



## CPP#6

Tox21 Face to Face Meeting  
December 4-5, 2017  
College Park MD



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# Predictive Modeling of Developmental Toxicity with Human Pluripotent Stem Cells

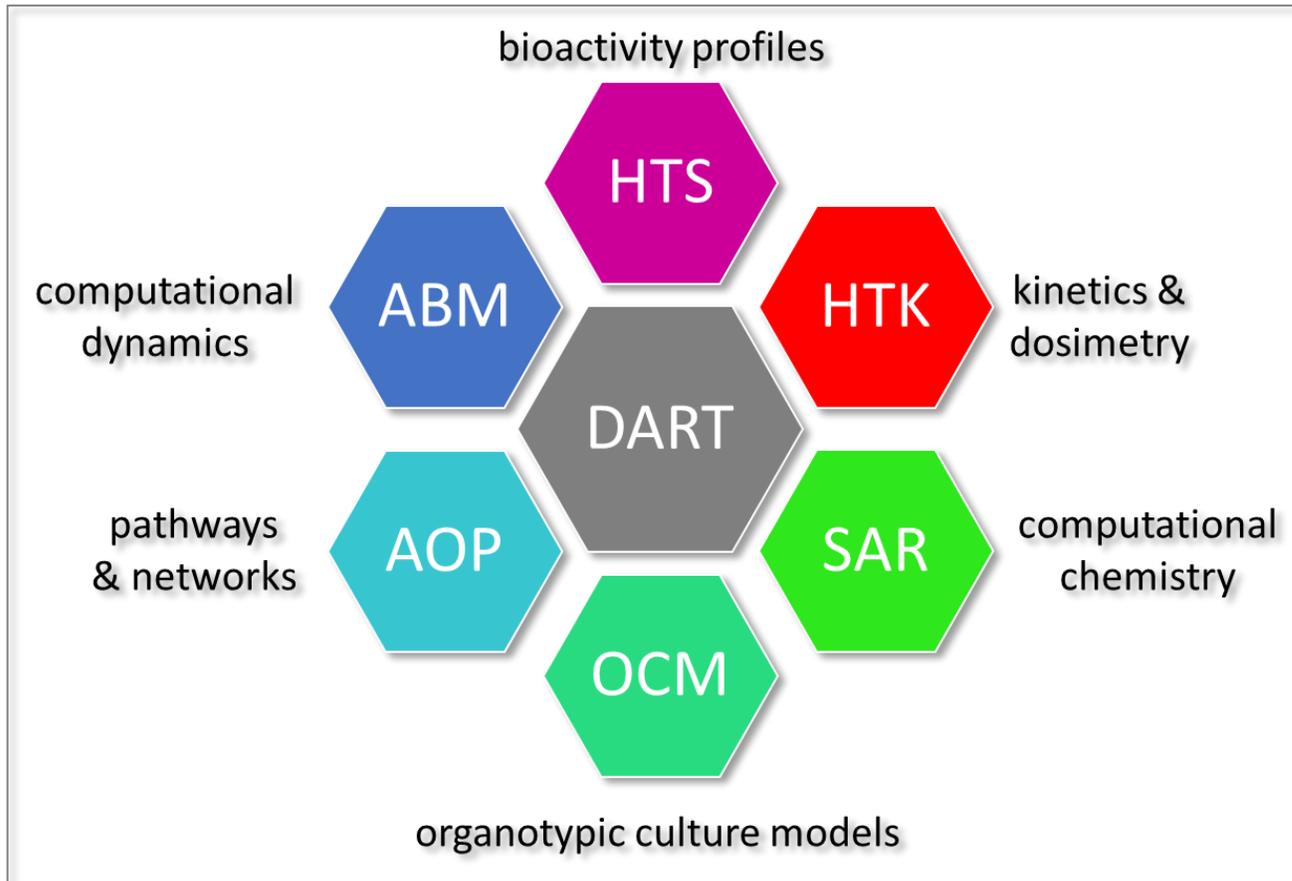
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DISCLAIMER: The views expressed are those of the presenters and do not reflect Agency policy.



# 1. Background



- Goal is computational modeling of ToxCast/Tox21 chemicals for **predictive DART** (prenatal developmental toxicity).
- Project focuses on applying fundamental principles of teratogenesis (birth defects) to human embryonic stem cells:
  - initiating mechanisms
  - dosimetry & bioavailability
  - stage vulnerability
  - chemistry
  - genetic susceptibility



## 2. Hypothesis and Specific Aims

**Hypothesis:** systematic integration of human pluripotent stem cell data into spatially-dynamic computer models of embryonic development will recapitulate the critical determinants of developmental toxicity based on the fundamental principles of teratology.

1. Data-mining: chemical-assay specific correlations to identify relevant mechanisms.
2. Validation: use in a prioritization schema for human teratological outcome.
3. Kinetics: reverse dosimetry for exposure-based hazard prediction in the human fetus.
4. Dynamics: system-specific vulnerabilities of the developing embryo.



# 3. Description of the overall project

Birth Defects Research Part B | **Developmental and Reproductive Toxicology** |  Explore this journal >

Original Article

## Establishment and Assessment of a New Human Embryonic Stem Cell-Based Biomarker Assay for Developmental Toxicity Screening

Jessica A. Palmer , Alan M. Smith, Laura A. Egnash, Kevin R. Conard, Paul R. West, Robert E. Burrier, Elizabeth L.R. Donley, Fred R. Kirchner

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View issue TOC  
Volume 98, Issue 4  
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Pages 343–363

*DevTOX<sup>qP</sup> platform (Stemina Biomarker Discovery)  
Palmer et al. 2013*

*Pluripotent H9 human embryonic stem cell metabolomics assay that “... identified the potential developmental toxicants in the test set with 77% accuracy (57% sensitivity, 100% specificity).”*

