

Computational Systems Biology: translational tools for data integration and modeling

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

Conflict of Interest Statement

The author declares no conflict of interest.

Abbreviations

- ToxCast** EPA's *Toxicity Forecaster* generates data and predictive models on thousands of chemicals of interest to the as part of the Tox21 federal collaboration. <https://www.epa.gov/chemical-research/toxicity-forecasting>
- HTS** ToxCast/Tox21 use *high-throughput screening* and computational modeling to rank chemicals based on their *in vitro* bioactivity profiles. <https://www.epa.gov/chemical-research/toxicity-forecaster-toxcasttm-data>.
- ToxRefDB** EPA's *Toxicity Reference Database* provides results from across thousands of animal toxicity studies effects and endpoints. <https://www.epa.gov/chemical-research/toxicity-forecaster-toxcasttm-data>
- AOP** *Adverse outcome pathway* is a formal representation of steps linking a molecular initiating event (MIE) to an adverse outcome (AO) relevant to risk assessment. <https://aopwiki.org/aops/15>
- ABM** *Agent-based model* is an *in silico* approach to simulate the dynamics of autonomous agents (eg, cells) that individually assess their situation and make decisions based on encoded rules and constraints.
- CC3D** *CompuCell3D* is an open-source simulation environment to execute cellular ABMs with a view to assessing critical state transitions and emergent properties in a self-organizing system. <http://www.compuccell3d.org/>

Problem statement

- Automated *in vitro* assays enable high-throughput screening (HTS) to ‘decode the toxicological blueprint of active substances’ that interact with pregnancy.
- Vast HTS data (ToxCast/Tox21) in hand [<https://comptox.epa.gov/dashboard>], the need arises for predictive models of developmental toxicity.
- Key challenge: model ‘critical phenomena’ in self-organizing embryonic systems that compute with complex genetic circuits and multi-cellular networks.

Outline

Two examples will demonstrate the predictive power of ToxCast HTS data when integrated with developmentally-conserved cell signaling pathways:

1. Profiling the ToxCast library with a pluripotent human (H9) embryonic stem cell assay



Objective: increase the diversity and relevance of assays in ToxCast that can be used to profile chemicals for potential adverse effects on human embryonic development.

2. Translating biomolecular lesion(s) into quantitative phenotypes for predictive developmental toxicology



Objective: build and test computer models of complex tissues that advance critical phenomena (specificity, canalization, plasticity) for quantitative prediction for virtual screening and *in silico* testing.

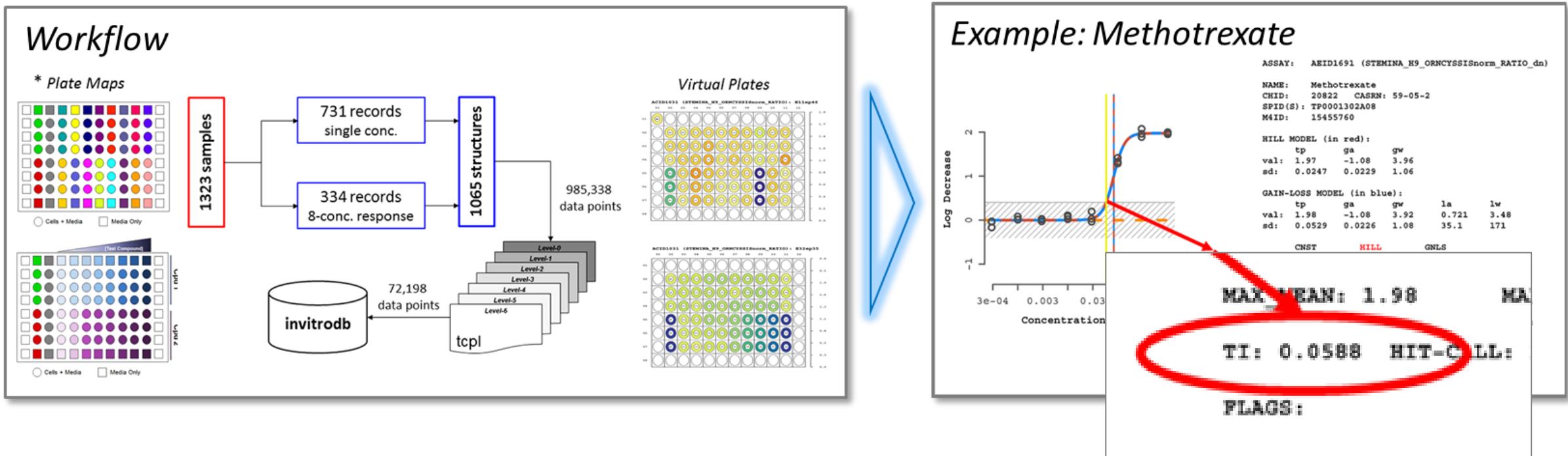
1. Profiling the ToxCast library with a pluripotent human (H9) embryonic stem cell assay



Objective: increase the diversity and relevance of assays in ToxCast that can be used to profile chemicals for potential adverse effects on human embryonic development.

ToxCast_STM assay

- devTOX^{qP} assay from Stemina Biomarker Discovery [Palmer et al. 2013]
- pluripotent stem cells exposed for 3-days
- critical drop in ornithine : cystine ratio is the targeted readout
- **Key point:** 183 of 1065 (17%) ToxCast chemicals tested positive



SOURCE: Zurlinden et al. (manuscript)

ToxCast_STM assay

- *in vitro* accuracy anchored to *in vivo* DevTox studies (ToxRefDB)
- **Key point:** balanced accuracy improves with evidence for DevTox

		Stringency Filter Applied to DevTox Anchor					
		Condition ²	Base	Low	Medium	High	
<i>in vivo</i>	<i>in vitro</i>	TP	85	60	35	19	
		FP	14	37	23	9	
		FN	217	127	51	11	
		TN	116	208	176	88	
		<i>n</i>	432	432	285	127	
		<i>sensitivity</i>	0.281	0.321	0.407	0.633	
		<i>specificity</i>	0.892	0.849	0.884	0.907	
		<i>PPV</i>	0.859	0.619	0.603	0.679	
		<i>NPV</i>	0.348	0.621	0.775	0.889	
		ACC	46.5%	62.0%	74.0%	84.3%	
		MCC	0.190	0.202	0.332	0.554	
							
			any dLEL rat OR rabbit	SOME evidence rat OR rabbit	CLEAR evidence rat OR rabbit	CLEAR evidence rat AND rabbit	

