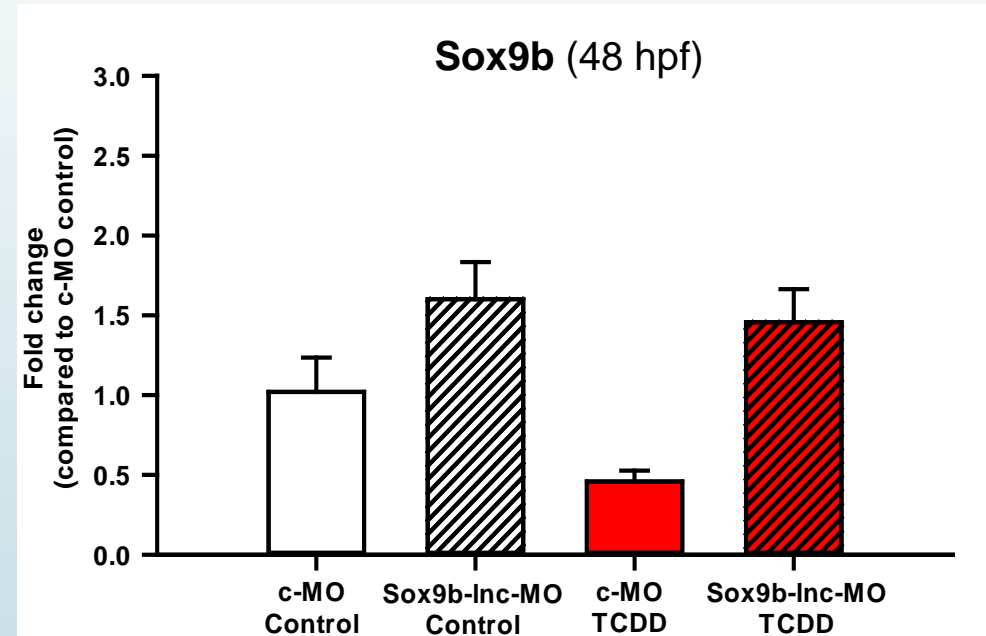
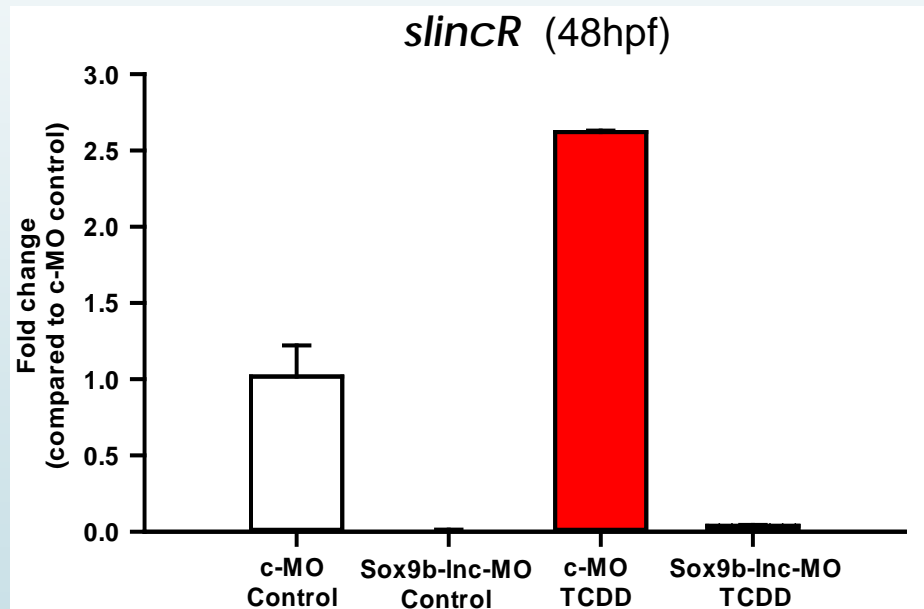