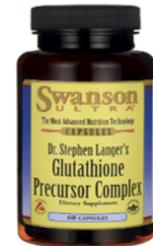


**Ornithine release**  
urea cycle, polyamine & pyrimidine synthesis.



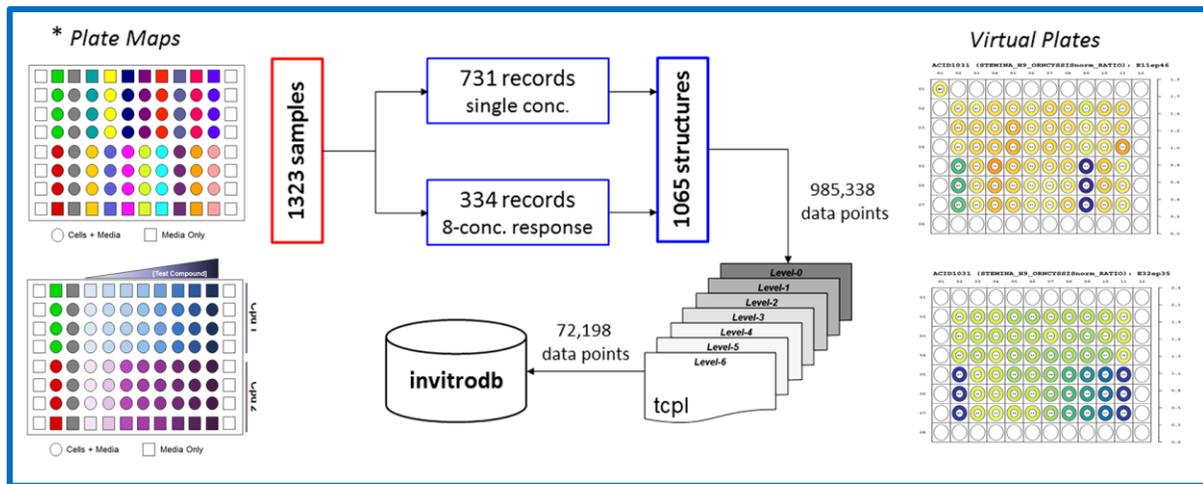
**TI = ORN/CYSS**

**Cystine utilization**  
glutathione synthesis, redox cycling.





# ToxCast Phase I/II library tested (EPA contract EP-D-13-055)



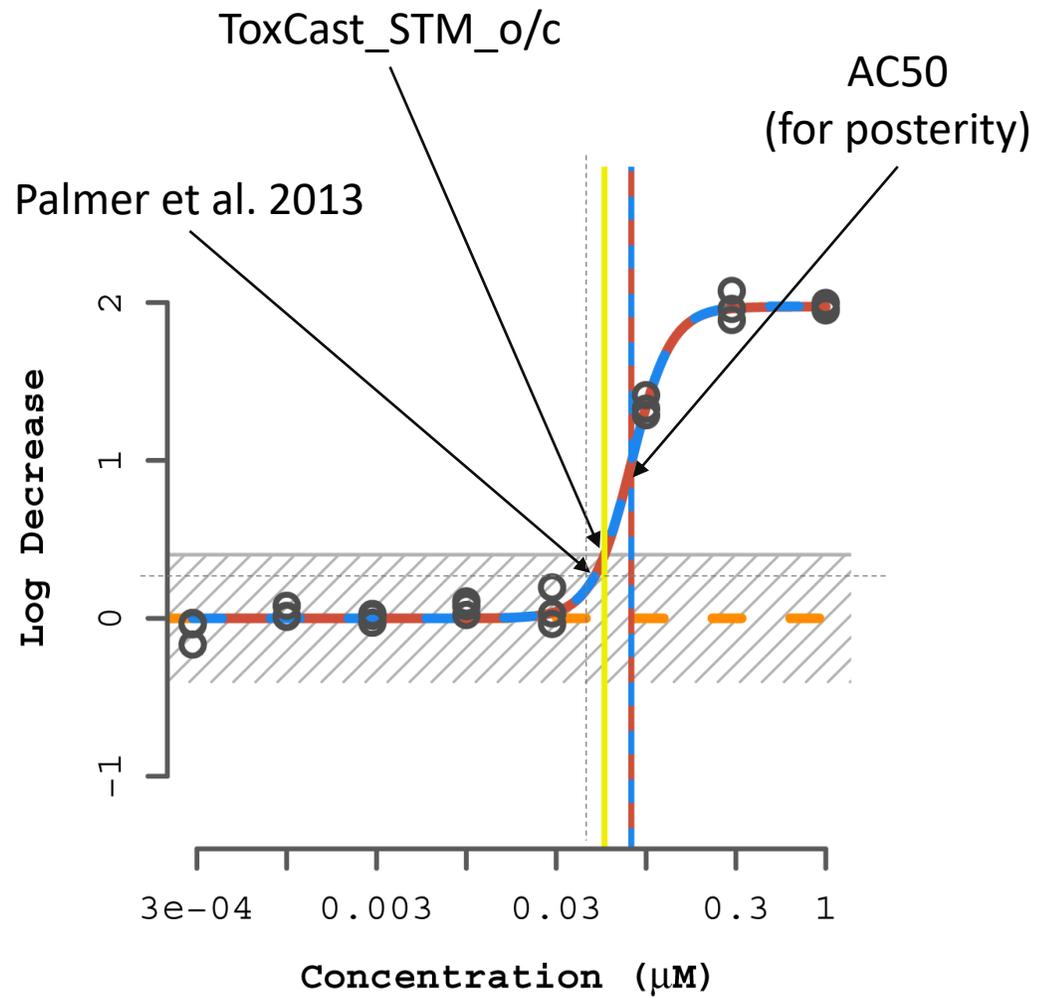
## Main STM assay component identifiers (*acid*):

- ORN in the secretome normalized to control (*acid 1029*)
- CYSS in the secretome normalized to control (*acid 1026*)
- ORN:CYSS ratio (*acid 1031*)
- Cell viability normalized to control (*acid 1114*)

## Invitrodb currently holds results for the STM dataset on 1065 chemicals tested to date:

- concentration-response profiles for 334 chemicals pushed through the ToxCast pipeline (tcpl)
- single-concentration screens for 731 chemicals (mostly negative under the specific test condition)
- dataset has been manually curated and essentially locked-in for data release upon publication (FY18 product)
- positive response on the targeted biomarker translates to a micromolar threshold predicting teratogenicity
- 181 of 1065 (17%) chemicals generated a STM-positive signal <200  $\mu$ M