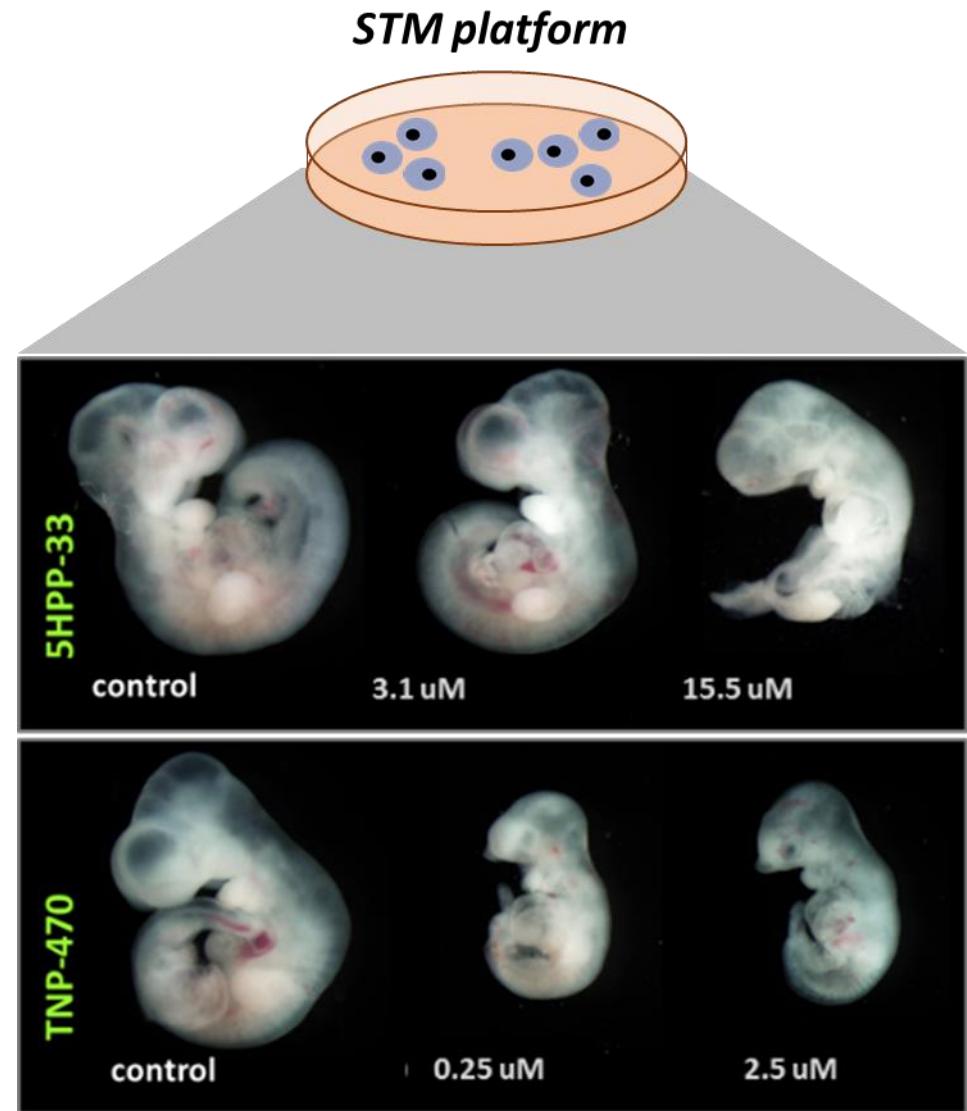
STM versus rat WEC

5HPP-33: synthetic thalidomide analog

- T.I. predicted 9.5 μM
- AC50 observed 21.2 μM (embryo viability)

TNP-470: synthetic fumagillin analog

- T.I. predicted 0.01 μM
- AC50 observed 0.04 μM (dysmorphogenesis)



Key point: exposure-based potential for DevTox predicted by STM assay (quantitative prediction).

SOURCE: Ellis-Hutchings et al. (2017) Reprod Toxicol

DATA SETS

ToxCast_NVS biochemical AC50

337 inhibited features
83 activated features

X

ToxCast_STM binary hit calls

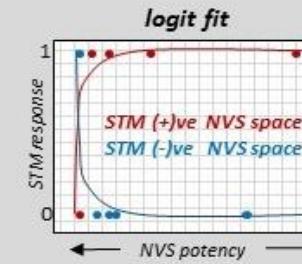
183 STM-positive
882 STM-negative

ASSAY SELECTION

Logistic Regression

Gene Potency Score (GPS)

- chemical-specific gene score
- consolidate AC50 homologs
- add up and down extensions



267

Phenotype Weighting

HMDC database

- 28 phenotype systems
- 233 GPS bins (0,1,2,3)
- log₂ normalization
- top 40 weighted correlations

STM(+)

5.675	PTPN11_up
4.003	NR3C1_down
2.134	RAF1_down
2.012	PDE5A_down
2.039	EGFR_down
1.921	IRAK4_up
1.821	TACR3_down
1.753	KDR_down
1.703	FLT4_down
1.662	FLT1_down
...	...

STM(-)

- **sensitive:** regulation of PI3K signaling, FoxO signaling pathway, and focal adhesion pathway.
- **insensitive domain:** GPCR signaling through G(q) and steroid hormone mediated signaling pathways.

Pathways and Processes

DAVID 6.8 bioinformatics resources

- GO Direct, KEGG, Reactome, INTERPRO
- Benjamini corrected $p \leq 0.05$
- redundancies resolved manually by FDR
- 144 category Pearson correlation matrix
- discuss keystone pathways/processes



FUNCTIONAL ANNOTATION

Keystone Pathways

2. Translating biomolecular lesion(s) into quantitative phenotypes for predictive developmental toxicology



Objective: build and test computer models of complex tissues that advance critical phenomena (specificity, canalization, plasticity) for quantitative prediction for virtual screening and *in silico* testing.

SOURCE: Andersen, Newman and Otter (2006) Am. Assoc. Artif. Intel.

Cell agent-based models (ABMs)

- **Approach:** build and test self-organizing morphogenetic systems *in silico* using CC3D modeling environment (www.compuccell3d.org).
- **Input:** A.I. cast into mathematically-defined cells (agents), synthetic gene circuits, and viscoelastic properties to emulate developmental progression.
- **Emergence:** simulation resolves into normal or perturbed phenotypes reading *in vitro* data input from specific ToxCast assays ([cybermorphs](#)).
- **Output:** probabilistic rendering of where, when and how a developmental defect might occur ([critical phenomena](#)).

