



Advisory Group on Endocrine Disruptors Testing and Assessment
(EDTA AG) - Virtual Meeting, May 26, 2021

RETINOID SIGNALING IN SKELETAL DEVELOPMENT: *SCOPING THE SYSTEM FOR PREDICTIVE TOXICOLOGY*

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



Evaluating prenatal developmental toxicity

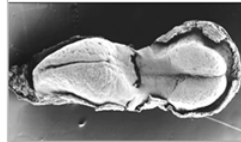


"The first trimester is the most crucial to your baby's development. During this period, your baby's body structure and organ systems develop."

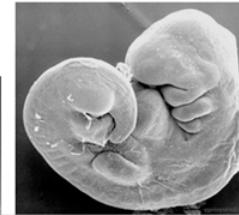
www.ucsfhealth.org

TIMELINE OF THE HUMAN EMBRYONIC PERIOD

Week 3
(Carnegie Stage 8)



Week 4
(Carnegie Stage 13)



Week 8
(Carnegie Stage 20)



peak sensitivity (3rd – 8th wk)

Embryonic Period

T1

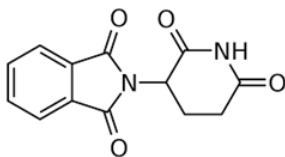
Fetal Period

T2

Fetal Period

T3

OECD TG 414
OPPTS 870.3700

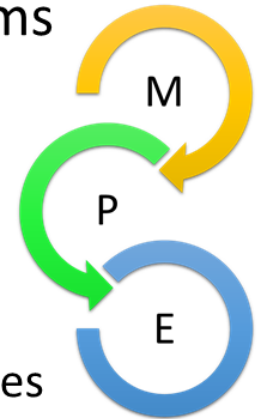


Adverse Birth Outcomes (CDC)

- neonatal mortality (1-2%)
- malformations (3-4% live births)
- premature births (10%)
- low birth weight (11%)
- functional deficits (17% children)

Complex Systems

- gene networks
- multiscale
- autopoiesis
- canalization
- temporality
- state trajectories
- and more ...





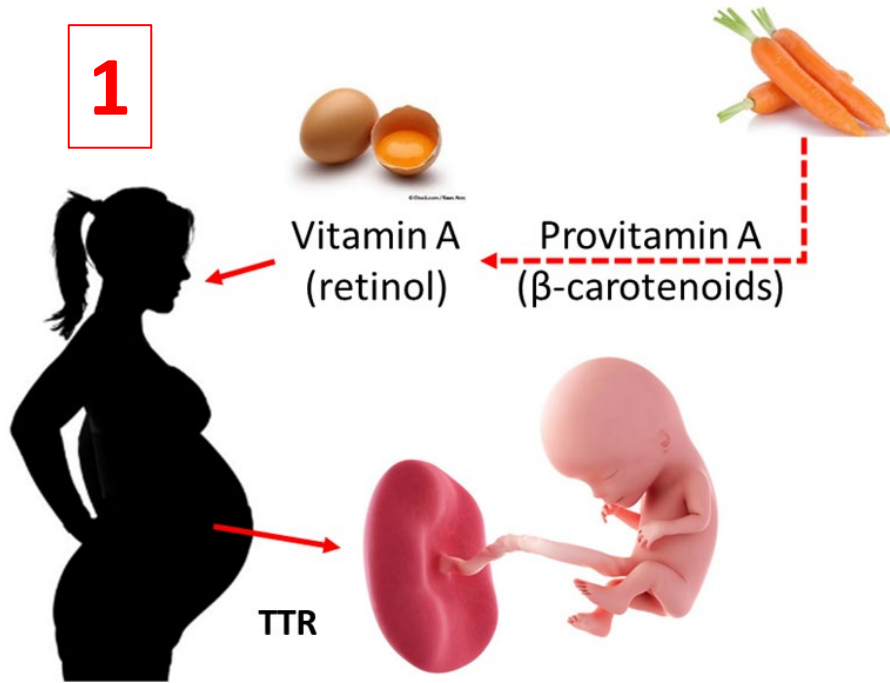
1. Key messages from DRP 4.97 (Annex-B)

- The retinoid pathway has parallels with endocrine signaling (EATS), but its many unique functions are essential to morphogenesis, growth and differentiation of the embryo.
- ATRA signaling collaborates with some of the most powerful morphogenetic signals known to the embryo (FGF, BMP, SHH, WNT, ...).
- Threshold ATRA responses are transduced by intracellular transport, liganding of nuclear RAR/RXR receptors bound to RARE genomic sequences, and recruitment of NCoA.
- ATRA-dependent gene expression is a molecular determinant of early growth, patterning and differentiation of the fetal skeleton (facial, vertebral, appendicular).

KEY POINT: Although it is clear that disruption of endogenous retinoid signaling has adverse consequences to skeletal development, from a broad DevTox regulatory context it is not clear which (if any) apical outcomes are attributable exclusively to retinoid-related mechanism(s).