# Example tcpl readout for the targeted biomarker (o/c ratio)



ASSAY: AEID1691 (STEMINA\_H9\_ORNCYSSISnorm\_RATIO\_dn)

NAME: Methotrexate

CHID: 20822 CASRN: 59-05-2

SPID(S): TP0001302A08

M4ID: 18146613

HILL MODEL (in red):

	tp	ga	gw
val:	1.97	-1.08	3.96
sd:	0.0247	0.0229	1.06

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	1.98	-1.08	3.92	0.721	3.48
sd:	0.0529	0.0226	1.08	35.1	171

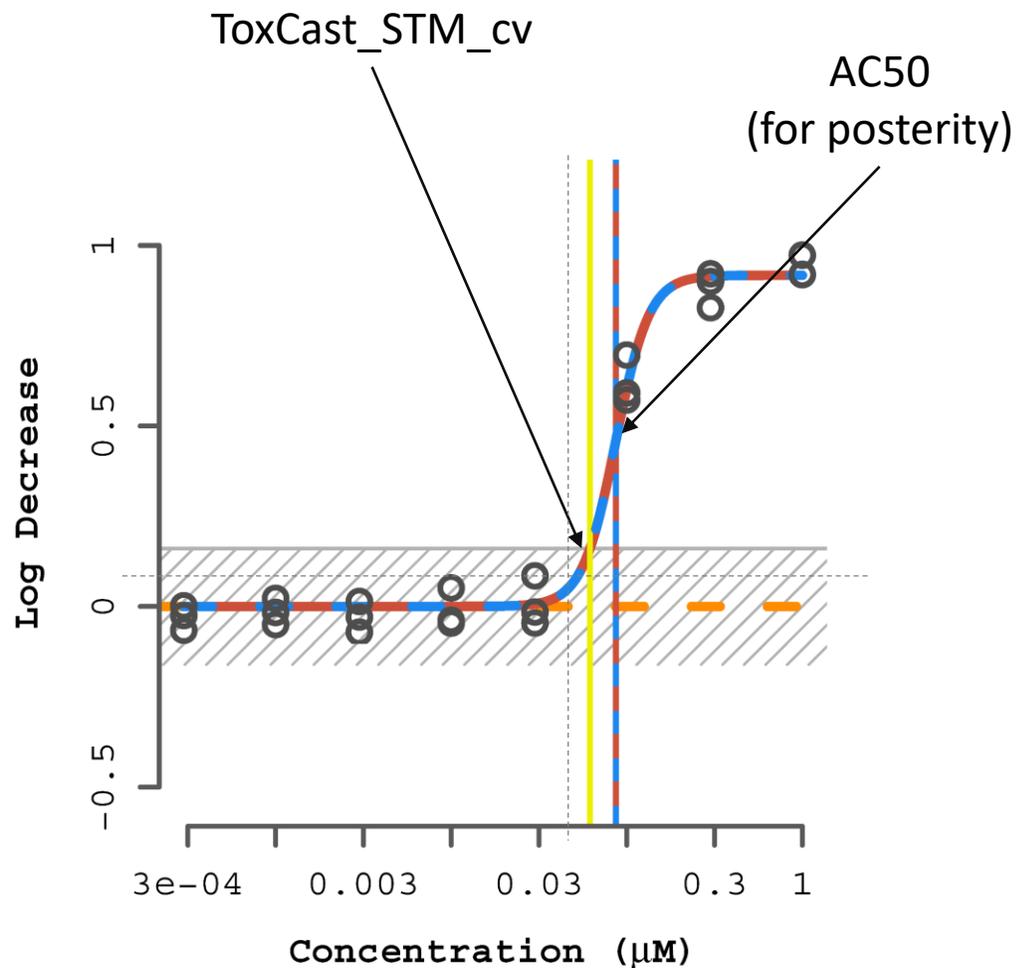
	CNST	HILL	GNLS
AIC:	74.32	-52.43	-48.44
PROB:	0	0.88	0.12
RMSE:	1.1	0.07	0.07

MAX\_MEAN: 1.98      MAX\_MED: 1.96      BMAD: 0.134

ACB: 0.0588      HIT-CALL: 1      FITC: 41      AC50: 0.0828

FLAGS:

Example tcpl readout for cell viability: calculated point of departure reflects an 11% loss



ASSAY: AEID1858 (STEMINA\_H9\_Viability\_Norm)

NAME: Methotrexate

CHID: 20822 CASRN: 59-05-2

SPID(S): TP0001302A08

M4ID: 15456100

HILL MODEL (in red):

	tp	ga	gw
val:	0.917	-1.06	4.56
sd:	0.0185	0.0404	3.05

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	0.917	-1.06	4.56	1.79	5.05
sd:	0.0185	0.0404	3.05	1650	4670

