



Computational Embryology:

Agent-Based Modeling and Simulation of External Genital Development

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Virtual Tissue Models (VTM) project

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NIH-NIDDK/AUA Workshop

"Congenital Anomalies of External Genitalia "

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DISCLAIMER: *The views expressed in this presentation are those of the presenter and do not necessarily reflect the views or policies of the US EPA*

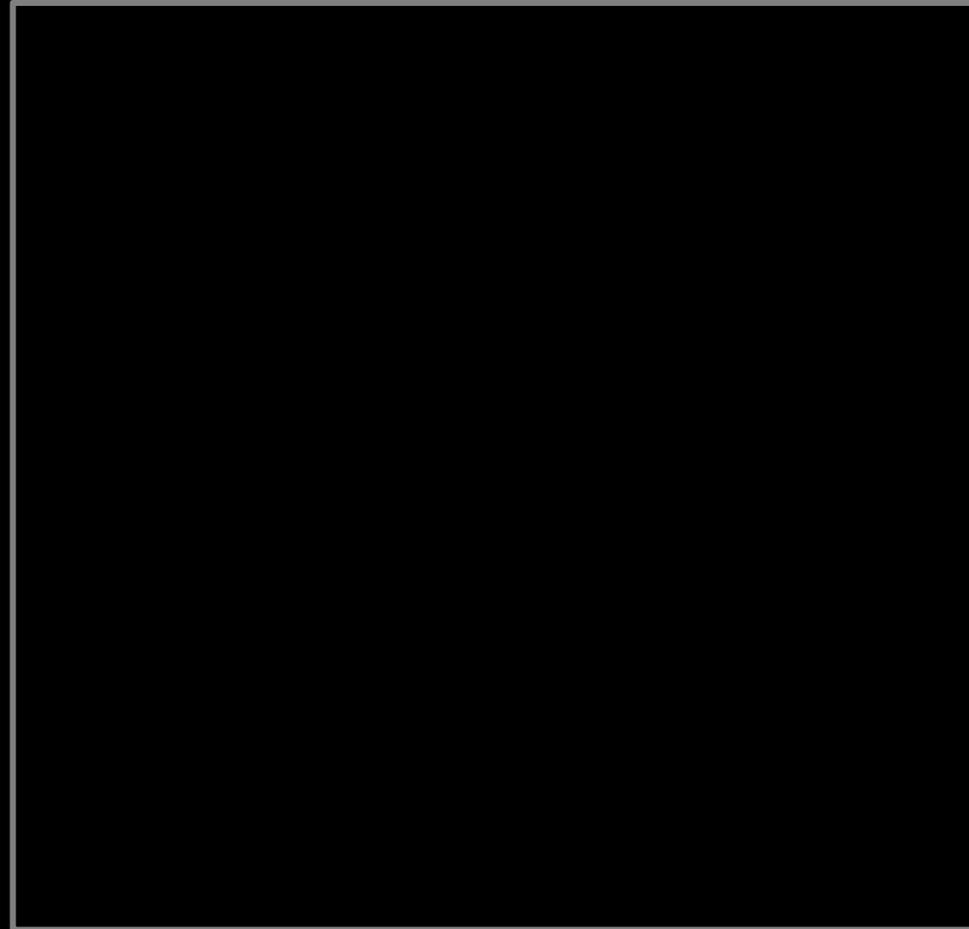
Scoping the Problem

Chemical regulation under the *Frank R. Lautenberg Chemical Safety for the 21st Century Act of 2016* ('amended TSCA') requires rapid affirmation of 'low' and 'high' priority substances based on unreasonable risk to vulnerable subpopulations and lifestages.

New approach methods (NAMs) based on HTS assays, complex *in vitro* systems, and *in silico* models can lead us in “*decoding the toxicological blueprint of active substances that interact with living systems*” [Sturla et al. 2014].



Anatomical homeostasis in a self-regulating Virtual Embryo



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

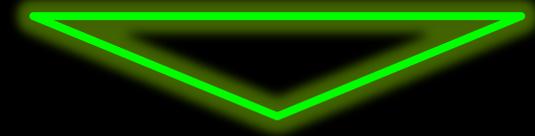
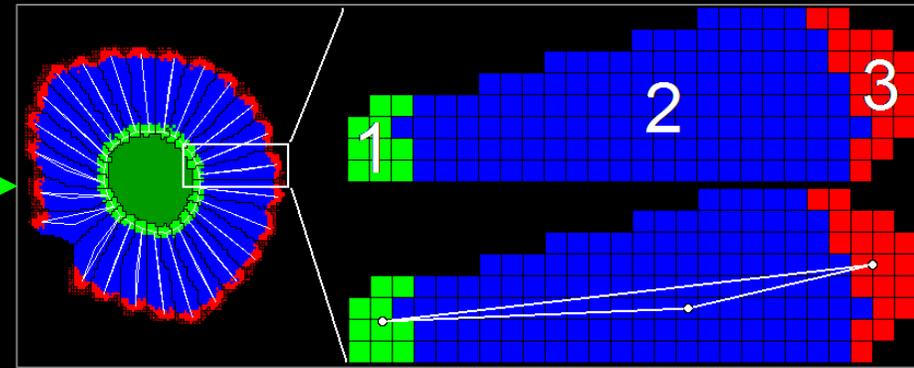
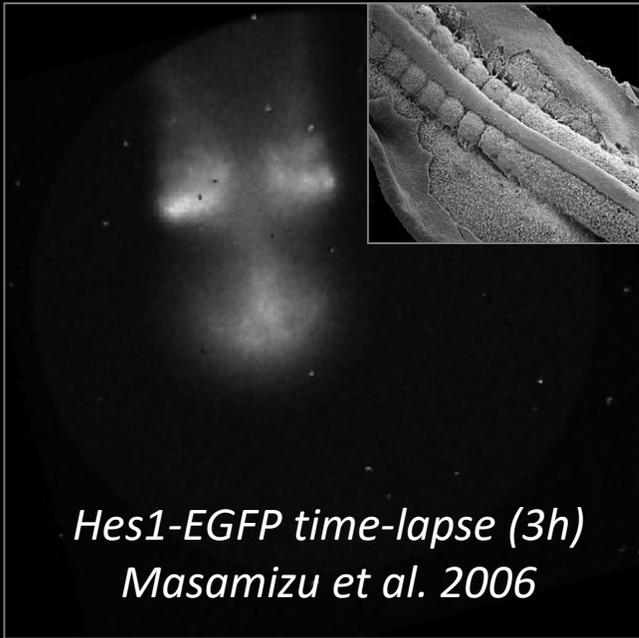
Agent-Based Models (ABMs):

in silico toxicology
is 3R's compliant!