AOP framework: *cleft palate* as an example



ToxCast Chemicals

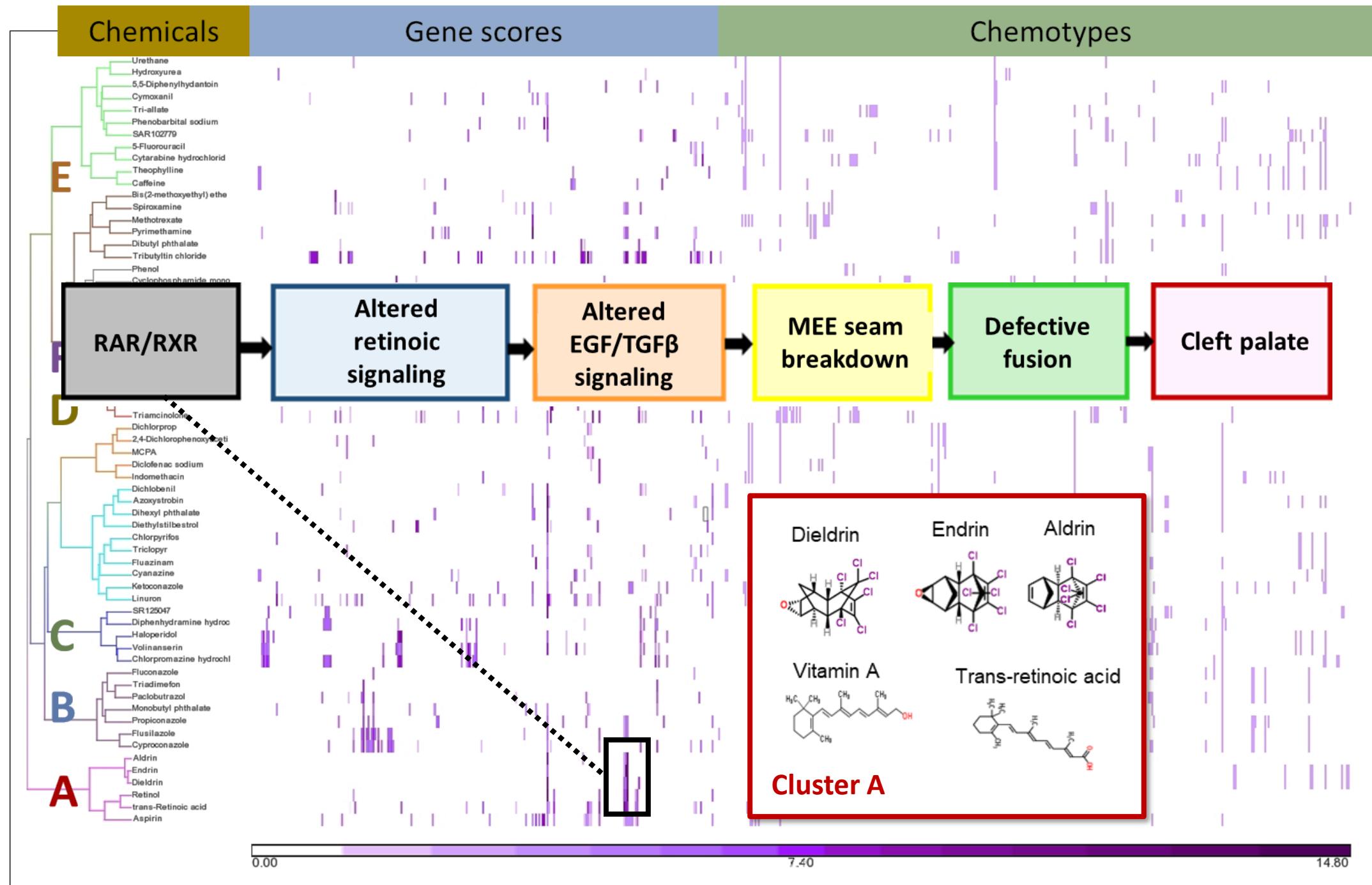
500 chemicals summarized by ToxCast gene score and chemotype for machine-learning

Animal studies

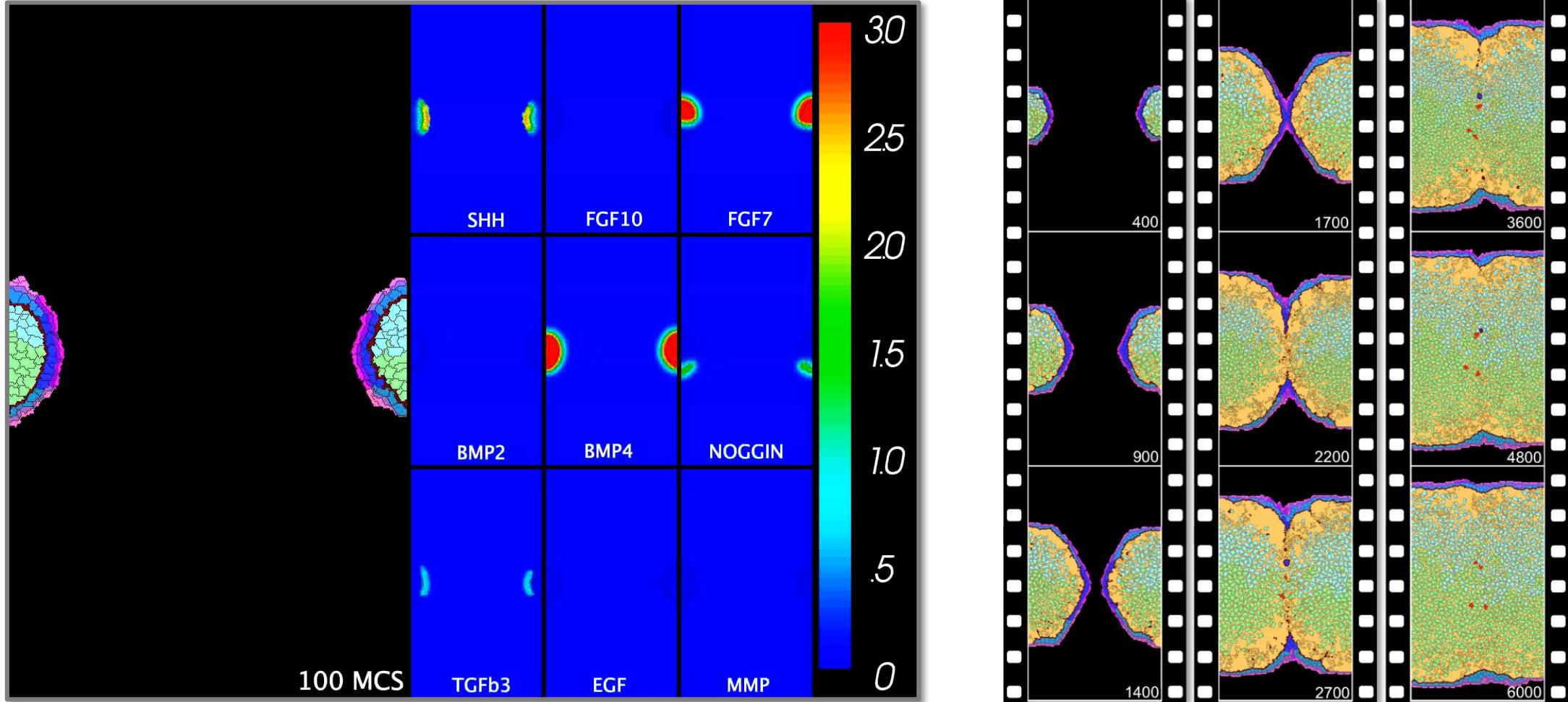
63 chemicals associated with cleft palate in ToxRefDB or open literature

AOP clusters

6 mechanistic pathways inferred from integration of HTS data with chemical structure.