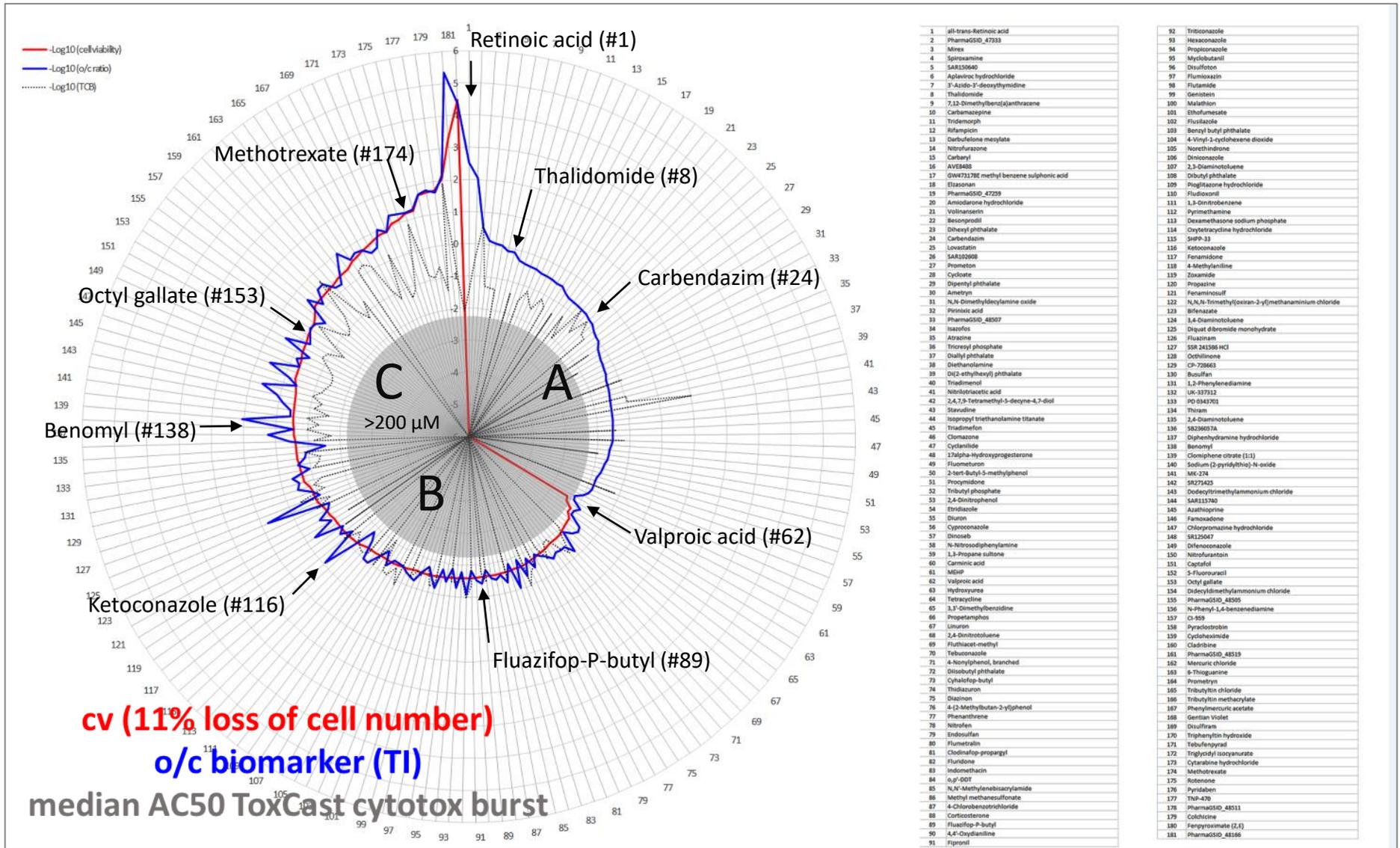
	CNST	HILL	GNLS
AIC:	37.29	-69.83	-65.83
PROB:	0	0.88	0.12
RMSE:	0.51	0.05	0.05

MAX\_MEAN: 0.938      MAX\_MED: 0.921      BMAD: 0.0537

ACB: 0.0617      HIT-CALL: 1      FITC: 41      AC50: 0.0867

FLAGS:

# Stratification of 181 STM-positives





# Performance Check

- ToxCast\_STM anchored to DevTox benchmark compounds aimed at assessing alternative models<sup>1</sup> and having information on human pregnancy risk.
- Overall accuracy 82.1% (71% sensitivity, 100% specificity, MCC = 0.695, n=39).
- Compares well to the Palmer et al. (2013) who reported an overall accuracy of 77% (57% sensitivity, 100% specificity).

<sup>1</sup> Genschow et al. 2002; West et al. 2010; Daston et al. 2014; Augustine-Rauch et al. 2016; Wise et al. 2016

NAME	CASRN	Benchmark	Pregnancy Risk Category	TI (uM)	Class
all-trans-Retinoic acid	302-79-4	tera togen	X	0.003	TP
Cytarabine hydrochloride	69-74-9	tera togen	D	0.054	TP
Methotrexate	59-05-2	tera togen	X	0.059	TP
Diphenhydramine hydrochloride	147-24-0	tera togen	B	0.588	TP
Thalidomide	50-35-1	tera togen	X	1.27	TP
5-Fluorouracil	51-21-8	tera togen	D	2.02	TP
Carbamazepine	298-46-4	tera togen	C	2.29	TP
Busulfan	55-98-1	tera togen	D	2.31	TP
Rifampicin	13292-46-1	tera togen	C	2.46	TP
Amiodarone hydrochloride	19774-82-4	tera togen	D	5.10	TP
Lovastatin	75330-75-5	tera togen	X	6.67	TP
Stavudine	3056-17-5	tera togen	C	32.5	TP
Dexamethasone sodium phosphate	2392-39-4	tera togen	C	37.7	TP
Indomethacin	53-86-1	tera togen	D	72.7	TP
Hydroxyurea	127-07-1	tera togen	D	74.9	TP
Valproic acid	99-66-1	tera togen	D	155	TP
MEHP	4376-20-9	tera togen	D	167	TP
5,5-Diphenylhydantoin	57-41-0	tera togen	D	1000	FN
6-Propyl-2-thiouracil	51-52-5	tera togen	D	1000	FN
Boric acid	10043-35-3	tera togen	N	1000	FN
Cyclophosphamide monohydrate	6055-19-2	tera togen	D	1000	FN
Diethylstilbestrol	56-53-1	tera togen	X	1000	FN
Phenobarbital sodium	57-30-7	tera togen	D	1000	FN
Warfarin	81-81-2	tera togen	X	1000	FN
Salicylic acid	69-72-7	non-tera togen	C	513	TN
Acetaminophen	103-90-2	non-tera togen	B	1000	TN
Acrylamide	79-06-1	non-tera togen	NTP	1000	TN
Aspirin	50-78-2	non-tera togen	C	1000	TN
Bisphenol A	80-05-7	non-tera togen	NTP	1000	TN
Butylparaben	94-26-8	non-tera togen	GRAS	1000	TN
Caffeine	58-08-2	non-tera togen	B	1000	TN
D-Camphor	464-49-3	non-tera togen	C	1000	TN
Dimethyl phthalate	131-11-3	non-tera togen	NTP	1000	TN
Folic acid	59-30-3	non-tera togen	A	1000	TN
Isoniazid	54-85-3	non-tera togen	C	1000	TN
Retinol	68-26-8	non-tera togen	A	1000	TN
Saccharin	81-07-2	non-tera togen	A	1000	TN
Sodium L-ascorbate	134-03-2	non-tera togen	A	1000	TN
Sulfasalazine	599-79-1	non-tera togen	B	1000	TN