- virtually reconstruct a morphogenetic field cell-by-cell and interaction-by-interaction (compucell3d.org)
- execute simulations that self-organize cells into higher-levels of biological structure and function (*emergent phenotypes*)
- simulate perturbations in the system - dose or stage response, critical pathways, non-chemical stressors, etc (*dynamics*)
- probabilistic rendering of where, when and how a developmental defect might occur (*mechanistic interpretation*)

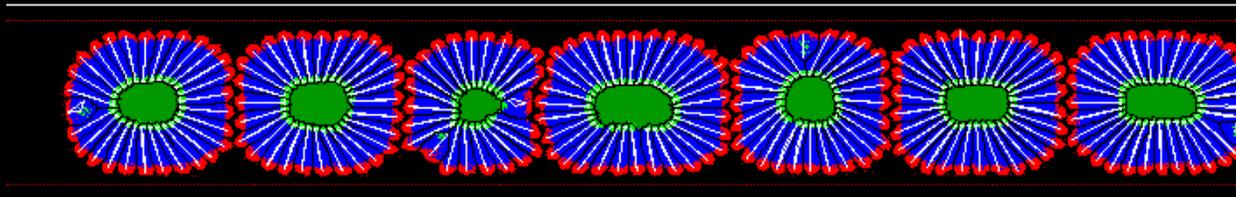


Modeling somite development

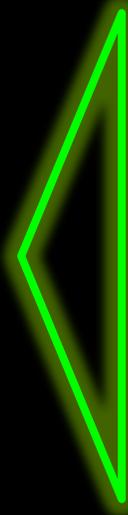


Differential cell adhesion

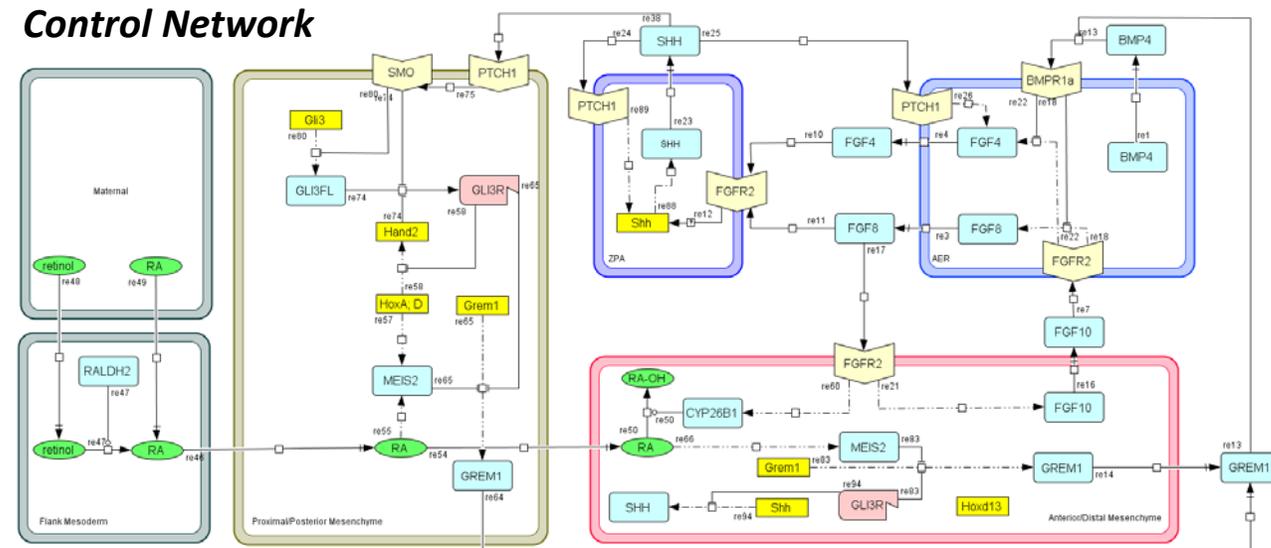
- clock genes do not oscillate
- somites form simultaneously