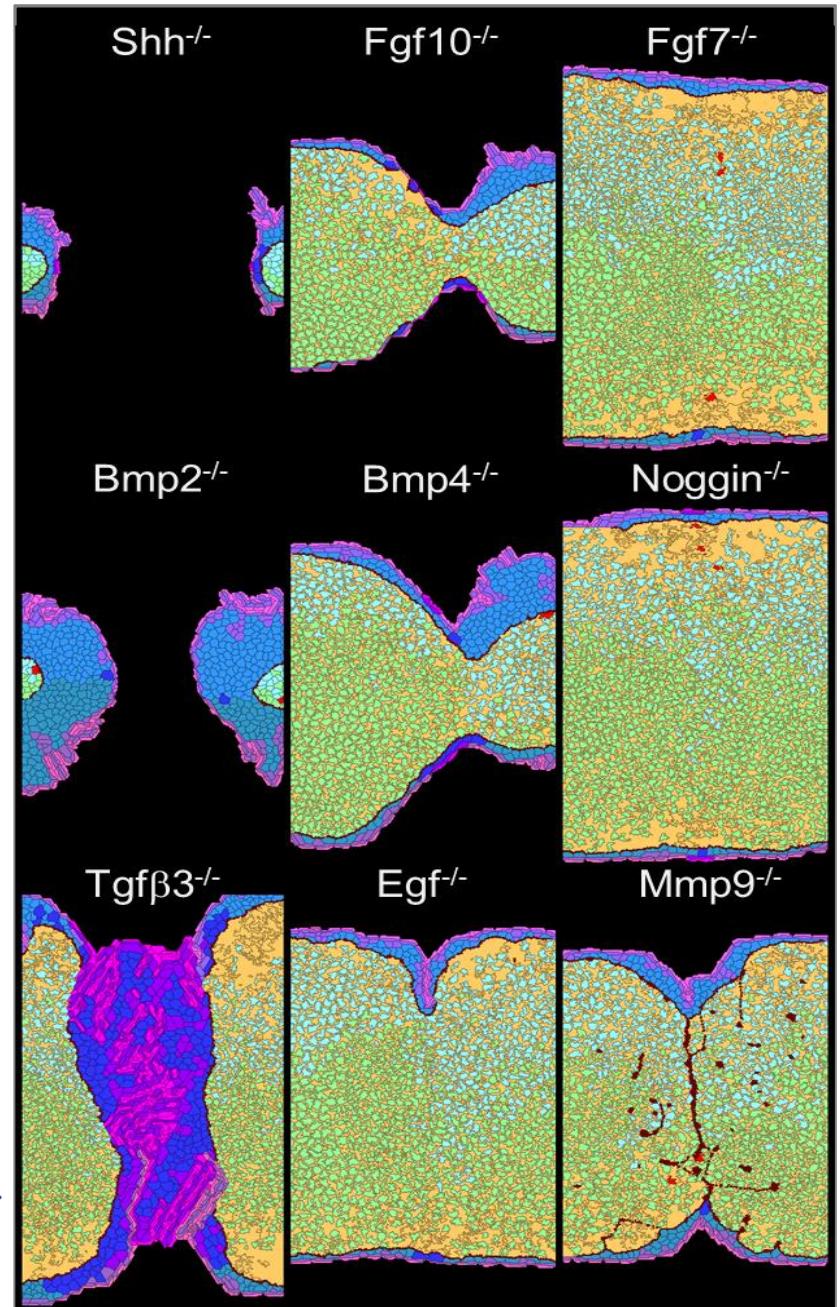
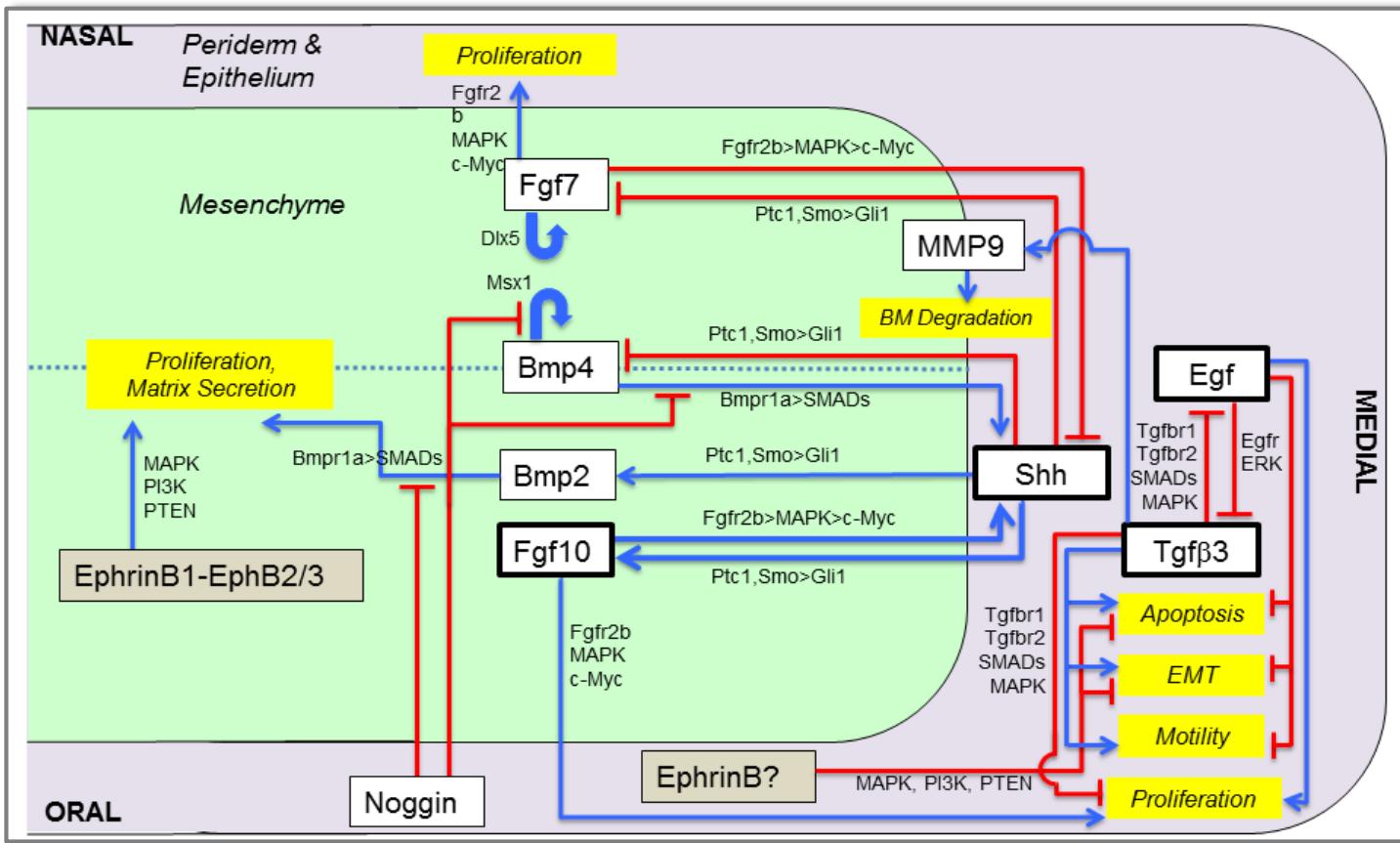