## 4. Timeline and Deliverables

### ✓ Year 0.5

- Issue task order 85-90 NTP/NICEATM chemicals for iPSC data generation (Aim 3)
- Build literature mining database for chemical-assay connectivity (Aim 1)

### Year 1.0

- Integrate results from iPSC data with extant H9 (ToxCast) data in tcpl (Aim 1)
- Run additional assays needed to strengthen concordance between iPSC and H9 (Aim 1)

### Year 1.5

- Comprehensive correlation analysis to ToxRefDB and NTP animal studies (Aim 3)
- Implement the human fetal PBPK exposure model (Aim 2)

### Year 2.0

- Quality Control (QC) and manual curation plan to evaluate the dataset (Aim 1)
- Draft relevant manuscripts for publication (Aims 1 & 2)

### Year 2.5

- Compendium of sensitivity-specificity predictors based on adverse outcomes (Aim 3)
- Develop 3 case studies for computational dynamics at windows of vulnerability (Aim 4)

### Year 3.0

- Virtual embryo agent-based models to simulate spatial dynamics (Aim 4)
- Draft relevant manuscripts for publication (Aims 3 & 4).



## 5. Progress to date (Year 0.5)

*Toward issuing a task order 85-90 NTP/NICEATM chemicals for iPSC data generation (Aim 3):*

- Nicole learned of a draft document “*ICH Reference Compounds for Qualifying Alternative Assays*” candidate list of 84 DevTox reference compounds with detailed dosimetry. She inquired whether the Expert Working Group of ICHS5(R3) could share kinetic and toxicity information, but it was not ready until Q1-2018.
- Finalized 81 chemicals based on : (a) NTP prenatal animal studies; (b) potential for human exposure and suspected developmental effects; (c) high priority to NTP, EPA, or other ICCVAM member agencies; and (d) inclusion in the ICH guidance: “*Detection of toxicity to reproduction for human pharmaceuticals.*”
- Almost ½ of the list shares overlap with ToxCast\_STM (H9 cells). Stemina kindly shared the list of compounds they already compared between H9 and iPSC versions of the platform.



# 5. Progress to date (Year 0.5)

*Toward building a literature mining database for chemical-assay connectivity (Aim 1):*

- Nancy Baker (NCCT) developed a sifter/literature miner knowledge mapping tool for the test compounds.

Feel free to delete columns after Column D and rows after Row 4.

Update Article Counts   Hide queries   View / edit queries   Heat Map by column   Heat Map by row

Notes	Preferred Name	Chemical / entity query	MeSH Link	Any article	Stem cells	(dna/drug effects OR DNA)
Expect: 1 : ICH List, no Stemina data	Almokalant	Almokalant OR 123955-10-2[RN]	<a href="#">almokalant</a>	79		
Expect: 0 : ICH List, no Stemina data	Chlorthalidone	Chlorthalidone OR 77-36-1[RN]	<a href="#">Chlorthalidone</a>	1777		
Expect: 1 : ICH List, no Stemina data	Diltiazem	Diltiazem OR 42399-41-7[RN]	<a href="#">Diltiazem</a>	9262		
Expect: 0 : ICH List, no Stemina data	Hydrochlorothiazide	Hydrochlorothiazide OR 58-93-5[RN]	<a href="#">Hydrochlorothiazide</a>	8546		
Expect: 1 : ICH List, no Stemina data	Topiramate	Topiramate OR 97240-79-4[RN]	<a href="#">topiramate</a>	4363		
Expect: 1 : ICH List, no Stemina data	Trimethadione	Trimethadione OR 127-48-0[RN]	<a href="#">Trimethadione</a>	606		
Expect: 1 : ICH List, no Stemina data	Cisplatin	Cisplatin OR 15663-27-1[RN]	<a href="#">Cisplatin</a>	65866	19	9211
Expect: 1 : ICH List, no Stemina data	Cyclophosphamide	Cyclophosphamide OR 6055-19-2[RN]	<a href="#">Cyclophosphamide</a>	68343	79	47851
Expect: 1 : ICH List, no Stemina data	Thiotepa	Thiotepa OR 52-24-4[RN]	<a href="#">Thiotepa</a>	3349	8	962
Expect: 1 : ICH List, no Stemina data	Aspirin	Aspirin OR 50-78-2[RN]	<a href="#">Aspirin</a>	61104	5	469
Expect: 1 : ICH List, no Stemina data	Captopril	Captopril OR 62571-86-2[RN]	<a href="#">Captopril</a>	12925	0	50
Expect: 1 : ICH List, no Stemina data	Enalapril	Enalapril OR 75847-73-3[RN]	<a href="#">Enalapril</a>	8244	0	40
Expect: 1 : ICH List, no Stemina data	Methimazole (Thiamazole)	Methimazole OR Thiamazol OR 60-56-0[RN]	<a href="#">Methimazole</a>	4225	2	63
Expect: 0 : ICH List, no Stemina data	Saxagliptin	Saxagliptin OR 361442-04-8[RN]	<a href="#">saxagliptin</a>	559	0	0

Abstract with highlights

Article: 21820460

Title: Distribution of bisphenol A into tissues of adult, neonatal, and fetal Sprague-Dawley rats.