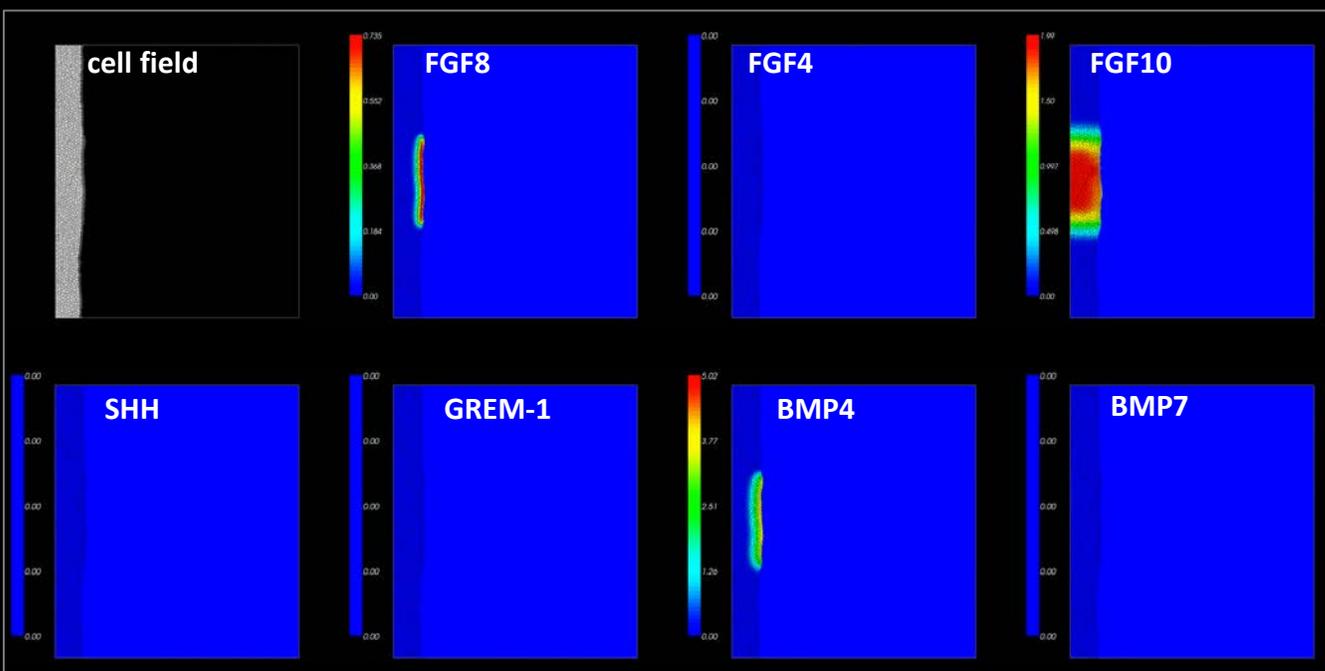
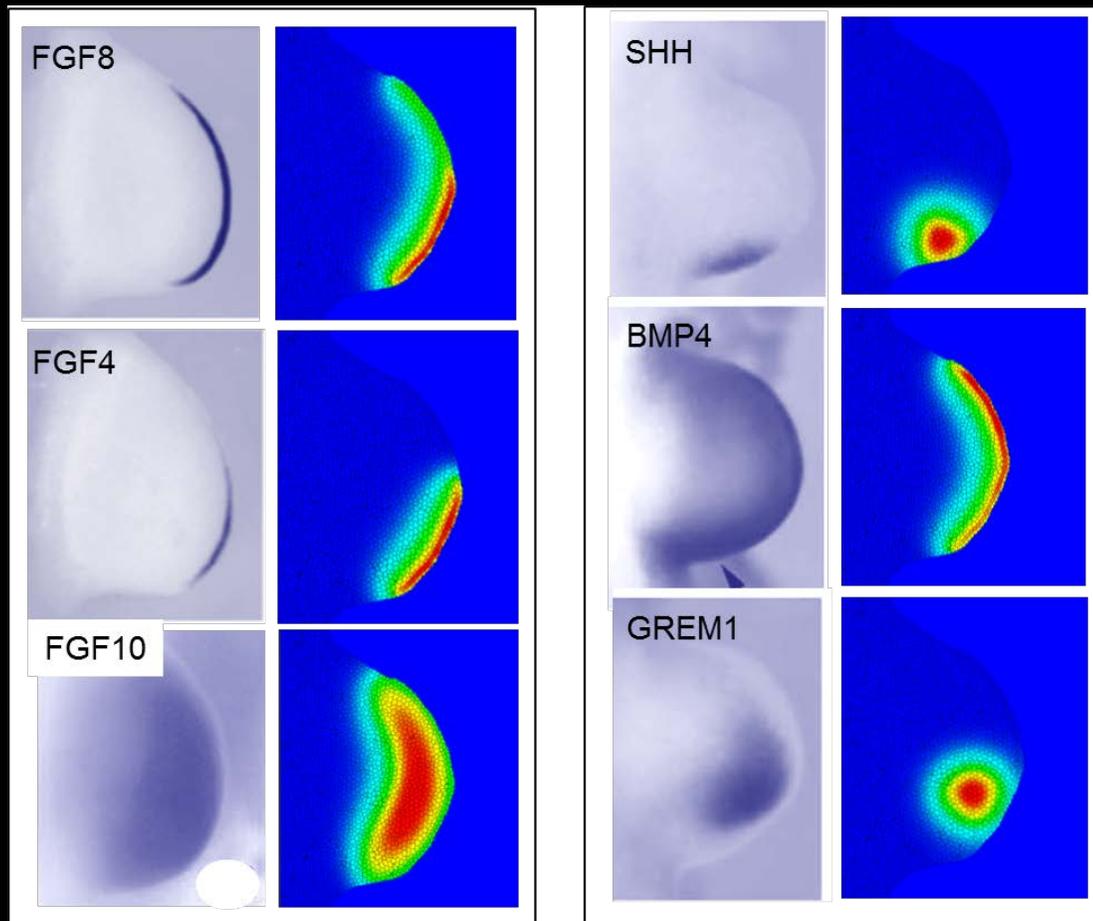
- *FGF8 wavefront restores sequentiality*
- *oscillatory clock improves regularity*