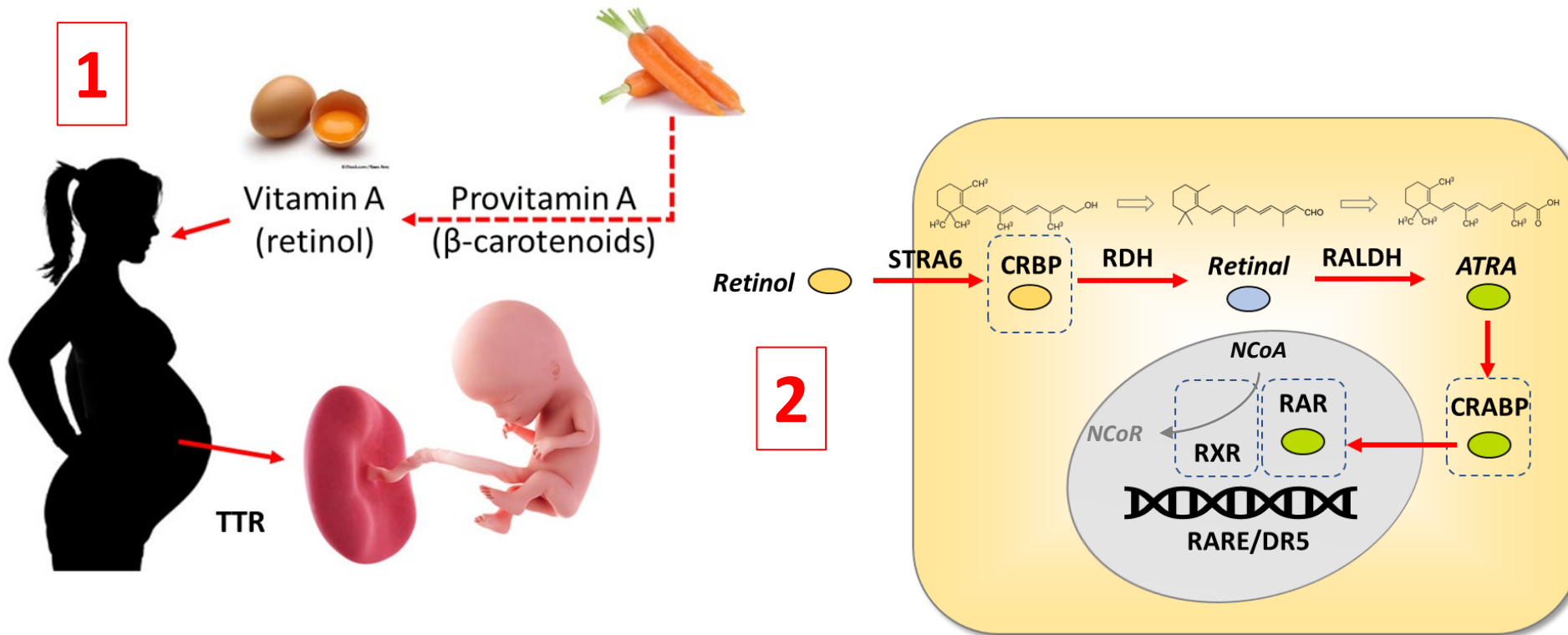
The retinoid pathway and morphogenetic signaling



- 1.** Retinol (vitamin A) is ingested by the mother and stored in her liver; it circulates to target tissues bound to serum transporters such as transthyretin (TTR).



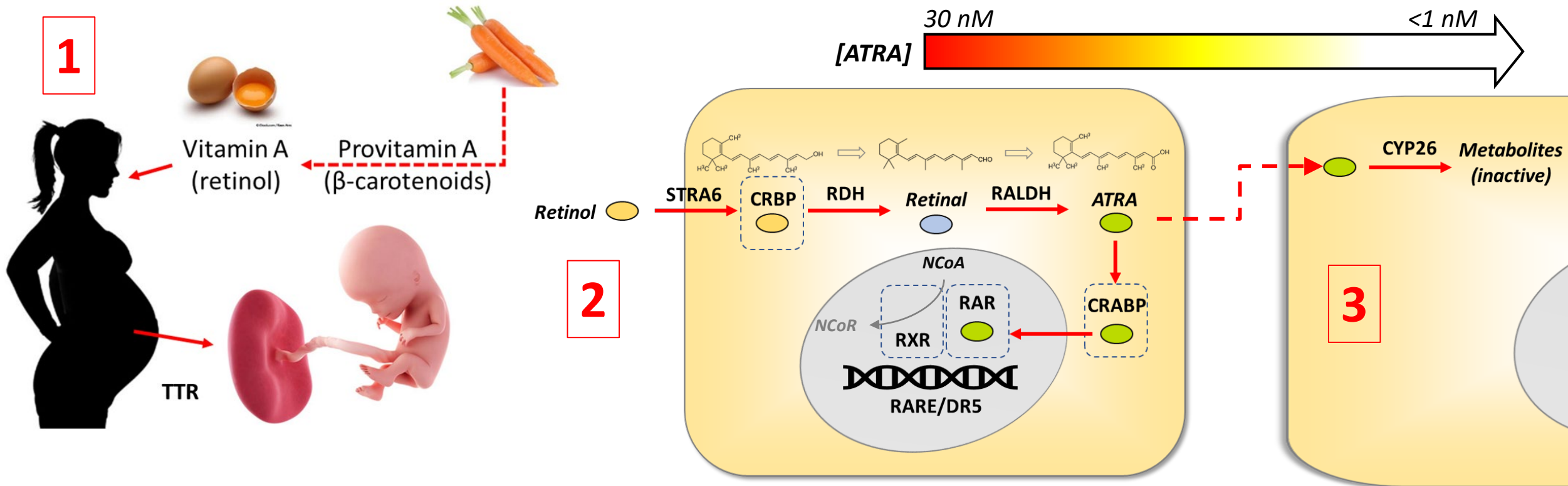
The retinoid pathway and morphogenetic signaling



- 2.** In the embryo, retinol is enzymatically converted to all-trans retinoic acid (ATRA). This capacity is acquired during gastrulation and continues through organogenesis.



The retinoid pathway and morphogenetic signaling



3. ATRA is fashioned into biochemical gradients by the cell-specific expression of enzymes for its biosynthesis (RDH10, RALDH2) and breakdown (CYP26a/b/c).