Abstract: Distribution of bisphenol A into **tissues** of adult, neonatal, and fetal Sprague-Dawley rats. Abstract: Bisphenol A (BPA) is an important industrial chemical used in the manufacture of polycarbonate plastic products and epoxy resin-based food can liners. The presence of BPA metabolites in urine of 90% of Americans aged 6-60 suggests ubiquitous and frequent exposure in the range of 0.02-0.2µg/kgbw/d (25th-95th percentiles). The current study used LC/MS/MS to measure placental transfer and concentrations of aglycone (receptor-active) and conjugated (inactive) BPA in **tissues** from Sprague-Dawley rats administered deuterated BPA (100µg/kg bw) by oral and IV routes. In adult female rat **tissues**, the **tissue**/serum concentration ratios for aglycone BPA ranged from 0.7 in liver to 5 in adipose **tissue**, reflecting differences in **tissue** perfusion, composition, and metabolic capacity. Following IV administration to dams, placental transfer was observed for aglycone BPA into fetuses at several gestational days (GD), with fetal/maternal serum ratios of 2.7 at GD 12, 1.2 at GD 16, and 0.4 at GD 20; the corresponding ratios for conjugated BPA were 0.43, 0.65, and 3.7. These ratios were within the ranges observed in adult **tissues** and were not indicative of preferential accumulation of aglycone BPA or hydrolysis of conjugates in fetal **tissue** in vivo. Concentrations of aglycone BPA in GD 20 fetal **brain** were higher than in liver or serum. Oral administration of the same dose did not produce measurable levels of aglycone BPA in fetal **tissues**. Amniotic fluid consistently contained levels of BPA at or below those in maternal serum. Concentrations of aglycone BPA in **tissues** of neonatal rats decreased with age in a manner consistent with the corresponding circulating levels. Phase II metabolism of BPA increased with fetal age such that near-term fetus was similar to early post-natal rats. These results show that concentrations of aglycone BPA in fetal **tissues** are similar to those in other maternal and neonatal **tissues** and that maternal Phase II metabolism, especially following oral administration, and fetal age are critical in reducing exposures to the fetus.

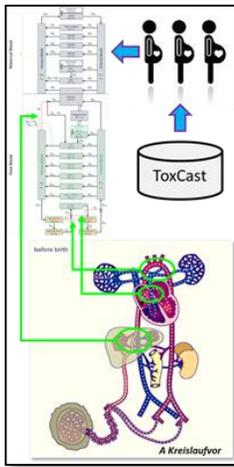
ReadMe | Main | Abstract | Log | Notes | SampleQueries | Landscape



## 5. Progress to date (Year 0.5)

### *Toward HTTK modeling (Aim 2):*

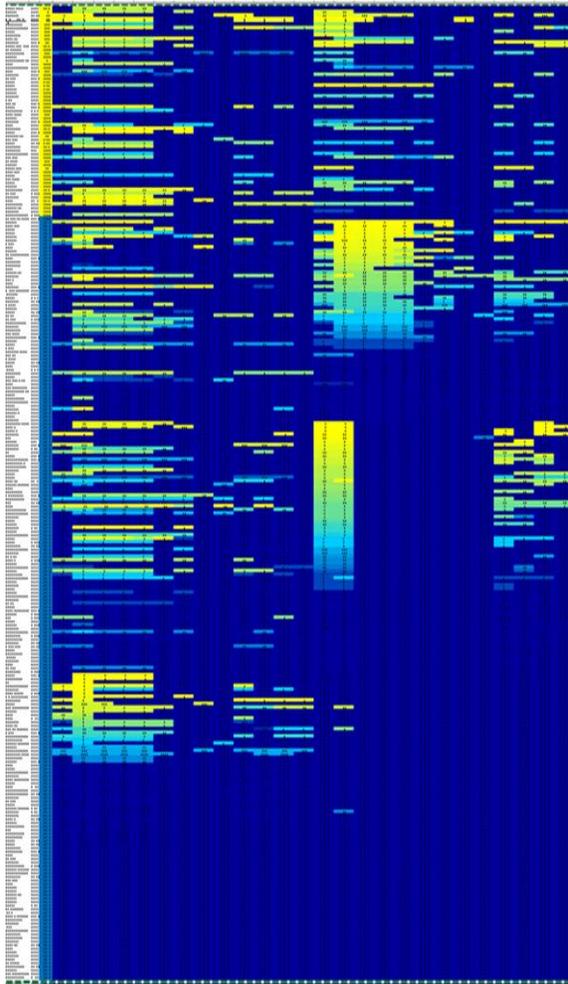
- Since STM outputs an exposure-based hazard prediction, we can translate TI to human pregnancy using a fetal PBPK adaptation of the HTTK model. Annie installed fetal htk package from NCCT and is running initial trials on maternal serum levels for retinoic acid (RA) utilizing parameters provided by NTP.
- RA has a well-defined role in human development and teratogenesis, and was the most potent compound of the 181 positives in ToxCast\_STM. Palmer et al. 2017 just published the bioactivity of several retinoids in the iPSC-version of the assay to address RA homeostasis.
- Annie compiled a list of the compounds (n=271) for which the htk package can be used and a slightly longer list of chemicals (n=349) for which a 3 compartment steady state model can be built. Since the STM prediction has **high specificity**, we can begin to make inferences on margins of exposure.





## 5. Progress to date (Year 0.5)

*Toward computational modeling with the stem cell data (Aim 3):*



Concordance model (rat & rabbit):

- 272 chemicals tested for prenatal DevTox in both species
- Positives = dLEL  $\leq$  125 mg/kg/d for fetal endpoints
- Negatives = no dLEL  $\geq$  1000 mg/kg/day

TP	9
FP	19
FN	15
TN	103
n	146
Sensitivity	0.375
Specificity	0.844
Accuracy	76.7%
Mathew's cc	0.206
F1 score	0.005

Low **sensitivity** raises a question of what is missing biologically or metabolically in the H9 cell platform that animal studies pick up.



## 5. Progress to date (Year 0.5)

### *Toward computational modeling with the stem cell data (Aim 3):*

- Todd Zurlinden (NCCT) built a logistic regression model to mine the strongest positive and negative correlates to 331 enzymatic and receptor signaling assays in the ToxCast\_NVS dataset. Initial machine-learning defined several functional annotation groups and pathways.
- Top sensitive pathways were kinase signaling, some neuroactive GPCRs, and corticotrophs. In contrast, top negative correlations were observed with other neuroactive GPCRs (dopamine, serotonin, endothelins), estrogen signaling, and RAR antagonism.
- These findings point to molecular processes that potentially account for the sensitivity of the STM model and set the stage for AOP integration and mechanistic *in silico* multiscale agent-based models (Aim 4).