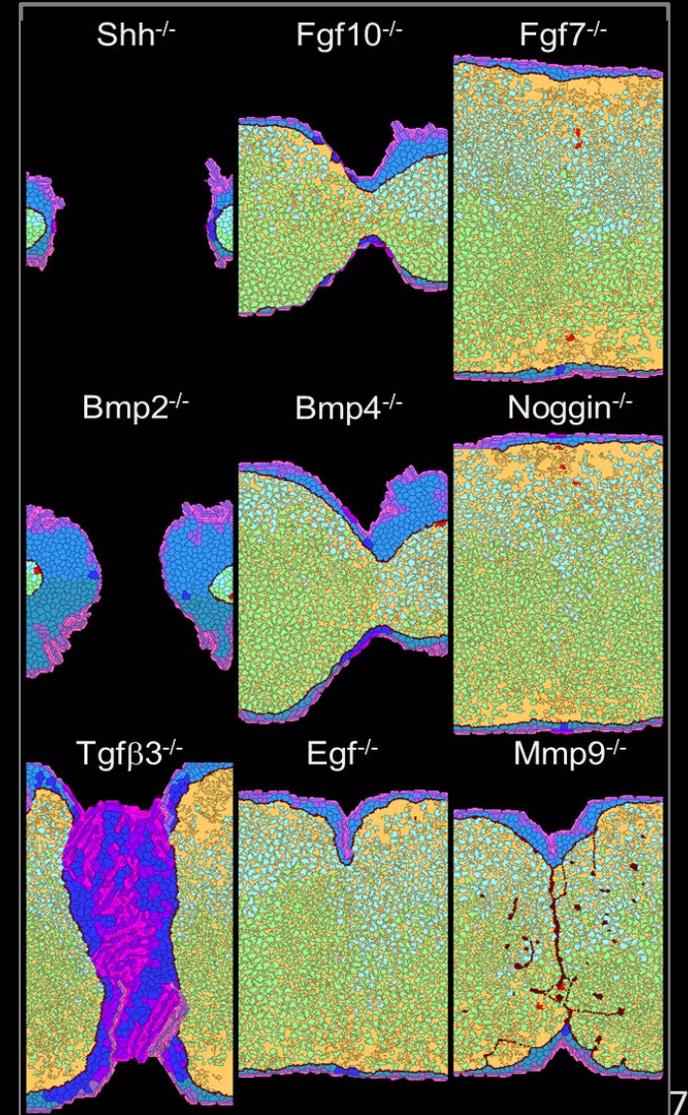
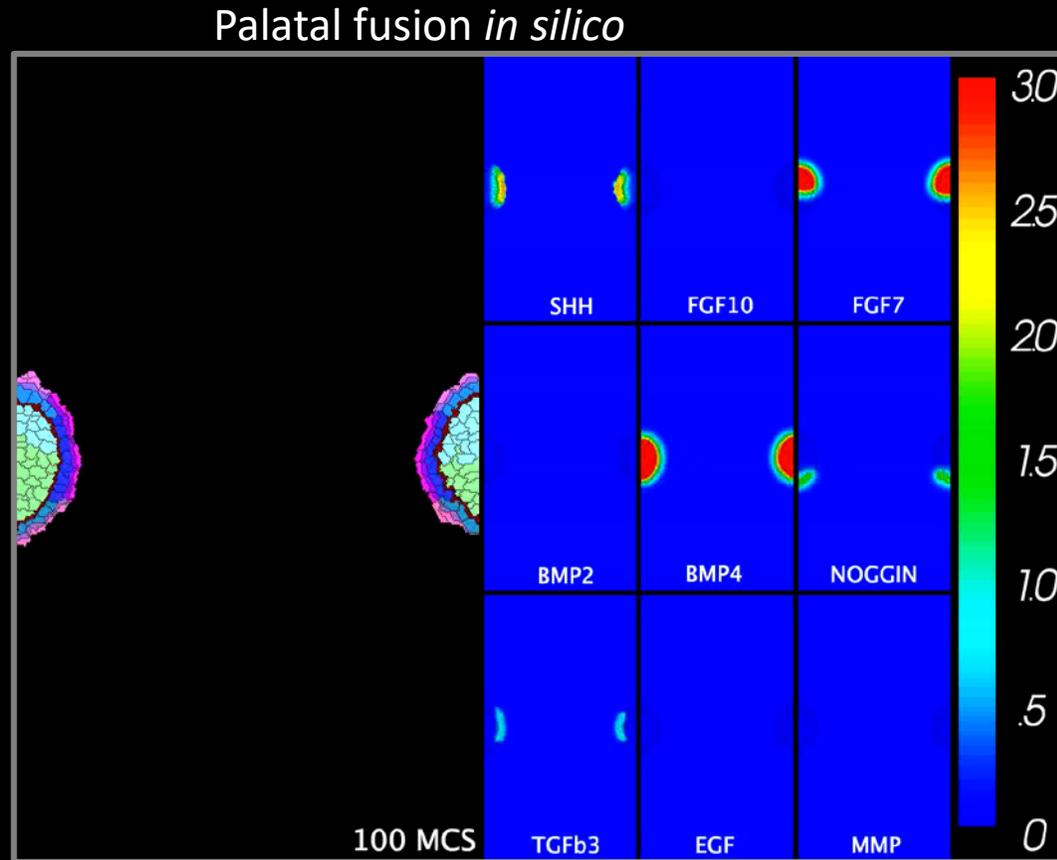
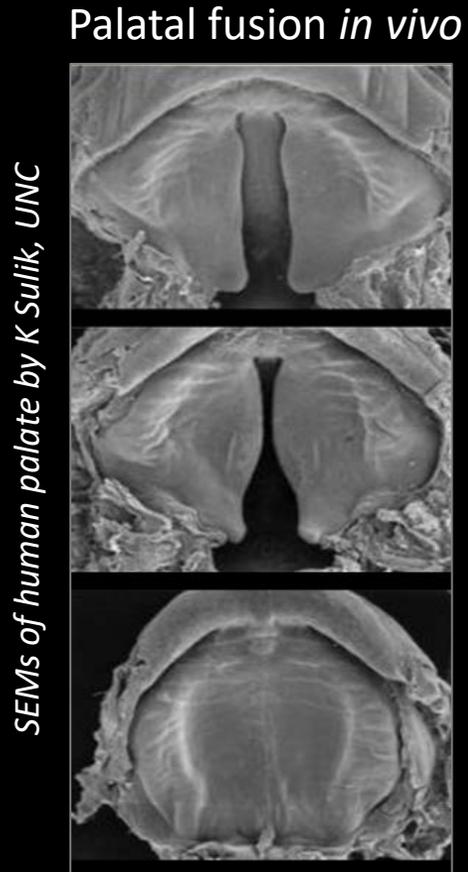
Control Network