The retinoid pathway and morphogenetic signaling



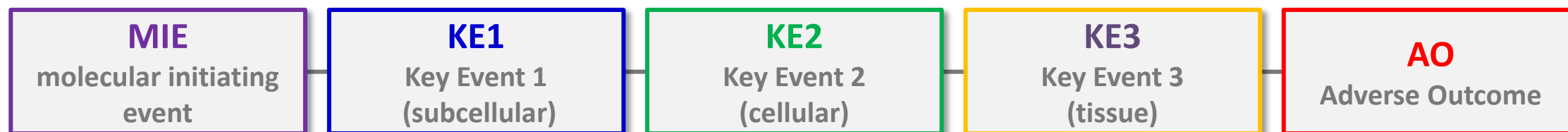
Dosimetric	Conc.	Indication	Reference
baseline ATRA (5 somite zebrafish embryo)	< 1 nM	non-morphogenetic	(Shimozono, Iimura et al. 2013)
maternal serum (animal study)	1.7 nM	non-teratogenic	(Daston, Beyer et al. 2014)
devTOX ^{qp} assay (pluripotent hESC)	3.0 nM	teratogenic threshold	(Zurlinden, Saili et al. 2020)
normal plasma concentration	5.0 nM	physiological (adult)	(Napoli, Posch et al. 1991)
axial gradient (5 somite zebrafish embryo)	6.0 nM	morphogenetic signal	(Shimozono, Iimura et al. 2013)
endodermal differentiation (h-iPSC)	17 nM	toxicological tipping point	(Saili, Antonijevic et al. 2019)
devTOX ^{qp} assay (pluripotent h-iPSC)	19 nM	DevTox potential	(Palmer, Smith et al. 2017)
genetic perturbation (mouse)	30 nM	altered homeostasis	(Helms, Thaller et al. 1994)
maternal serum (animal study)	30 nM	teratogenic potential	(Daston, Beyer et al. 2014)
limb-bud (GD 10.5 mouse embryo)	30 nM	physiological (embryo)	(Horton and Maden 1995)
pharmacological kinetics	1,000 nM	efficacious (therapeutic)	(Helms, Thaller et al. 1994)
limb-bud (GD 11 mouse embryo)	1,500 nM	weakly teratogenic dose	(Satre and Kochhar 1989)
limb-bud (GD 10.5 mouse embryo)	12,500 nM	fully teratogenic dose	(Horton and Maden 1995)

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4. It is not clear which (if any) apical skeletal outcomes in chemically-induced ‘retinoid embryopathy’ are attributable exclusively to retinoid-related mechanism(s).



2. Postulated AOPs and status of development

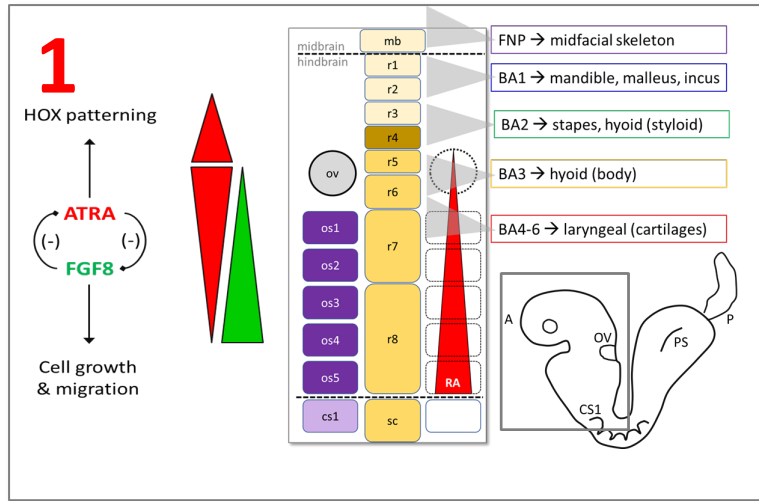


REGION	MIE	KE1	KE2	KE3	KE4	KE5	AO
Anterior Neural Tube	Inhibition of CYP26A1 enzymatic activity	Local increase in endogenous ATRA levels	Hyperactivation of the RAR/RXR heterodimer	Repression of <i>Fgf8</i> limits FGF8 signaling	Mis-specification of CNC cell fate and behavior	Maxillary arch dysplasia alters palatal outgrowth	Cleft palate
Paraxial Mesoderm	Reduction in RDH/RALDH2 activity	Local decrease in endogenous ATRA levels	Hypoactivation of the RAR/RXR heterodimer	Overextension of FGF8 signaling	Disruption of the periodic somitic wavefront	Altered somite number, shape, and alignment	Hemivertebra
Limb-Bud Mesoderm	Hyperactivation of the RAR/RXR heterodimer	Underextension FGF8 signaling from the AER	Dysregulation of <i>Meis1/2</i> and <i>Hox</i> gene expression	Proximalization of the limb-bud mesenchyme	Mis-specification of precartilage blastema	Malformed cartilaginous bone rudiment	Phocomelia

Postulated AOPs for each skeletal domain based on literature for dietary, genetic or chemical disruption of endogenous ATRA signaling during embryogenesis.

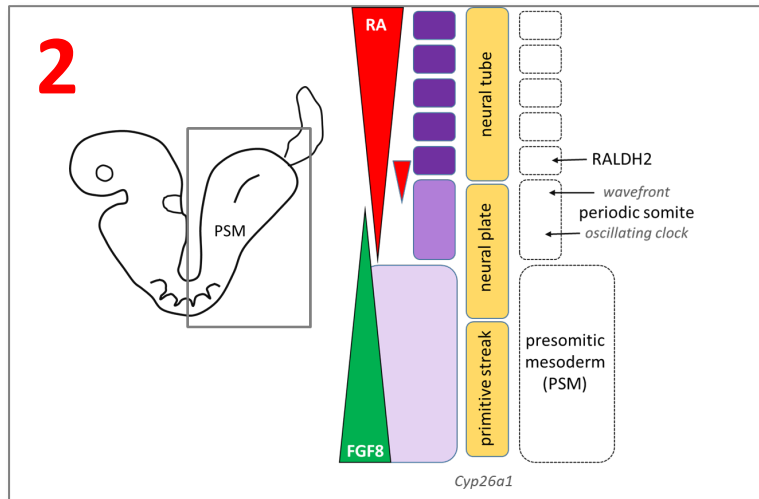


Domain-specific AOPs (provisional examples)



1. Facial skeleton: positional information of premigratory neural crest cells destined for branchial arches (5- to 11 somite stage).