# Pathway sensitivity against 337 biochemical targets in ToxCast

NVS ASIDs selected from invitroDB v2 (June, 2017)

Filter NVS+ by AC50 cutoffs (50 nM, 20 nM, 10 nM)

Feature selection with Scikit (Python) classification ANOVA 40<sup>th</sup> percentile

Build logistic regression model based on STM+ (1) and STM- (0) calls

**AD score** (weighted NVS potency and discretized targeted biomarker)

Target	AD Score	AD Score	AD Score	AD Score	AD Score	AD Score	AD Score
GPCR_AdrA1A	Adra1a	adrenergic	0.098	0.091	0.117	0.025	1.551
ENZ_NIN2	MAPK9	Ser/Thr kinase	0.002	1.426	0.097	1.064	1.548
ENZ_H2C2	COM2	Ser/Thr kinase	0.018	1.144	0.449	1.467	1.038
ENZ_NAH1	ABL1	non-receptor tyrosine kinase	0.053	0.038	1.353	0.635	1.000
MP_PBR	Tspo	TSPD	0.391	0.035	1.459	0.305	1.094
GPCR_H3	Hs3	histamine	0.028	0.009	1.321	0.006	1.061
DR_gSDMA_NonSelective	Smor1	2nd messenger	0	0	1.344	0.437	1.124
GPCR_H2A2	TRKA2	2nd messenger	0	0	1.24	0.288	0.915
ENZ_H2MA2_Activator	H2MA2	deacetylase	0.006	0.006	1.121	0.337	0.861
ADME_H2PC239	CYP2C19	CYP inhibition	0.562	0.014	1.31	0.433	0.487
GPCR_INK3	Tao3	neurokinin	0.532	0.04	1.287	0	-0.114
NR_AR	Nr3a2	coreceptor	0.048	0.017	0.26	0.074	0.664
NR_NFR_Antagonist	Nr3a4	ROR	0	0.029	0.722	0.186	0.532
ADME_H2PC341	CYP3A1	CYP inhibition	0.044	0.034	0.001	0.001	0.001
NR_SGR	Nr3c1	coreceptor	0.121	0.54	0.113	0.041	0.091
NR_ICAR_Antagonist	Nr3b3	ROR	0.465	0.141	0.251	0.39	0.541
ENZ_H2A2	JAK2	non-receptor tyrosine kinase	0.265	0.026	0.427	0.238	0.369
ENZ_H2A1	Raf1	Ser/Thr kinase	0.002	0.362	0.229	0.23	0.485
GPCR_gS1ZC	HTR2C	serotonin	0.195	0.013	0.418	0.2	0.372
ENZ_INK2	Nrk2	Ser/Thr kinase	0.074	0.032	0.341	0.228	0.303
GPCR_H2A2A	HTR2A	serotonin	0	0	0.324	0	0.002
NR_AR	Ar	steroid hormone receptor	0	0	0.406	0.16	0.139
ENZ_H2GR	EGFR	receptor tyrosine kinase	0	0.029	0.342	0.002	0.366
ENZ_H2Yn	Fyn	non-receptor tyrosine kinase	0	0.346	0.002	0.366	0.331
GPCR_H1	Hrh1	histamine	0.211	0.03	0.424	0.16	0.116
ENZ_H2MA3_Activator	PAK2	deacetylase	0.148	0.001	0.17	0.17	0.19
ENZ_H2A2	PAK2	Ser/Thr kinase	0.148	0.001	0.17	0.17	0.19
ENZ_H2A2_Activator	TEK	receptor tyrosine kinase	0.148	0.001	0.17	0.17	0.19
ENZ_H2A4	PAK4	Ser/Thr kinase	0	0	0.325	0.319	0
ENZ_H2A72	Akt2	Ser/Thr kinase	0	0	0.001	0.001	0.001
ENZ_H2KA	PRACA	receptor tyrosine kinase	0	-0.007	0.34	0.084	0
ENZ_H2A1	INSR	receptor tyrosine kinase	0.148	0.001	0.17	0.17	0.19
ENZ_H2P133_Activator	PTPN13	non-receptor tyrosine phosphatase	0	0	0.001	0.001	0.001
GPCR_H2A2B	ADRB2	adrenergic	0.148	0.001	0.17	0.17	0.19
ADME_H2PC242	Cyp2a2	CYP inhibition	0	0.029	0	0.219	0.499
NR_H2A2	ROR	ROR	-0.028	0.225	0.017	0	0.18
ENZ_H2R7	SIRT3	deacetylase	0	0	0.472	0.17	0.18
ENZ_H2MAK3_Activator	MAPK3	Ser/Thr kinase	0	0	0.001	0.001	0.001
ADME_H2PC12	CYP1B2	CYP inhibition	0.281	0	0	0.48	0
ENZ_H2P14	PTPN14	non-receptor tyrosine phosphatase	0	-0.016	0.317	0.078	0
ENZ_H2P12	PTPN12	non-receptor tyrosine phosphatase	0	-0.015	0.295	0.073	0
ADME_H2PC26	Cyp1b1	CYP inhibition	0.796	0	0.796	0	0.796
ENZ_H2CAS2_Activator	CASP2	protease	0.713	0.042	0	0.259	0
ADME_H2PC28	CYP2C8	CYP inhibition	0.021	0	0.295	0	0.334
ENZ_H2VGR3	F12	receptor tyrosine kinase	0.062	0.027	0.125	0.007	0
ENZ_H2GR1	FGFR1	receptor tyrosine kinase	0.466	0.025	0.272	0.042	0
ENZ_H2P122	PTPN12	non-receptor tyrosine phosphatase	0.693	-0.03	0.344	-0.113	0
GPCR_gS1T_NonSelective	Hs1a	serotonin	0.008	0	0.266	0	0.008
GPCR_gS1A	Nopa	vaso-peptide	0	-0.027	0	0	0
ENZ_H2M1	MMP1	protease	0	-0.054	0	-0.229	0
ADME_H2PC34	CYP3A4	CYP inhibition	0.078	0	0.303	0	0.322
GPCR_gS1T4	Hs4	serotonin	-0.713	0	-0.168	0	0.667
ENZ_H2R72	SIRT2	deacetylase	0.091	0.04	0	0	-0.105
ENZ_H2MAK2_Activator	RAC1	protease	0	0.02	0.228	0	0
NR_AR	Ar	steroid hormone receptor	0.004	-0.049	0	0.363	0.47
GPCR_AdrA2B	Adra2b	adrenergic	0	0.004	-0.268	-0.11	0
C_C2M1	Ca2m1b	channel	0.001	0	0.274	0.31	0
GPCR_H2R7	Htr7	serotonin	0	0	0.547	0	0
ENZ_H2AKT1	Akt1	Ser/Thr kinase	0	0.009	-0.446	-0.037	-0.039
ENZ_H2R2B	Ser/Thr kinase	Ser/Thr kinase	0	0.4	-0.114	-0.061	-0.229
ENZ_H2S1T1	SIRT1	deacetylase	0.691	-0.049	0	0.315	-0.584
ENZ_H2MAK3	MAPK3	Ser/Thr kinase	0	-0.043	-0.446	-0.121	-0.102
ENZ_H2P111	PTPN11	non-receptor tyrosine phosphatase	0.642	0.04	-0.44	-0.168	0.116
ENZ_H2S1T2_Activator	SIRT2	deacetylase	0	-0.04	-0.375	-0.228	-0.231
GPCR_INK2	TACR2	neuropeptide	0	0.012	-0.607	0	-0.319
GPCR_H2A	Chrm3	muscarinic	0	0.091	0.403	0.14	0
NR_SBR	ESR1	steroid hormone receptor	0	0.122	-0.543	-0.157	-0.067
ENZ_H2AOP	Maoa	mt-oxidase	0.086	-0.053	-0.542	-0.227	-0.226
ENZ_H2S	BDHE	cholinergic	0	0.049	-0.246	-0.217	0
NR_H2RA	Esr1	steroid hormone receptor	0	-0.052	-0.568	-0.102	-0.212
ENZ_H2P11	PTPN11	non-receptor tyrosine phosphatase	0.635	-0.045	-0.623	-0.072	-0.349
ENZ_H2P113_Activator	PTPN13	non-receptor tyrosine phosphatase	0	0.01	0	0.058	0
GPCR_H2A	EDNRA	endothelin	0.677	-0.044	-0.569	-0.269	-0.226
ENZ_H2P119	PTPN19	non-receptor tyrosine phosphatase	0	-0.581	-0.396	-0.241	-0.249
NR_SBR	ESR1	steroid hormone receptor	0	-0.548	-0.025	-0.62	-0.461
GPCR_H2D1	Drd1	dopamine	0	-0.045	0	0	0
ENZ_H2P119_Activator	PTPN19	non-receptor tyrosine phosphatase	0.071	-0.08	-0.571	-0.396	-0.355
ENZ_H2MAK4_Activator	MAP4	Ser/Thr kinase	0	-0.603	-0.371	-0.541	-0.541
ENZ_H2PDE4A1	PDE4	2nd messenger	0.803	-0.089	-0.563	-0.396	-0.389
NR_SGR	MAPK9	steroid hormone receptor	0.817	0.046	-0.697	-0.10	0
ENZ_H2M1P	MMP9	protease	0.078	-0.045	-0.368	-0.37	-0.621
NR_H2AR_Antagonist	RARA	retinoic acid	-0.079	-0.007	-0.745	-0.393	-0.462
GPCR_H2D2a	Drd2	dopamine	0	0.036	0.046	0	0
NR_H2R2	ROR	ROR	-1.245	-0.717	-0.707	0	0
ADME_H2PC246	CYP24A	CYP inhibition	0.151	-0.067	-1.46	-0.311	-0.507