Epithelial seam breakdown and mesenchymal confluence during palatal fusion



SOURCE: Hutson et al. (2017) Chem Res Toxicol

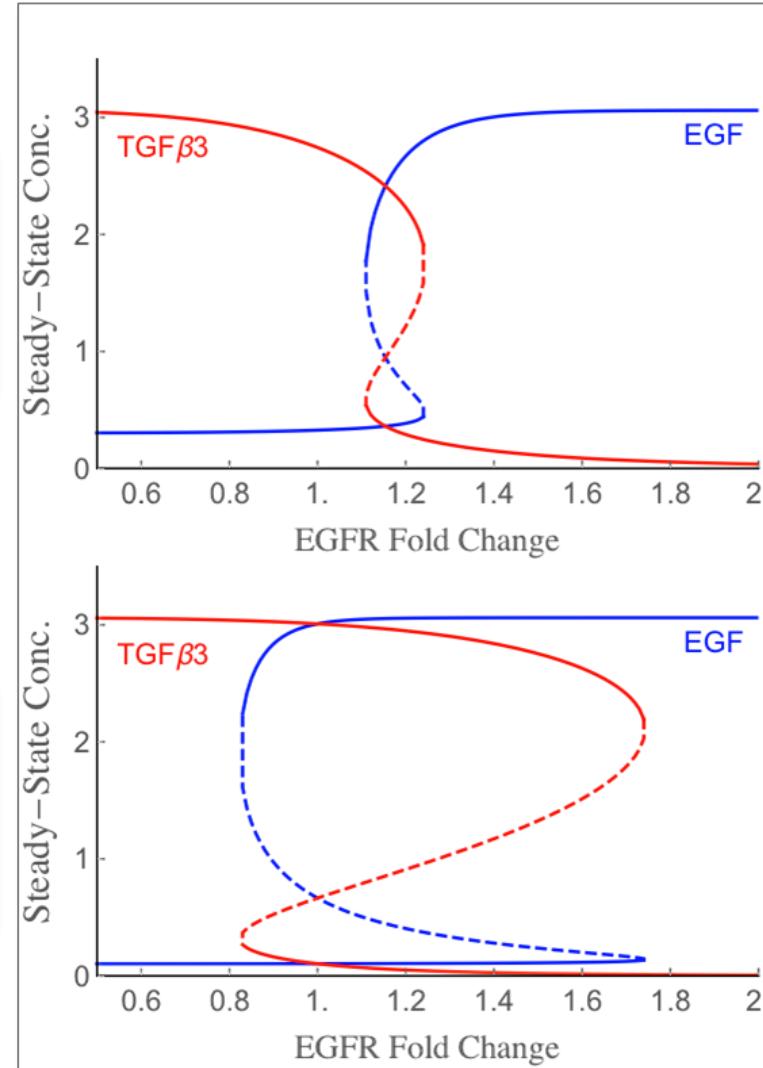
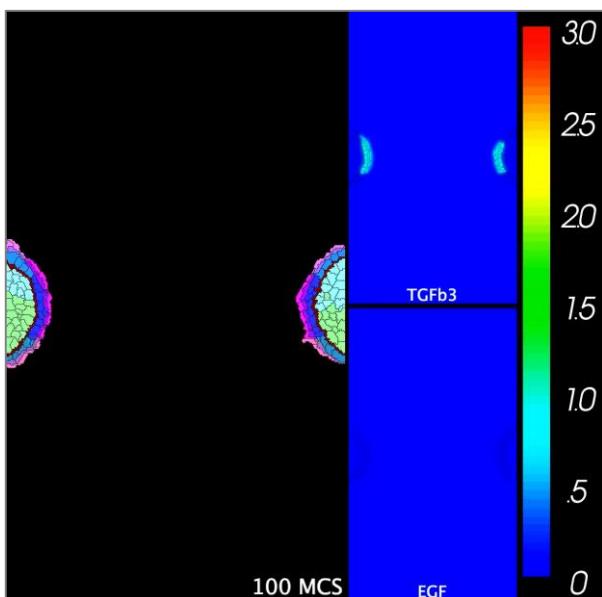
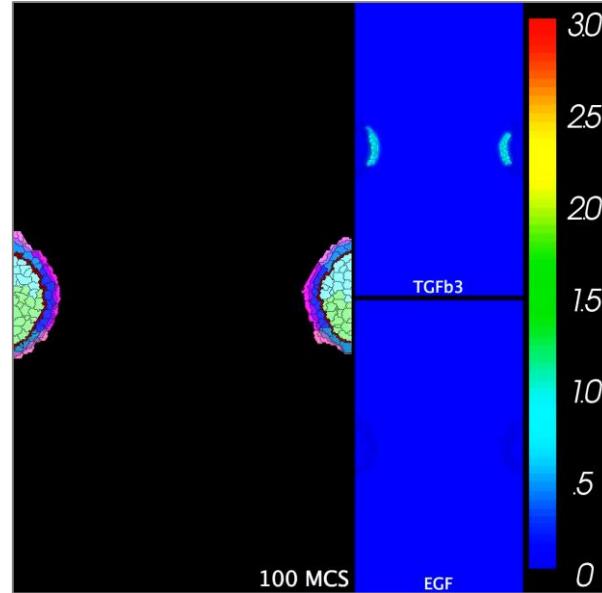
Hacking the control network