Limb-bud outgrowth

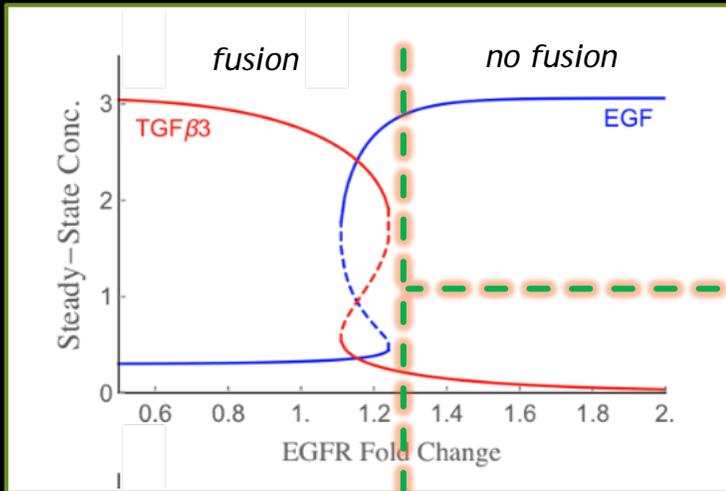


Palatal closure: driven by medial edge epithelium (MEE) seam breakdown



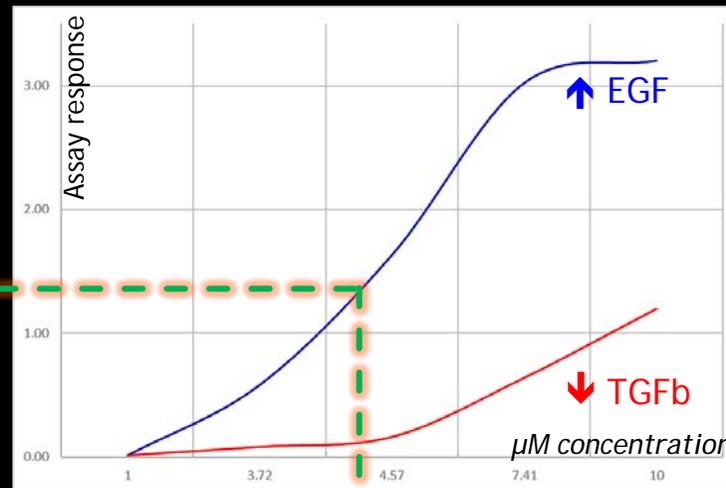
TGF-beta/EGF latch switch: *controls MEE breakdown*

INPUT: switch dynamics



tipping point predicted by
computational dynamics
(hysteresis switch)

Captan in ToxCast



OUTPUT: tipping point
mapped to concentration
response (4 μ M)

Captan in ToxRefDB

NOAEL = 10 mg/kg/day

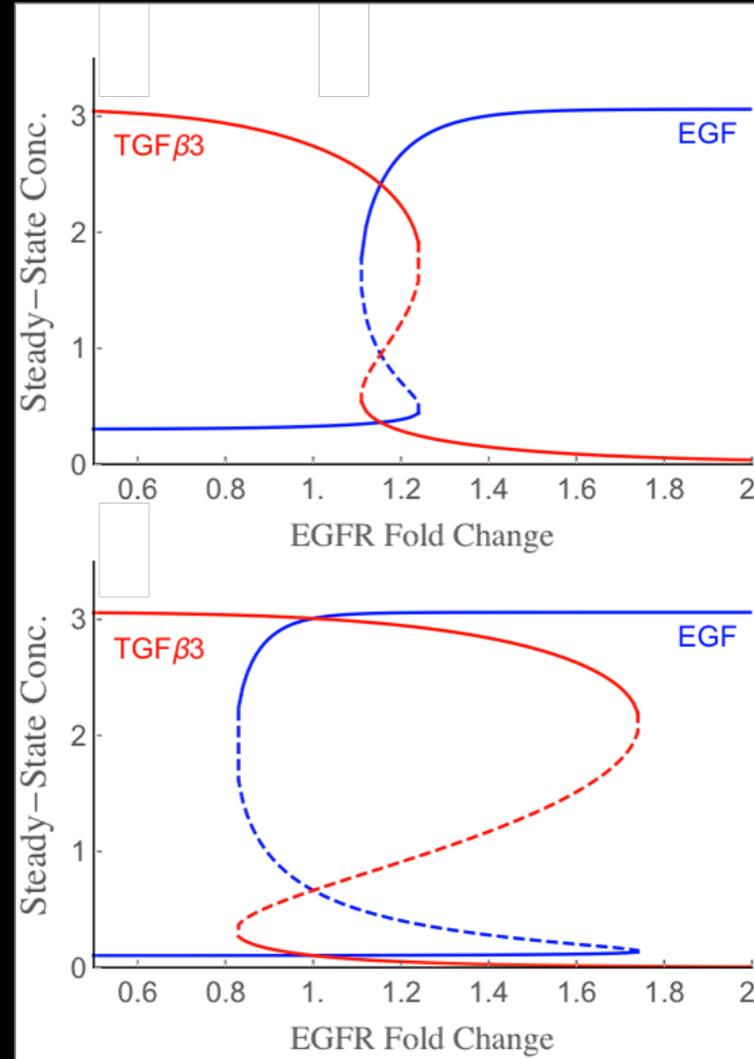
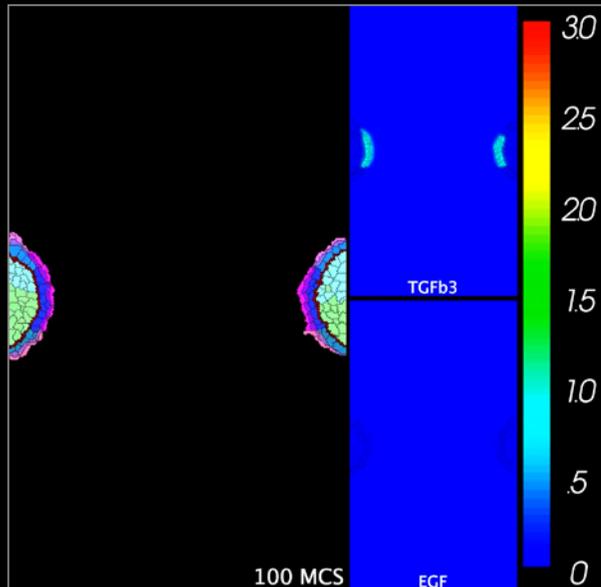
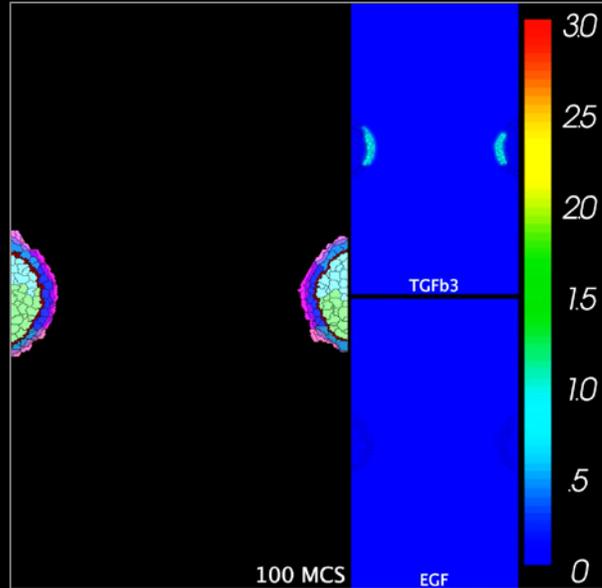
LOAEL = 30 mg/kg/day