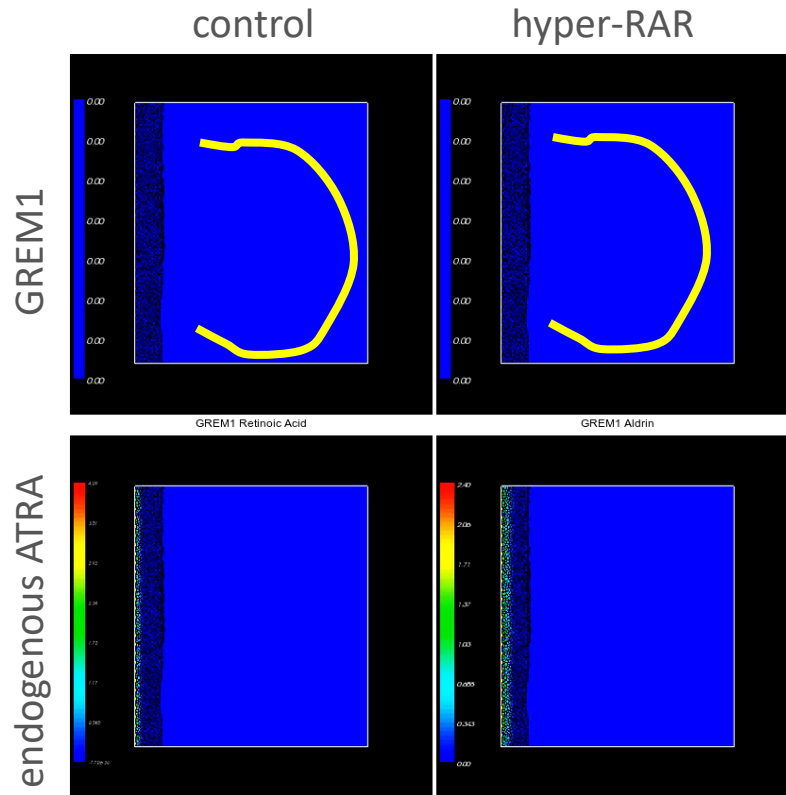
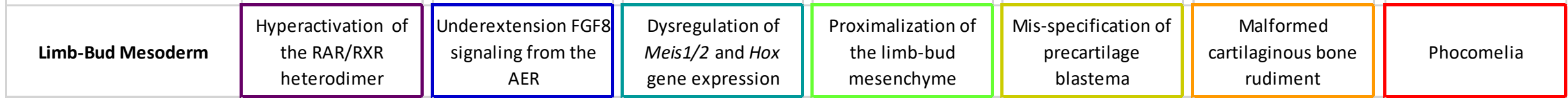
2. Vertebral skeleton: size, registration, and specification of somites giving rise to individual vertebrae/ribs (0- to 36 somite stage).



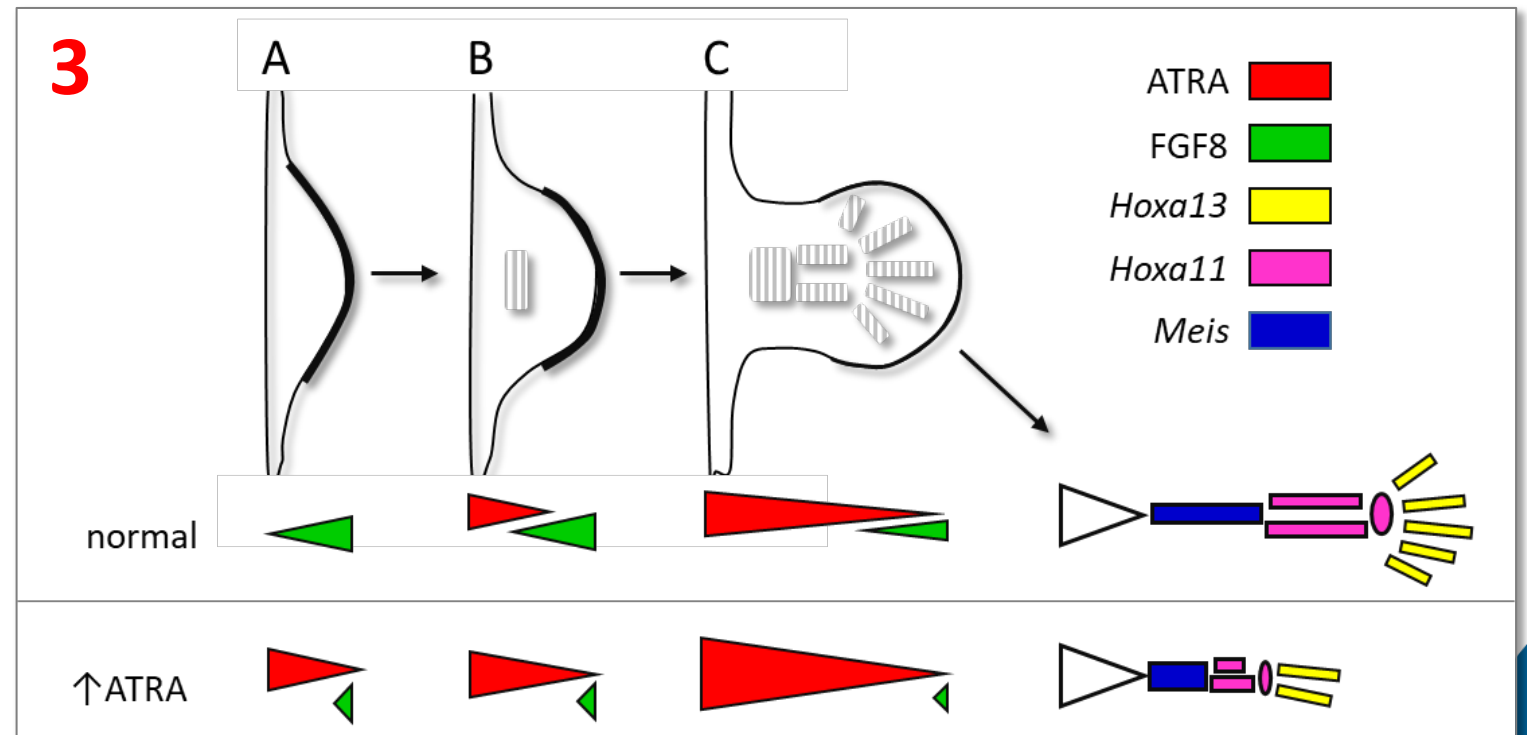
3. Appendicular skeleton: limb-bud initiation, outgrowth, patterning, and differentiation (12- to 36+ somite stage).