48 hpf



# Knockdown of *slincR* Results In a Relief in Repression of Sox9b

## Confirmation of Knockdown:

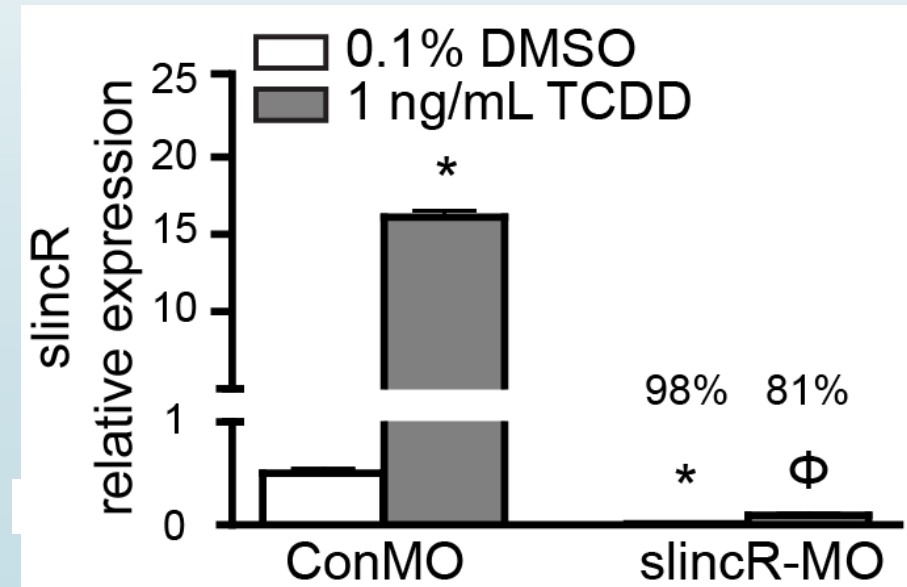
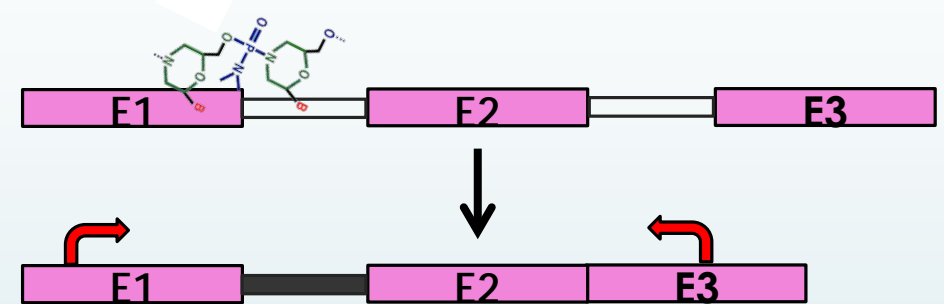


# What is the transcriptional impact of knocking slincR down?

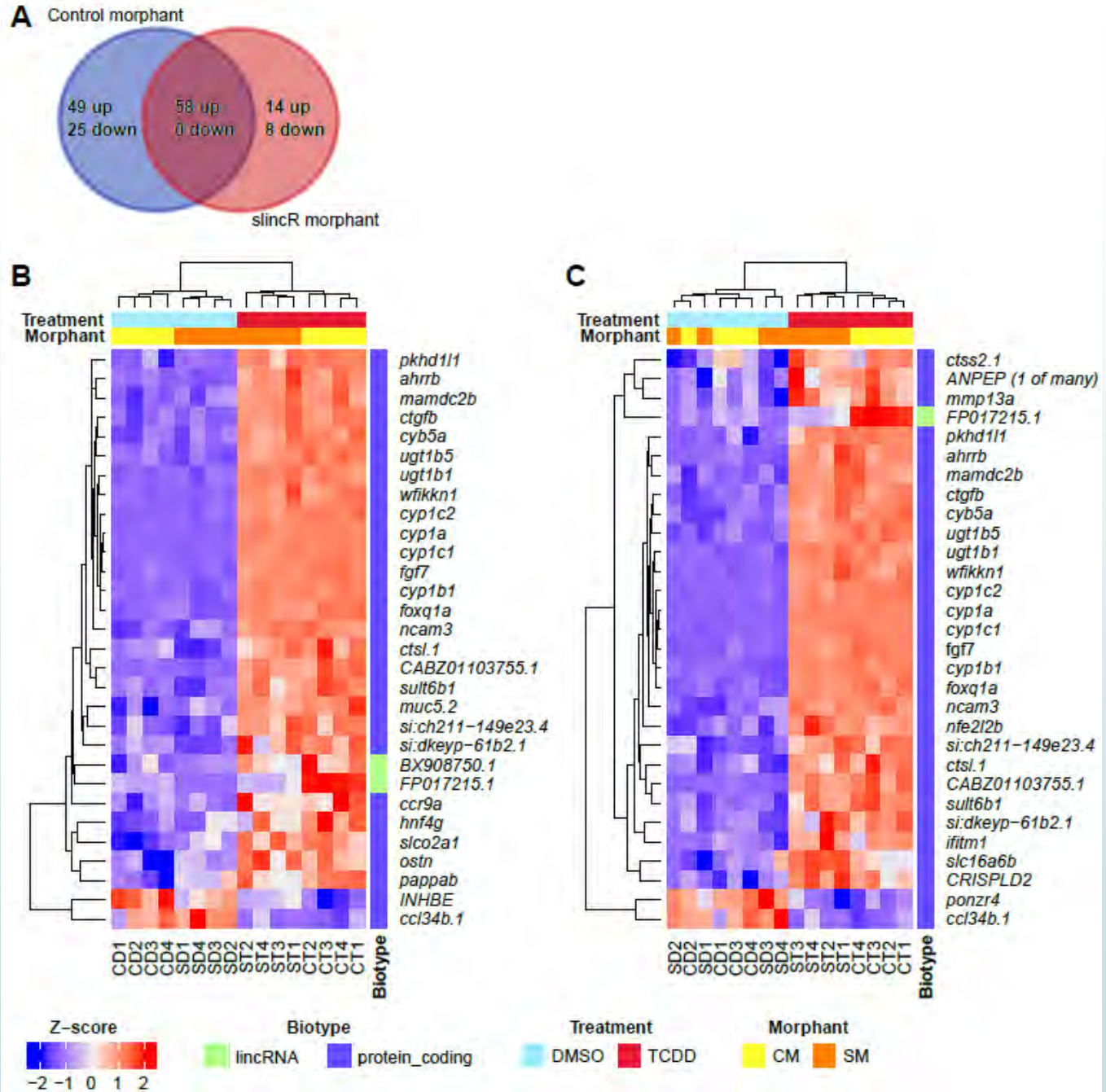
## RNA-Seq Experimental Objectives:

Identify the transcripts, pathways, and biological processes altered or regulated by slincR expression during normal development and during AHR2 activation

## slincR splice blocking morpholino design:

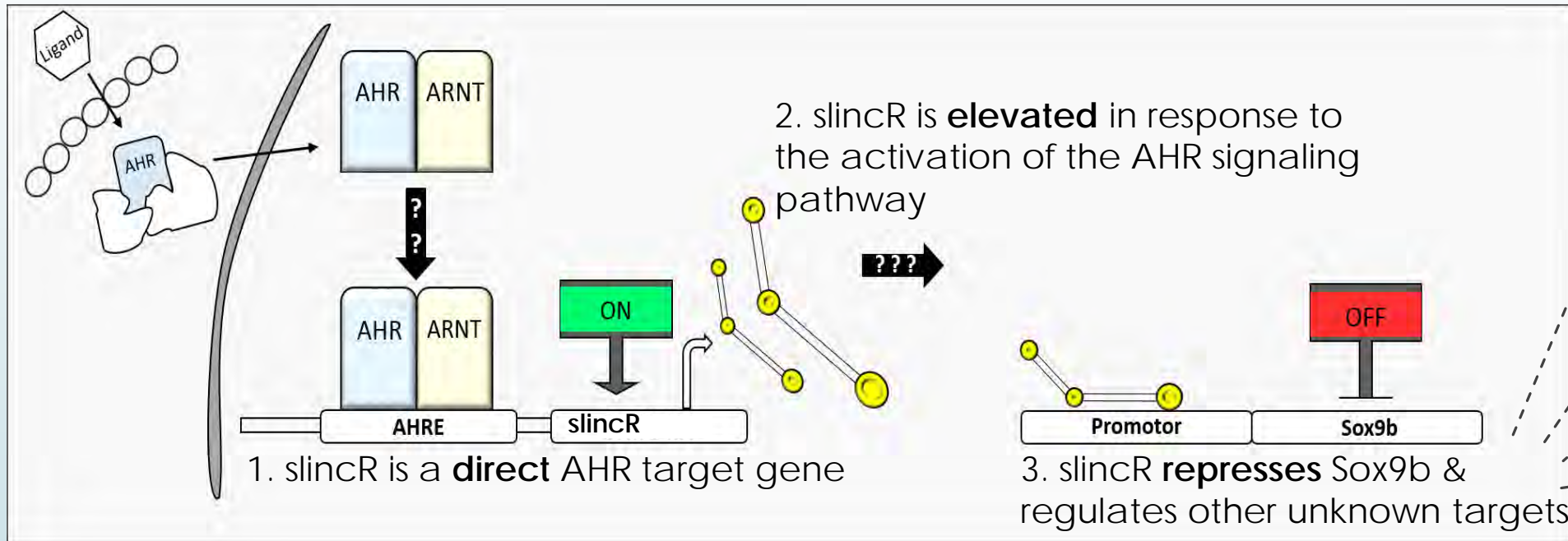


# SlincR Transcriptional regulation



Garcia et al. Signaling events downstream of AHR activation that contribute to toxic responses: The functional role of an AHR-dependent long non-coding RNA (slincR) using the zebrafish Model In press EHP

# Proposed mechanism of AHR-dependent control of gene expression during normal development



4. Altered biological processes

Translation (↓)

Inflammation (↓)

Cell adhesion (↑)

Notch signaling (↑)

# To Summarize

- AHR gene duplication – zebrafish have three functional AHRs
- Increasingly possible to predict ligand/AHR binding
- Platform for structure-based PAH screening
- Complex downstream AHR targets
- AHR/SincR/Sox9b relationship
- Highly amenable for mixture assessments

# Acknowledgements

## Tanguay Lab

Dr. Andrea Knecht

Dr. Britton Goodale

Dr. Mitra Geier

Gloria Garcia

Dr. Lisa Truong

Dr. Michael Simonich

Mike Garland

Laura Holden

Pamela Noyes

Jane LaDu

Hao Truong

Eric Johnson

Greg Gonnerman

Carrie Barton

Annika Swanson

## Engineering Team

Corwin Perrin

Dylan Thrush

David Mandrell

Mushfiq Sarker

Caleb Jephson

Chris Lang

Drew Gabler

## Funding (NIEHS)

- P42 ES016465
- R21 R21ES025421
- RC4 ES019764
- P30 ES000210

## Collaborators

NC State University

David Reif

OSU

Dr. William Bisson

Dr. Siva Kolluri

Ed O'Donnell

Susan Tilton

Kim Anderson

Staci Simonich

PNNL

Katrina Waters