human HTTK model

2.39 mg/kg/day would

achieve a steady state of 4
 μ M in fetal plasma

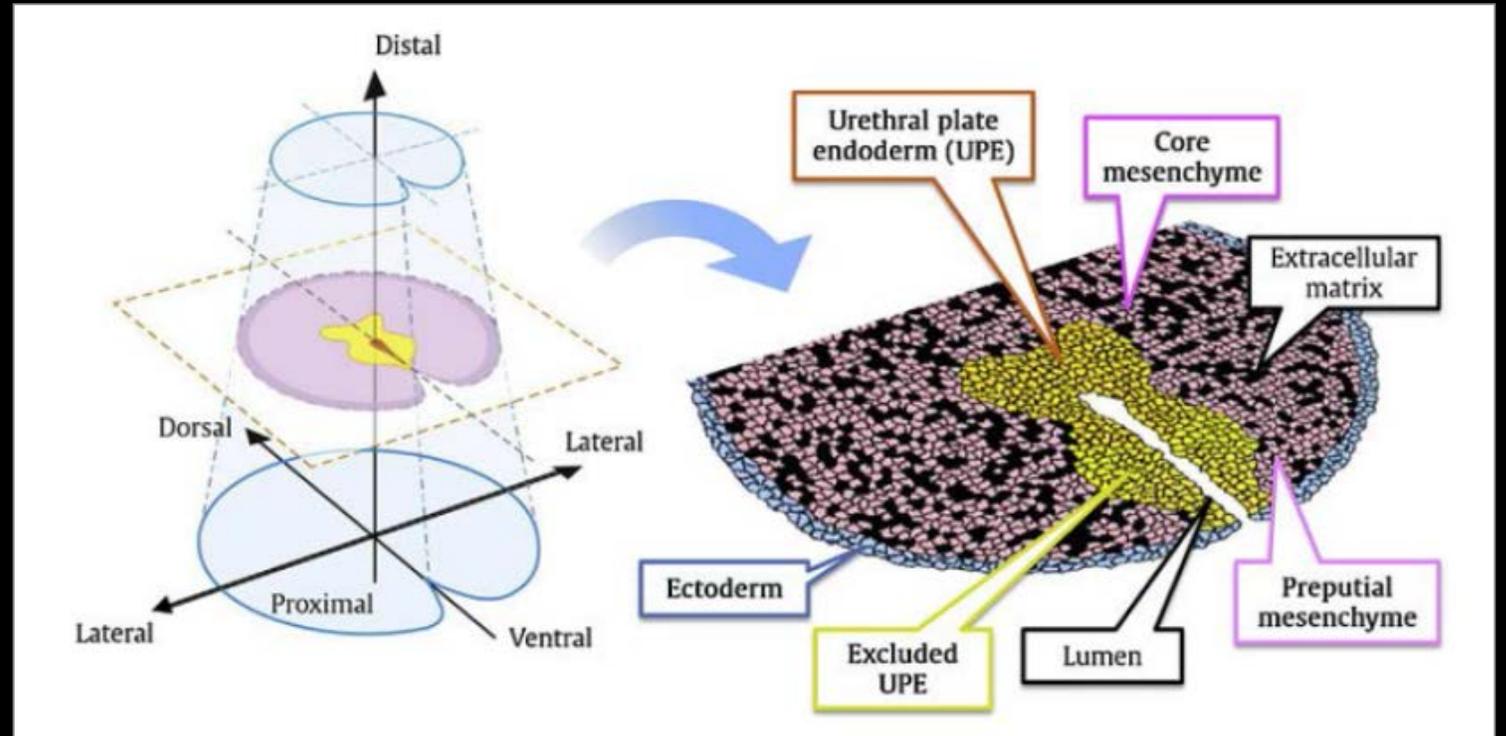
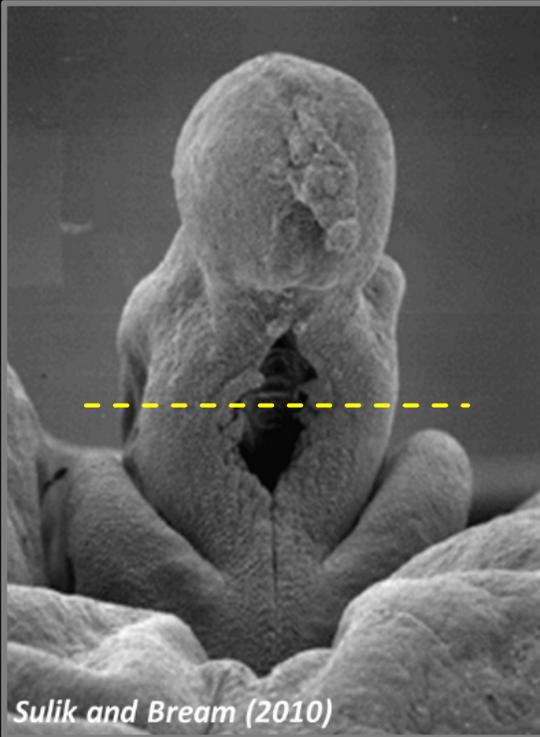
Messin' with the switch: *two scenarios for bistable dynamics*