**NIH\_DAVID annotation:**  
functional clusters

**Strongest (-) enrichment clusters**

DRD2	dopamine receptor D2(DRD2)
DRD1	dopamine receptor D1(DRD1)
Ca2m1b	calcium voltage-gated channel subunit alpha1 B(CACNA1B)
Htr1a	5-hydroxytryptamine receptor 1A(HTR1A)
TACR2	tachykinin receptor 2(TACR2)
Adra1b	adrenoceptor alpha 1B(ADRA1B)
HTR7	5-hydroxytryptamine receptor 7(HTR7)
CHRM3	cholinergic receptor muscarinic 3(CHRM3)
Htr4	5-hydroxytryptamine receptor 4(HTR4)
BACE1	beta-secretase 1(BACE1)
EDNRA	endothelin receptor type A(EDNRA)
ESR1	estrogen receptor 1(ESR1)
RARA	retinoic acid receptor alpha(RARA)
PGR	progesterone receptor(PGR)
Ar	androgen receptor(AR)
NR1I2	nuclear receptor subfamily 1 group I member 2(NR1I2)



## 6. Science challenges

- The **high specificity** of the STM assay provides confidence in predictivity; however, it's **low sensitivity** raises the science challenge of making the assay more health protective.
  - Understanding the dynamics that potentially account for sensitivity of the STM model set the stage for AOP integration and mechanistic *in silico* agent-based models (Aim 4).
  - Reducing uncertainty via computational synthesis and integration to stratify connections between the most relevant pathways and link DSSTox structural descriptors with the biology.
  - This cycles back to the principles of teratogenesis (initiating mechanisms, dosimetry & bioavailability, stage vulnerability, chemistry, and genetic susceptibility).
- Next steps: (a) NCCT will continue pushing the ToxCast\_STM dataset forward on H9 cells and could expand chemicals to meet needs for amended TSCA; (b) NTP will pilot the iPSC version of the assay on 81 chemicals and focus on data processing and integration; (c) NCTR will continue with exposure reconstruction on RA as a case study for the fetal htk.