Example: phocomelia due to hyperactivity of RAR



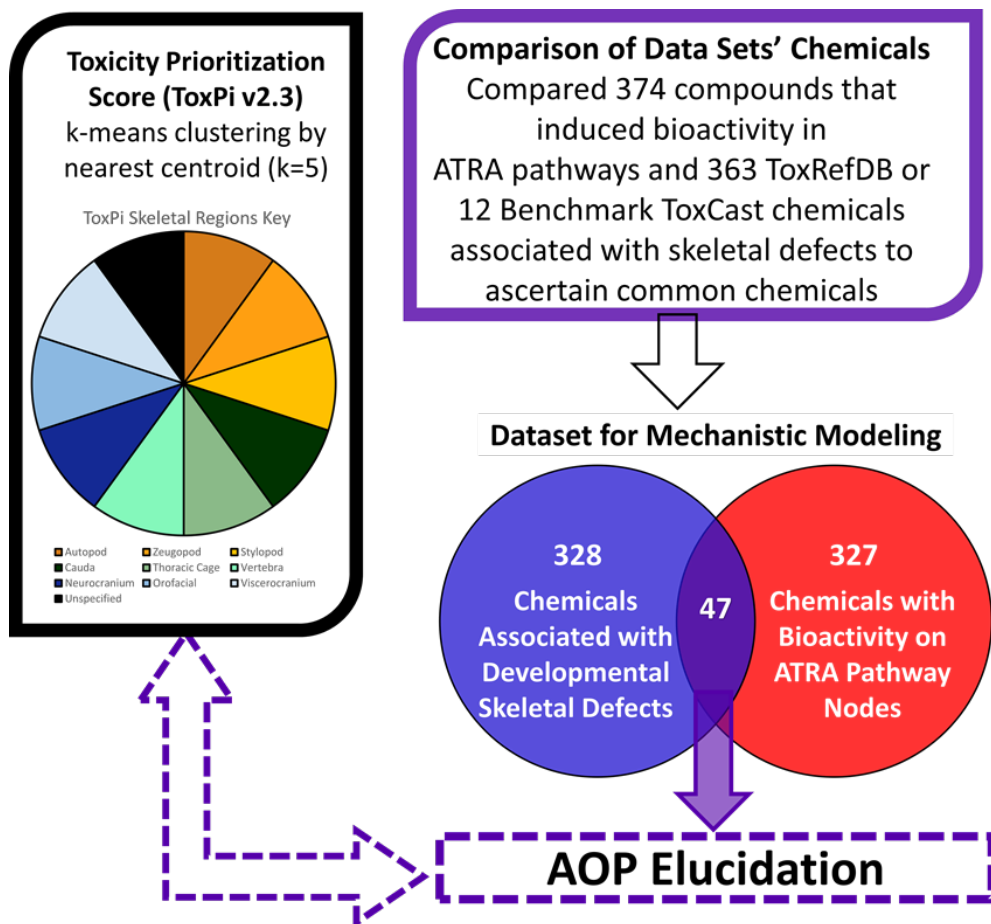
Systems model, work in progress



Based on Uzkudunet al. (2015) Mol Syst Biol



Broadening the landscape



- 2946 developmental studies in EPA's ToxRefDB with an adverse skeletal outcome (328 chemicals x 57,198 features).
- ToxPi slices annotated per chemical by composite and potency (mg/kg dosage) across these anatomical domains:

Cranium

- viscerocranium (oral, face)
- neurocranium (base, vault)

Axial skeleton

- vertebra (cervical, thoracic, lumbar)
- thoracic cage (ribs, sternum)
- cauda (tail)