A.I. = synthetic cell signaling networks

Cybermorphs = simulated loss of function

Messin' with the switch: two scenarios for bistable dynamics



**Narrow
hysteresis:**
*less resilient
but reversible*

**Broad
hysteresis:**
*more resilient
but irreversible*

ToxCast dataset

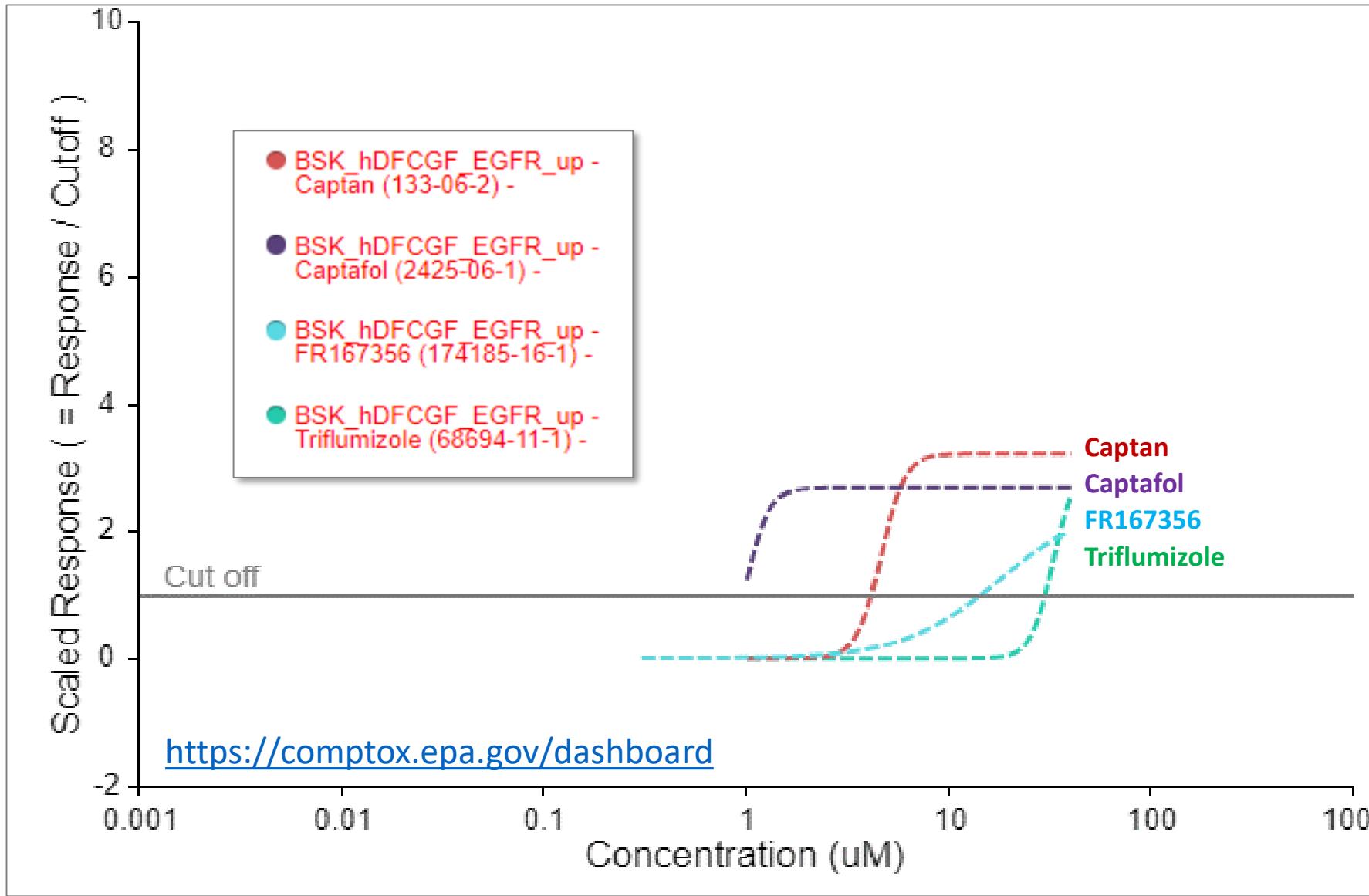
ChemicalName	FR_up	iFR_up	1_down	b1_down	ToxRefDB
Methylene bis(thiocyanate)	1.14	2.13	5.93	4.26	NEG
Zoxamide	14.22	1.85	17.37	9.69	NEG
2-(Thiocyanomethylthio)benzothiazole	2.28	1.54	6.48	7.21	NEG
Diphenylamine	32.71	1.49	5.95	1.63	NEG
Azamethiphos	0.89	1.81	1000.00	1000.00	NEG
Bromacil	20.50	1.57	1000.00	1000.00	NEG
Forchlorfenuron	0.02	1.53	1000.00	1000.00	NEG
Methyl isothiocyanate	4.60	1.44	1000.00	1000.00	NEG
Diuron	16.51	1.44	1000.00	1000.00	NEG
Rotenone	0.82	1.42	1000.00	1000.00	NEG
Captan	4.59	2.57	7.15	7.25	POS
Triflumizole	32.71	2.48	19.88	19.88	POS
Butachlor	32.71	2.47	17.85	17.85	POS
Captafol	1.02	2.20	3.76	3.25	POS
Thiram	4.45	1.96	6.95	5.38	POS
Raloxifene hydrochloride	12.40	1.91	15.94	10.94	POS
Fluazinam	2.39	1.61	2.48	4.84	POS
Carbaryl	0.07	1.55	1000.00	1000.00	POS
Linuron	10.91	1.46	1000.00	1000.00	POS
Maneb	0.01	1.46	1000.00	1000.00	POS
Bendiocarb	8.75	1.43	1000.00	1000.00	POS
Fipronil	1.18	1.43	1000.00	1000.00	POS
Propoxur	1.67	1.43	1000.00	1000.00	POS
TNP-470	7.78	1.57	3.97	3.61	x
1-(2,3,8,8-Tetramethyl-1,2,3,4,5,6,7,8-octal	8.33	2.10	9.74	1.88	x
Trimethylolpropane triacrylate	2.02	1.80	5.17	1.41	x
Diiodomethyl 4-methylphenyl sulfone	3.15	1.77	3.74	17.68	x
1,2-Benzisothiazolin-3-one	8.22	1.74	11.91	14.70	x
Tralopyril	18.30	1.68	0.87	1.08	x
Bis(trichloromethyl)sulfone	1.95	1.61	4.49	5.74	x
N,N,N-Trimethyloctadecan-1-aminium chl	2.22	1.56	1.77	1.45	x
beta-Nitrostyrene	7.12	1.52	2.01	2.34	x
4,5-Dichloro-3H-1,2-dithiol-3-one	2.71	1.47	6.42	6.56	x
Tri-o-cresyl phosphate	8.95	1.45	9.54	1.56	x
Isobornyl methacrylate	13.66	1.44	21.86	1.97	x
SAR102779	0.05	1.43	12.95	14.97	x
PharmaGSID_48511	12.19	1.37	11.22	17.33	x
Perfluoroundecanoic acid	6.81	1.35	4.76	5.04	x
FR167356	17.65	2.06	1000.00	1000.00	x
Monobutyl phthalate	0.01	1.35	1000.00	1000.00	x
Niclosamide	0.58	2.14	1000.00	1000.00	x
Tripropylene glycol diacrylate	26.52	2.09	1000.00	1000.00	x
CP-457920	3.50	1.92	1000.00	1000.00	x
Trimethylolpropane trimethacrylate	32.85	1.81	1000.00	1000.00	x
alpha-Terpinal acetate	39.18	1.64	1000.00	1000.00	x
3-(4-tert-Butylphenyl)-2-methylpropanal	35.26	1.62	1000.00	1000.00	x
1,4-Dinitrobenzene	2.95	1.54	1000.00	1000.00	x
SB281832	34.72	1.54	1000.00	1000.00	x
2-(Morpholin-4-yldithio)-1,3-benzothiazole	5.61	1.52	1000.00	1000.00	x
Tolclofos-methyl	7.71	1.49	1000.00	1000.00	x
1,1':3',1"-Terphenyl	11.98	1.38	1000.00	1000.00	x
Estrone	0.03	1.35	1000.00	1000.00	x

μM effect in vitro	↑EGFR		↓TGFβ1		ToxRefDB DevTox
	AC50	top	AC50	top	
Captan	4.59	2.57	7.15	7.25	POS
Triflumizole	32.71	2.48	19.88	19.88	POS
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- 54 chemicals ↑EGFR density
- some also ↓TGF-beta signaling

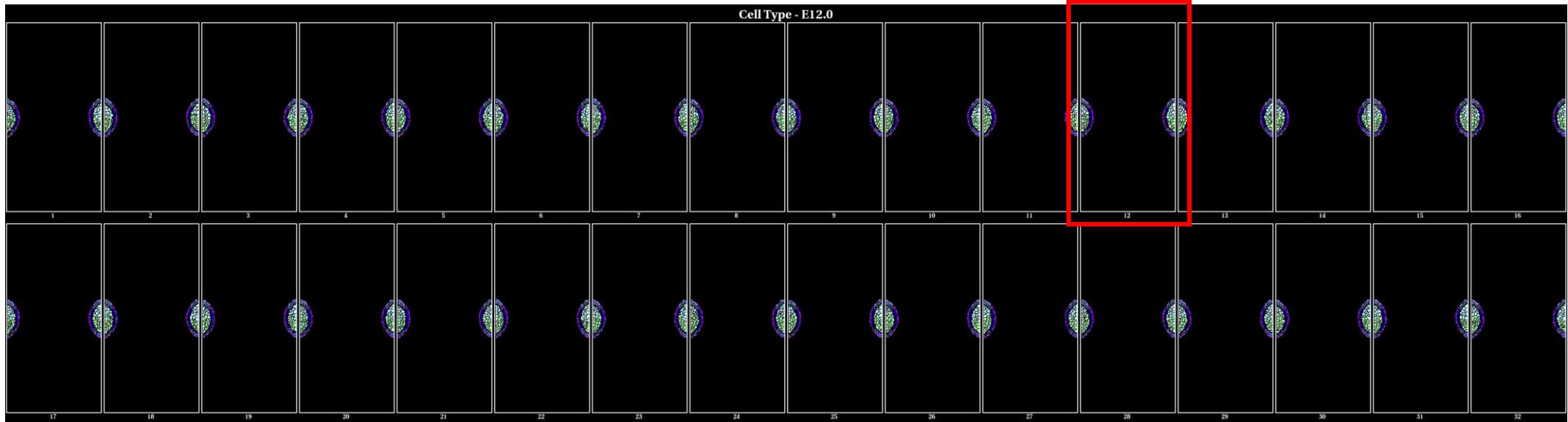
- negative for developmental toxicity in ToxRefDB
- positive for developmental toxicity in ToxRefDB
- positive for developmental toxicity in ToxRefDB