Appendicular skeleton

- stylopod, zeugopod, autopod

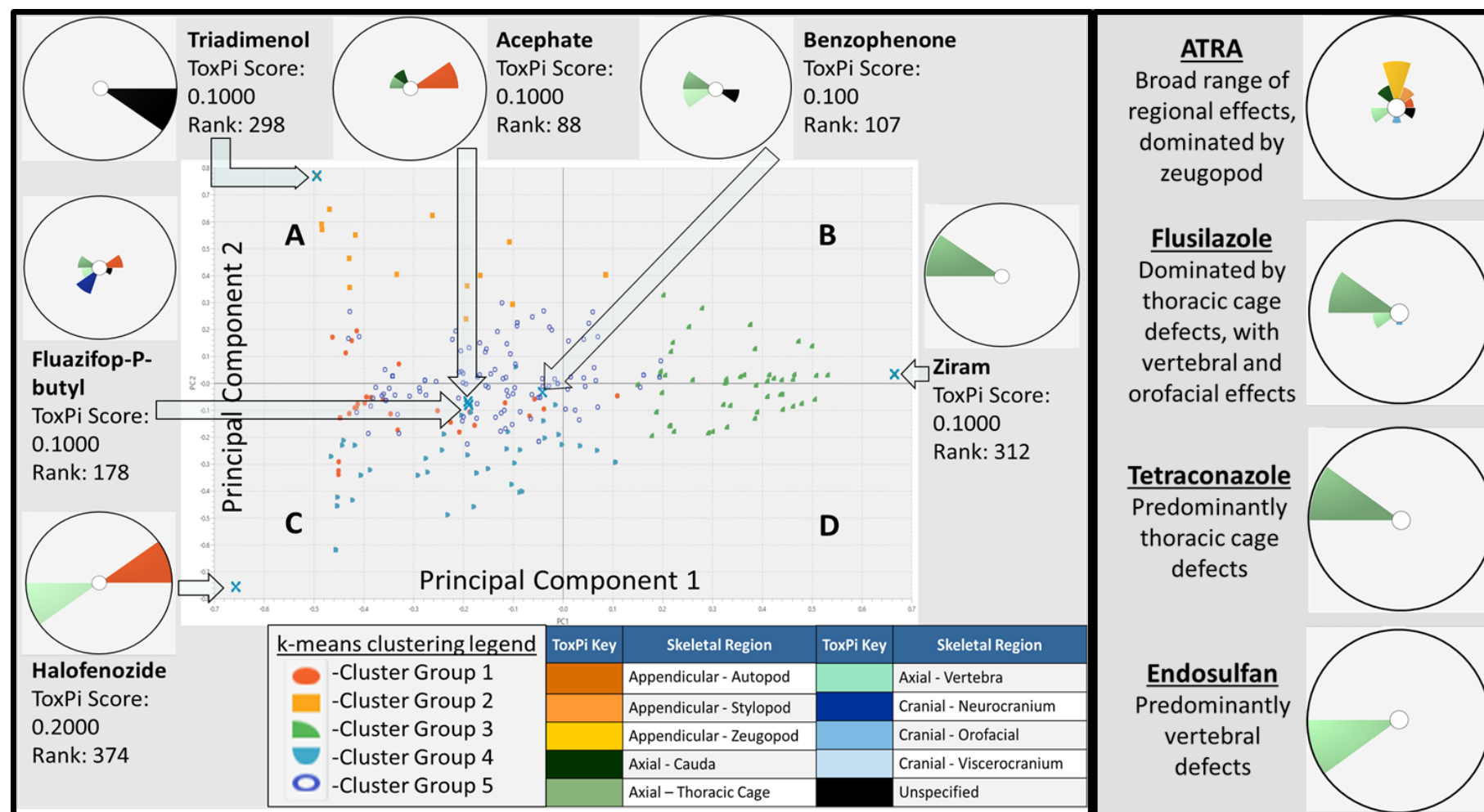
Other - insufficient detail for mapping





Chemicals (375) clustered by ToxPi phenotype (k=5)

- **Phenotyping:** *are defects of the fetal skeleton consistent with AOPs linked to ATRA pathway?*
- **Chemotyping:** *are distinct structural features shown by chemicals in each phenotype cluster?*





3. Candidate assays/endpoints and value for standardization



- New Approach Methods (NAMs): *in vitro* data and *in silico* models for hazard identification with low reliance on animal testing.
- An important consideration is building NAMs for early lifestages, reflecting the best knowledge of human developmental biology.
- Integrated Approaches to Testing and Assessment (IATA): integrating information with data generated with new testing strategies.
- The ATRA system provides an excellent opportunity to establish reliability of NAMs and defining IATA case studies.



ToxCast class distribution for retinoid target assays

1. Retinoids (ATRA, retinol, bexarotene).

2. Triazoles (CYP inhibition – CYP1A1 as a surrogate to CYP26a/b/c).

3. persistent organic pollutants (selective RAR activation).

4. tert-butyl and organotin compounds (selective RXR activation)

Other: mitochondrial disrupters, anticonvulsants (VPA), flame retardants (PBDEs).

DSSTOXID	PREFERRED_NAME	CYP1A1 (72)	RA Ra (65)	RARb (17)	RARg (49)	RXRa (69)	RXRb (299)	RXRg (0)	DR5 (250)
DTXSID7021239	all-trans-Retinoic acid	1.317	NA	NA	NA	NA	1.036	NA	0.006
DTXSID1040619	Bexarotene	NA	7.539	NA	2.655	0.009	0.009	NA	0.014
DTXSID3023536	Retinol	NA	0.076	NA	0.227	2.142	0.464	NA	0.197
DTXSID1020807	2-Mercaptobenzothiazole	0.164	NA	NA	NA	NA	NA	NA	NA
DTXSID2040363	Diniconazole	0.674	NA	NA	NA	NA	NA	NA	NA
DTXSID0032655	Triticonazole	0.793	NA	NA	NA	NA	NA	NA	16.741
DTXSID8024151	Imazalil	1.413	0.908	NA	NA	NA	NA	NA	5.888
DTXSID4032372	Difenoconazole	1.459	NA	NA	NA	NA	NA	NA	2.124
DTXSID3023897	Triademifon	2.085	41.462	NA	NA	NA	NA	NA	11.223
DTXSID7029871	Clotrimazole	2.306	NA	NA	NA	NA	NA	NA	NA
DTXSID3024235	Flusilazole	3.704	8.155	NA	NA	NA	NA	NA	7.718
DTXSID2032500	Triflumizole	4.134	1.453	NA	NA	NA	NA	NA	0.298
DTXSID0021337	Thiabendazole	4.721	NA	NA	NA	NA	NA	NA	NA
DTXSID8024280	Propiconazole	9.010	23.801	NA	NA	NA	NA	NA	6.253
DTXSID9020453	Dieldrin	NA	0.770	NA	1.679	NA	22.531	NA	0.579
DTXSID9037539	Endosulfan I	NA	1.384	NA	NA	NA	NA	NA	1.827
DTXSID6020561	Endrin	NA	NA	1.606	1.698	NA	24.982	NA	0.806
DTXSID1020560	Endosulfan	NA	NA	NA	NA	NA	NA	NA	0.894
DTXSID7020267	Chlordane	NA	NA	NA	6.878	71.470	21.422	NA	1.784
DTXSID7042065	Isodrin	NA	NA	NA	1.077	NA	NA	NA	2.111
DTXSID8020040	Aldrin	NA	NA	NA	0.912	NA	7.167	NA	3.085
DTXSID3042500	Triphenyltin fluoride	NA	NA	NA	NA	0.004	0.001	NA	0.655
DTXSID5034981	Tributyltin benzoate	NA	NA	NA	NA	0.005	0.036	NA	0.023
DTXSID9044796	(Acryloyloxy)(tributyl)stannane	NA	NA	NA	NA	0.015	0.026	NA	0.022
DTXSID2040733	Triphenyltin chloride	NA	NA	NA	NA	0.081	0.037	NA	0.356
DTXSID9035204	Tributyltin methacrylate	NA	NA	NA	NA	0.147	0.025	NA	0.005
DTXSID3027403	Tributyltin chloride	NA	NA	NA	NA	0.176	0.078	NA	0.003
DTXSID4022153	Tetrabutyltin	NA	NA	NA	NA	0.741	0.033	NA	0.279
DTXSID1021409	Triphenyltin hydroxide	NA	NA	NA	NA	NA	0.013	NA	NA
DTXSID9040712	Triethyltin bromide	NA	NA	NA	NA	4.029	0.252	NA	NA



Baker et al. (manuscript in preparation)

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Direct assays for modeling the ATRA pathway

Assays that Detect Potential RA-related MIEs			
ChEMBL Assay ID	Chem. Count	Assay description	PMID
ChEMBL entries for retinoic acid binding proteins (CRABP) related assays			
ChEMBL Assay 1252	10	Inhibition of CRABP1 with Cytoplasmic retinoic acid binding protein at 100-fold excess (ligand)	6801513
ChEMBL Assay 1253	7	Inhibition of CRABP1 with Cytoplasmic retinoic acid binding protein at 100-fold excess (ligand)	7798895
ChEMBL Assay 1254	7	Inhibition of CRABP1 with Cytoplasmic retinoic acid binding protein at 100-fold excess (ligand)	7798895
ChEMBL Assay 1255	9	Inhibition of CRABP1 with Cytoplasmic retinoic acid binding protein at 100-fold excess (ligand)	6801513
ChEMBL entries for retinol dehydrogenase (RDH) related assays			
ChEMBL Assay 1310	10	Inhibition of human retinol dehydrogenase (RDH) activity	9172895
ChEMBL Assay 1311	10	Inhibition of human retinol dehydrogenase (RDH) activity	7005009
ChEMBL entries for ALDH1A related assays			
ChEMBL Assay 1312	37	Inhibition of full length recombinant human ALDH1A1 expressed in Sf9 cells using 10-100 nM of test compound. ALDH1A1 activity was measured by measuring NAD(P)H levels at 10 min post incubation for 2 min. Inhibitors were added at the end of the incubation period. ALDH1A1 activity was measured by measuring NAD(P)H levels at 10 min post incubation for 2 min. Inhibitors were added at the end of the incubation period.	28219111
ChEMBL Assay 1313	22	Inhibition of human ALDH1A1 using recombinant ALDH1A1 as substrate. Inhibitors were added at 10 min post incubation for 2 min. Inhibitors were added at the end of the incubation period.	24660056
ChEMBL entries for CYP2B related assays			
ChEMBL Assay 1314	24	Inhibition of recombinant human CYP2B1 expressed in Sf9 cells using 10-100 nM of test compound. CYP2B1 activity was measured by measuring NAD(P)H levels at 10 min post incubation for 2 min. Inhibitors were added at the end of the incubation period.	26018320
ChEMBL Assay 1315	16	Inhibition of CYP2B1 in ATRA-induced human HL60 cells. Recombinant CYP2B1 activity was measured by measuring NAD(P)H levels at 10 min post incubation for 2 min. Inhibitors were added at the end of the incubation period.	26307100
ChEMBL Assay 1316	18	Inhibition of human CYP2B1 activity using recombinant CYP2B1 as substrate. Inhibitors were added at 10 min post incubation for 2 min. Inhibitors were added at the end of the incubation period.	21836326
ChEMBL Assay 1317	31	Inhibition of human CYP2B1 activity using recombinant CYP2B1 as substrate. Inhibitors were added at 10 min post incubation for 2 min. Inhibitors were added at the end of the incubation period.	15745819

ChEMBL has limited data on 12 upstream assays (ATRA metabolism) for drug-like compounds

ToxCast has target-specific data on 11 assays for <2K Ph-I/II chemicals.

Tox21 has 2 assays on the integrated retinol pathway (LOPAC and Tox21 10K chem) libraries.

ToxCast/Tox21 Assay ID	Chem. Active	Assay description	PMID
ToxCast assays for CYP3A1			
WIS_A001_CYP3A1*	72	Using a type of enzyme reporter, loss-of-signal activity can be used to understand changes in the enzymatic activity as they relate to the gene CYP3A1	
WIS_A001_CYP3A1*	6	Using a type of enzyme reporter, loss-of-signal activity can be used to understand changes in the enzymatic activity as they relate to the gene CYP3A1	
ToxCast assays for RARalpha, RARbeta, and RXRgamma			
ATC_DR5_C15_u*	56	Measures of mRNA for gain-of-signal activity can be used to understand the reporter gene as the transcription factor level as they relate to the gene RARalpha and RARbeta	
ATC_DR5_C15_u*	80	Measures of mRNA for gain-of-signal activity can be used to understand the reporter gene as the transcription factor level as they relate to the gene RARalpha and RARbeta	
ATC_DR5_C15_u*	26	Measures of mRNA for gain-of-signal activity can be used to understand the reporter gene as the transcription factor level as they relate to the gene RARalpha and RARbeta	
ATC_DR5_C15_u*	63	Measures of mRNA for gain-of-signal activity can be used to understand the reporter gene as the transcription factor level as they relate to the gene RARalpha and RARbeta	
WIS_M001_Antagonist*	61	Using a type of binding reporter, loss-of-signal activity can be used to understand changes in the binding as they relate to the gene RARalpha	
WIS_M001_Antagonist*	62	Using a type of binding reporter, loss-of-signal activity can be used to understand changes in the binding as they relate to the gene RARalpha	
Tox21 assays for reporter assays RARalpha			
TOX21_RAR_LUC_Antagonist*	3015	Using a type of luciferase reporter, loss-of-signal activity can be used to understand changes in the reporter gene as they relate to the gene RARalpha	
TOX21_RAR_LUC_Antagonist*	9815	Using a type of luciferase reporter, loss-of-signal activity can be used to understand changes in the reporter gene as they relate to the gene RARalpha	
ToxCast assays for RARalpha, RXRbeta, and RXRgamma			
ATC_DR5_C15_u*	19	Measures of mRNA for gain-of-signal activity can be used to understand the reporter gene as the transcription factor level as they relate to the gene RARalpha	