ToxCast dataset: EGFR: ↑ immunoreactivity relative to DMSO

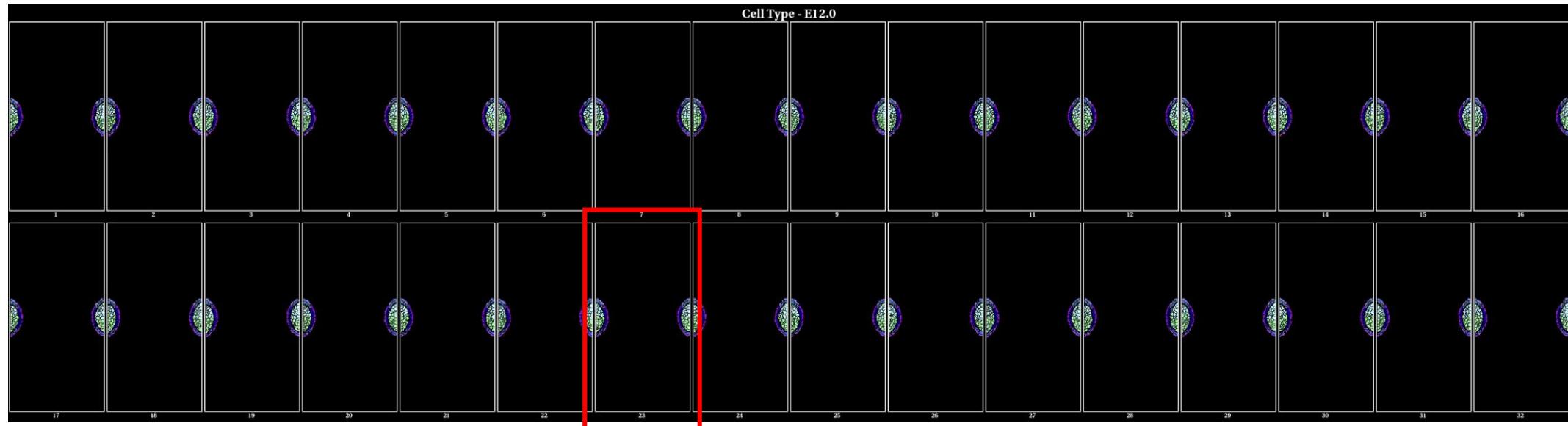


In silico dose-response: translating ↑EGFR conc. profile into a critical dose

Captan

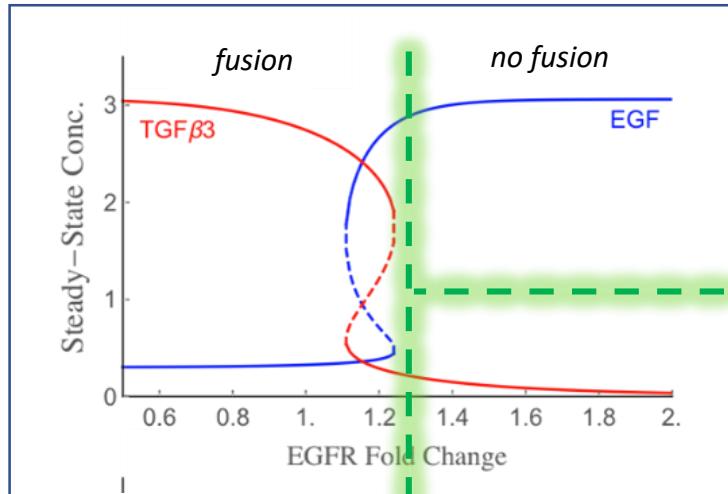


FR167356



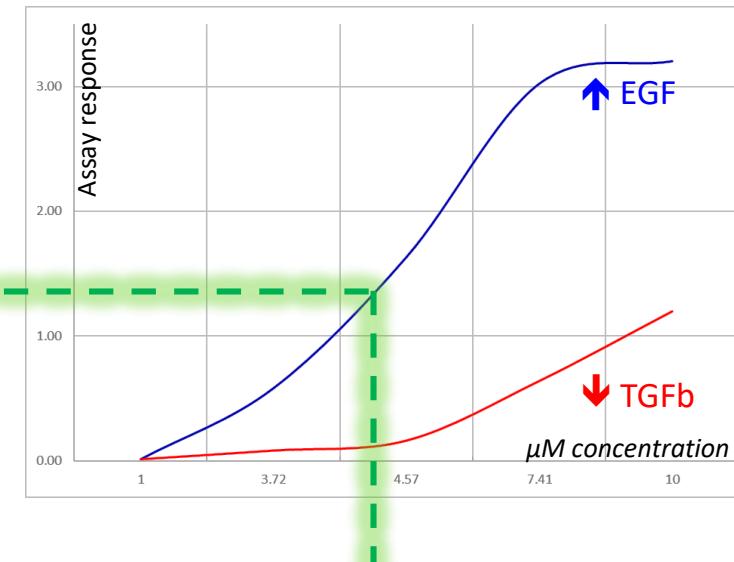
Predictive model: *modeling the critical phenomenon*

INPUT: switch dynamics



tipping point predicted by
computational dynamics
(hysteresis switch)

Captan in ToxCast



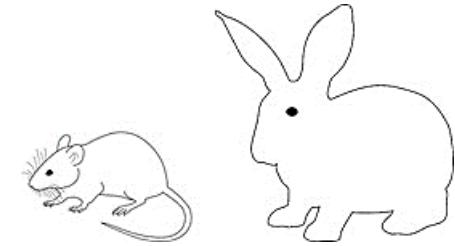
OUTPUT: tipping point
mapped to concentration
response (4 μM)

Captan in ToxRefDB
NEL = 10 mg/kg/day
LEL = 30 mg/kg/day

human HTTK model
2.39 mg/kg/day would
achieve a steady state of
4 μM in fetal plasma