ATRA processes that can be modeled

Pregnancy

- exposure (diet, pharma, chemicals)
- hepatic mobilization (retinol)
- transplacental kinetics (RBP)

Microphysiology

- bioactivation (RDH, RALDH2)
- molecular transporter (Stra6)
- metabolism (CYP26a/b/c)

Cell signaling

- endogenous ATRA gradients
- ATRA kinematics (retinoid dosimetry)
- nuclear delivery (CRABP)

Nuclear signaling

- RAR/RXR liganding
- NCOR – NCOA switching
- DR5 mediated transactivation



4. Connections with other on-going projects at OECD and elsewhere

Knowledgebase
(skeletal development)

AOP-WIKI
(limb defects)

HTS-based signatures
(ToxPi classifier)

HTS data analysis
(ToxCast/Tox21/ChEMBL)

Pregnancy IVIVE models
(targeted case studies)

Performance-based prediction
(ATRA pathway in Devtox)

Morphoregulatory simulation
(Limb ABM)

1. CPP 13 Predictive Toxicology of the Retinoid Signaling Pathway.

US EPA, CCTE

Nancy Baker, Leidos
Richard Judson
Thomas Knudsen
Jocelyn Pierro
Ann Richard
Laura Taylor

NCTR/FDA

Annie Lumen

NTP/NIEHS

Nicole Kleinstreuer,

NIH/NCATS

Srilatha Sakamuru
Menghang Xia

OECD

Patience Browne



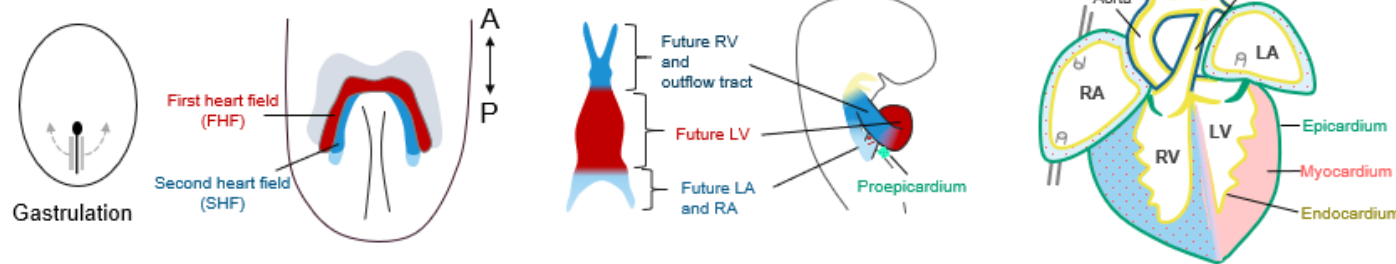
Aim: mine HTS bioactivity profiles for retinoid transporters, metabolism, receptors, and responsive pathways that can be formally integrated with embryological knowledge on the embryo-fetal skeletal system as a potential IATA case study.



4. Connections with other on-going projects at OECD and elsewhere

2. Retinoid Signaling and Cardiovascular Development.

	Cardiac Precursors	Cardiac Crescent	Linear Heart Tube	Cardiac Looping	4-Chambered Heart
Mouse:	6.5	E7.5	E8.0	E8.5	E14.5
Human:	CS6	CS7	CS8	CS8	CS20



Roles for Retinoic Acid Signaling	- Restricting heart field size - Patterning the SHF	Addition of SHF cells to growing heart tube	- Left-right asymmetry - Outflow tract development	- Myocardial expansion - Formation of fibroblast and vascular smooth muscle cell populations
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US EPA

Nancy Baker
Rachel Brunner, OPP
Sid Hunter
Thomas Knudsen
Jocelyn Pierro
John Rogers
Other experts (TBD)



Aim: comprehensive review of the scientific literature on endogenous ATRA signaling in the regulation of cardiovascular (CV) development, functioning and health focusing on the heart and great vessels. To be a new annex in DRP 4.79.



4. Connections with other on-going projects at OECD and elsewhere

3. DRP on Stem Cell Assays for Developmental Toxicity (EST).

- **1,250 chemicals:**
18 studies tested ≥ 10 chems, 174 studies 1-9 chems.
- **Most overlap:** ATRA (17 studies), and the most potent of 1065 ToxCast chemicals.

EST team

Nancy Baker - Leidos
George Daston – Procter & Gamble Co.
Burkhard Flick – BASF (Berlin)
Michio Fujiwara – Astellas Pharma Inc. (Japan)
Thomas Knudsen – USEPA
Hajime Kojima – NIEHS/JaCVAM (Japan)
Aldert Piersma – RIVM (Netherlands)
Horst Spielmann – Berlin (retired)
Noriyuki Suzuki – Sumitomo Chemical Co. (Japan)
Katya Tsaion – Johns Hopkins University



Aim: systematic scoping review on pluripotent stem cell assay modalities and applications for predictive developmental toxicity: chemical and biological domains, readout endpoints, standardized protocols, reproducibility and predictive performance.



Outlook

- Harmonized protocol(s) assessing the retinoid pathway would be useful for developmental toxicity hazard prediction in AOPs leading to fetal skeletal defects.
- Translatability will depend on the integration of *in vitro* data with embryological knowledge (*in vivo*) using multifaceted computational tools, approaches, and models.
- Due to complexity of the retinoid pathway and related networks, computer modeling will be useful for quantitative simulation for animal-free hazard evaluation.