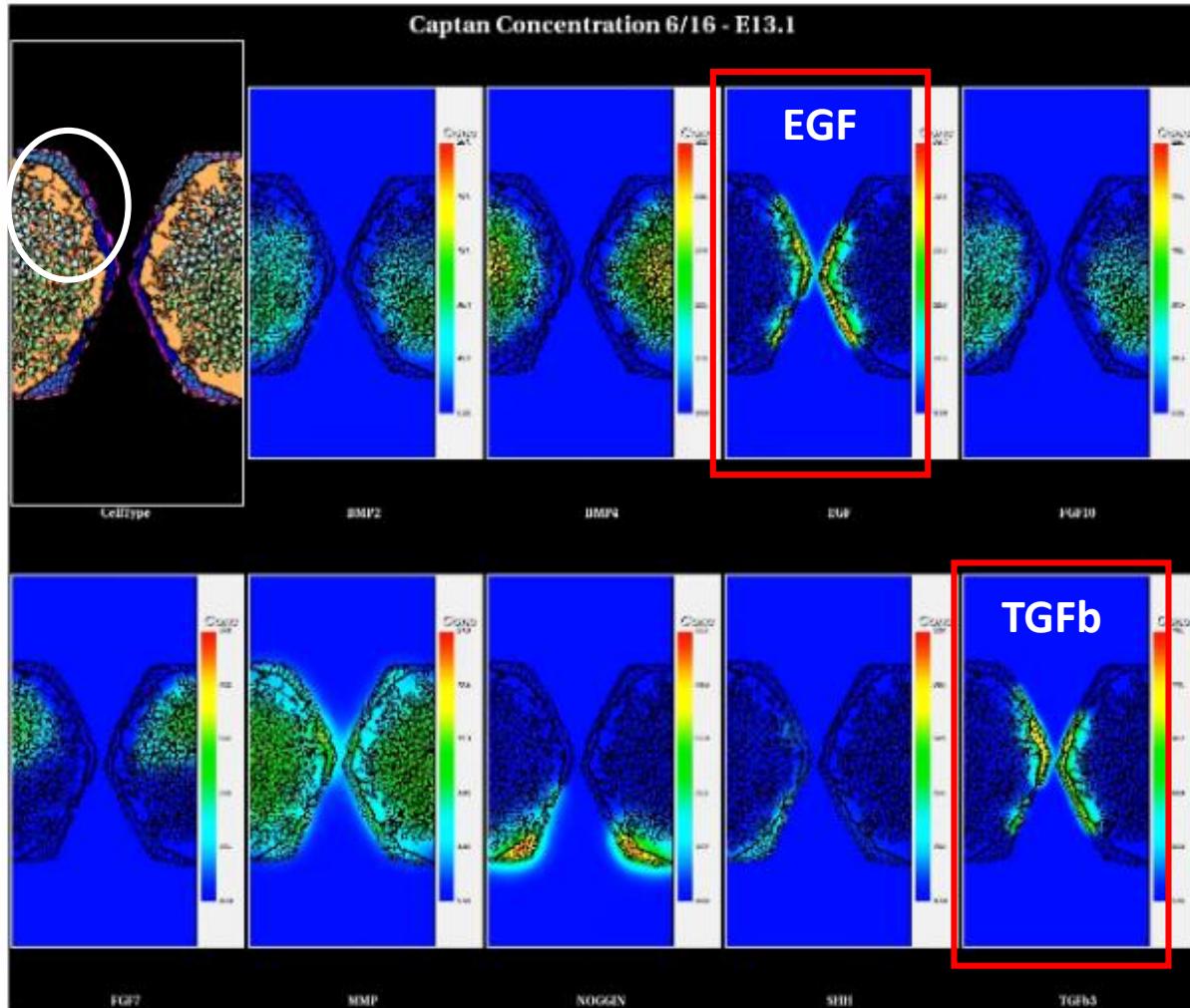


CompTox Chemicals Dashboard
exposure prediction
 $0.88 \times 10^{-7} \text{ mg/kg/day}$

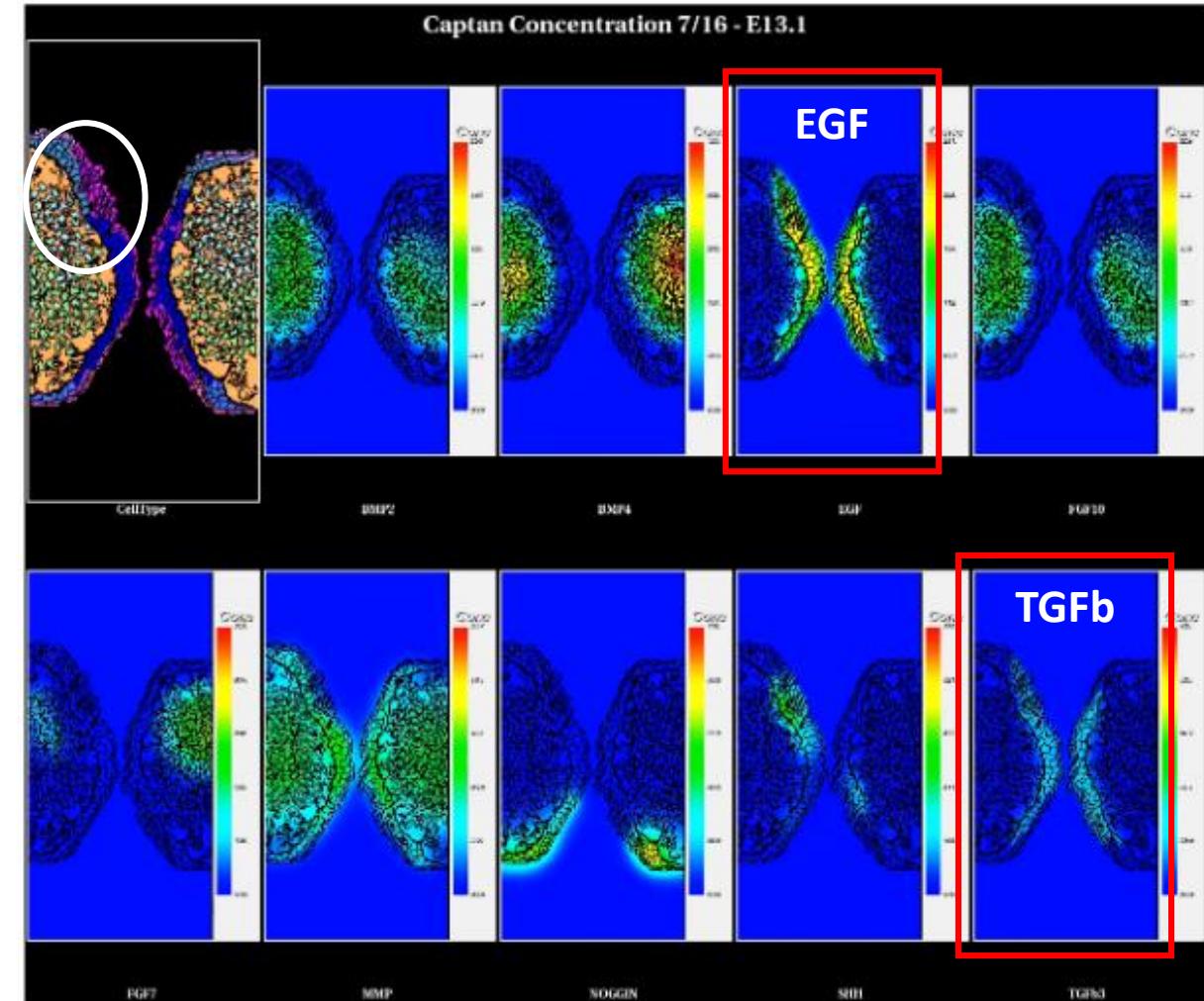


Pathogenesis: simulating the prefusion alterations

pre-critical dose



post-critical dose



Summary and Conclusions

*Computer modeling
is 3R's compliant!*

1. Several new approach methods (NAMs) are available for high-throughput screening chemical inventories for DevTox potential.

- STM assay in ToxCast gives an exposure-based readout of a chemical's DevTox hazard potential with 84% balanced accuracy.
- Assay sensitivity predicted high for kinase signaling converging on FoxO signaling but weak for estrogenic (ESR1) and G(q) signaling.

2. Cell ABMs recapitulate morphogenesis cell-by-cell and interaction-by-interaction as an embryonic system advances in time.

- Computer models simulate key events in AOPs to render mechanistic predictions and critical phenomena for DevTox.



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