Narrow hysteresis:
less resilient but reversible

Broad hysteresis:
more resilient but irreversible

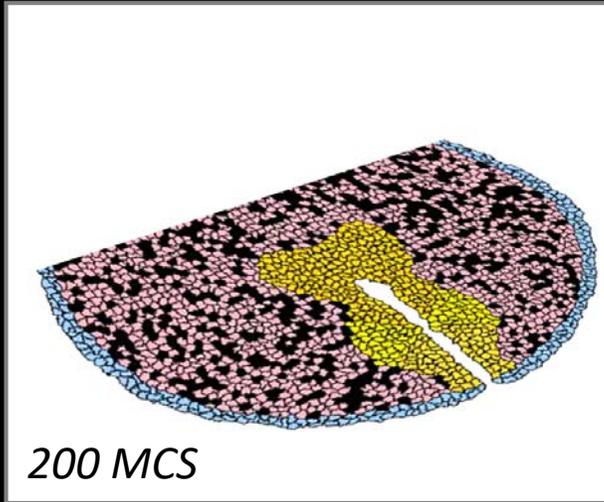
Genital tubercle: abstraction for ABM-simulated urethral closure



GT development modeled @ 1000 MCS/day of gestation (mouse):

- *E13.5 - urethral plate endoderm (UPE)*
- *E15.5 - ventral elevation of urethral folds*
- *E17.5 - urethral fusion and septation*

Control network: minimal requirements for dual-reciprocal signaling



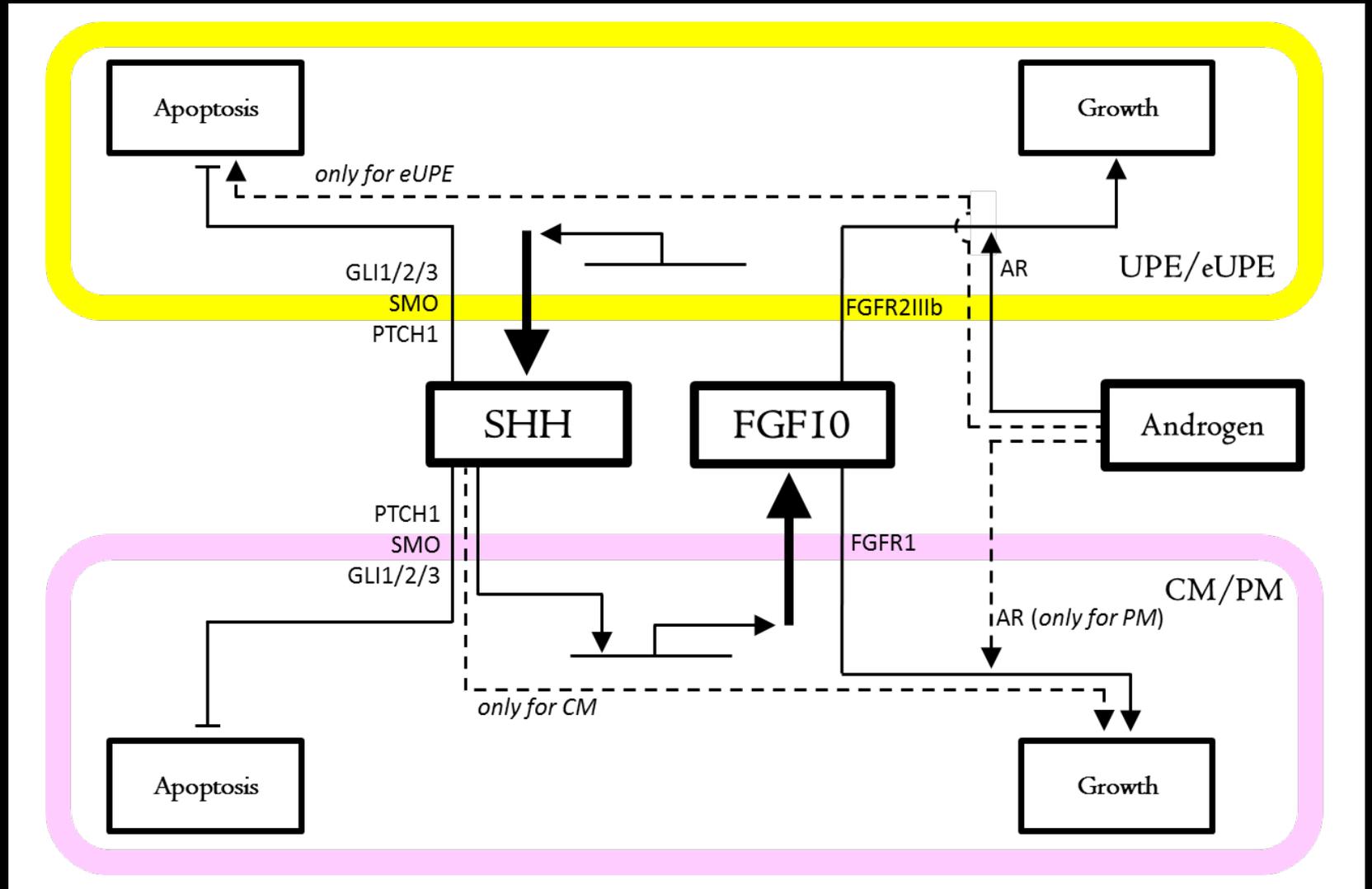
Biokinetic gradients:

FGF10 – simple diffusion

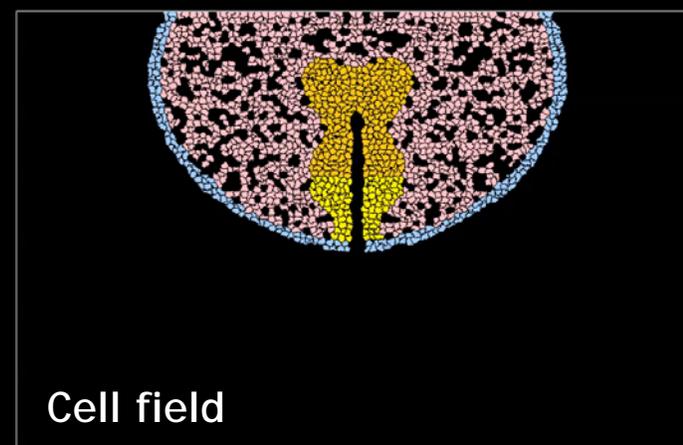
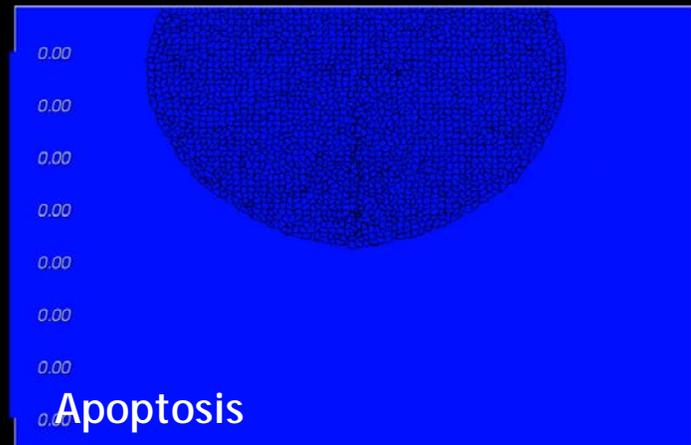
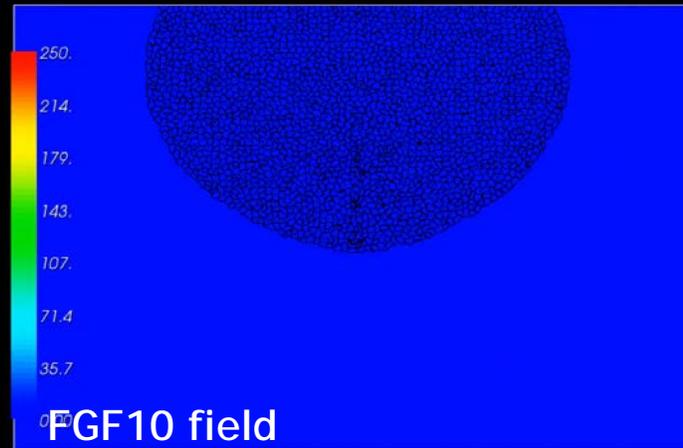
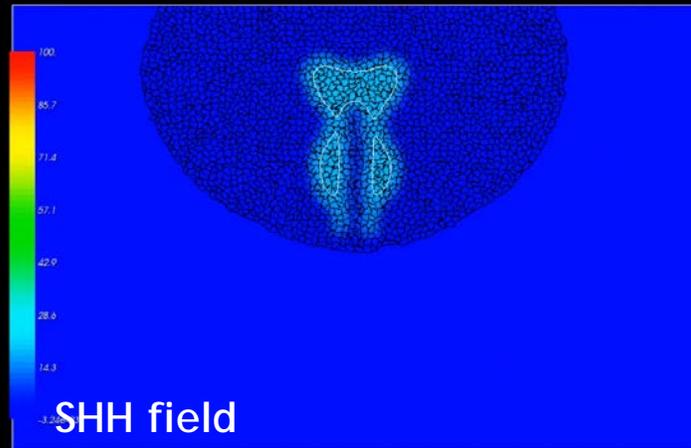
SHH – transcytosis (1/8 rate)

An – latched @2000 MCS (E15.5)

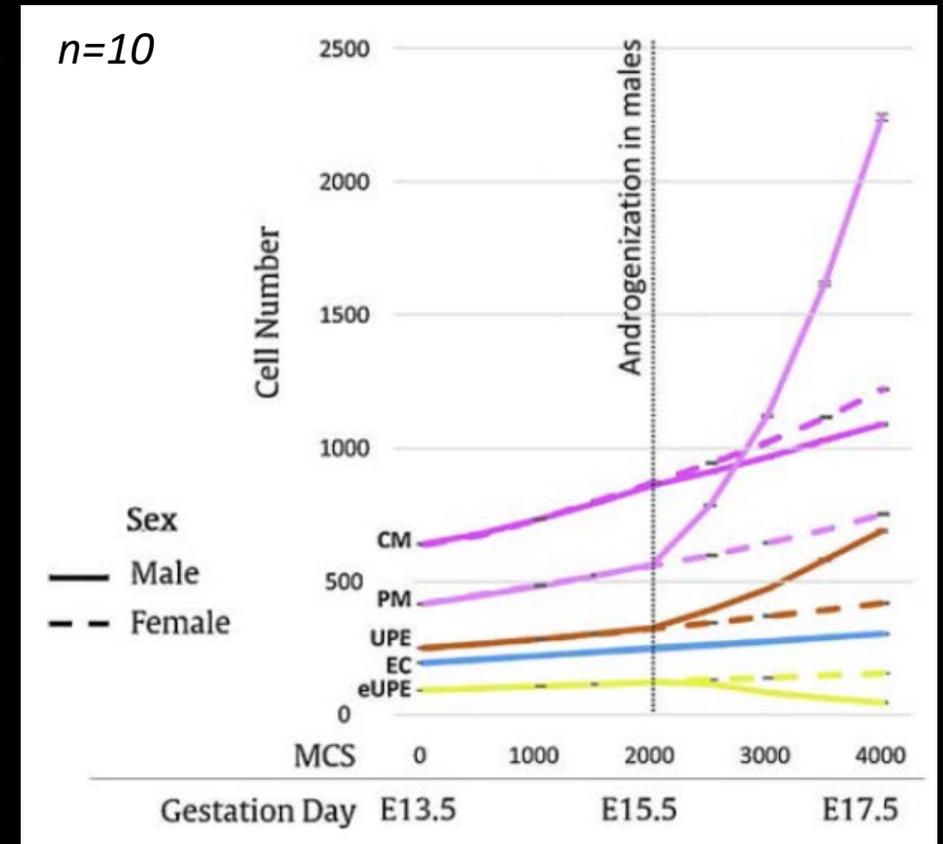
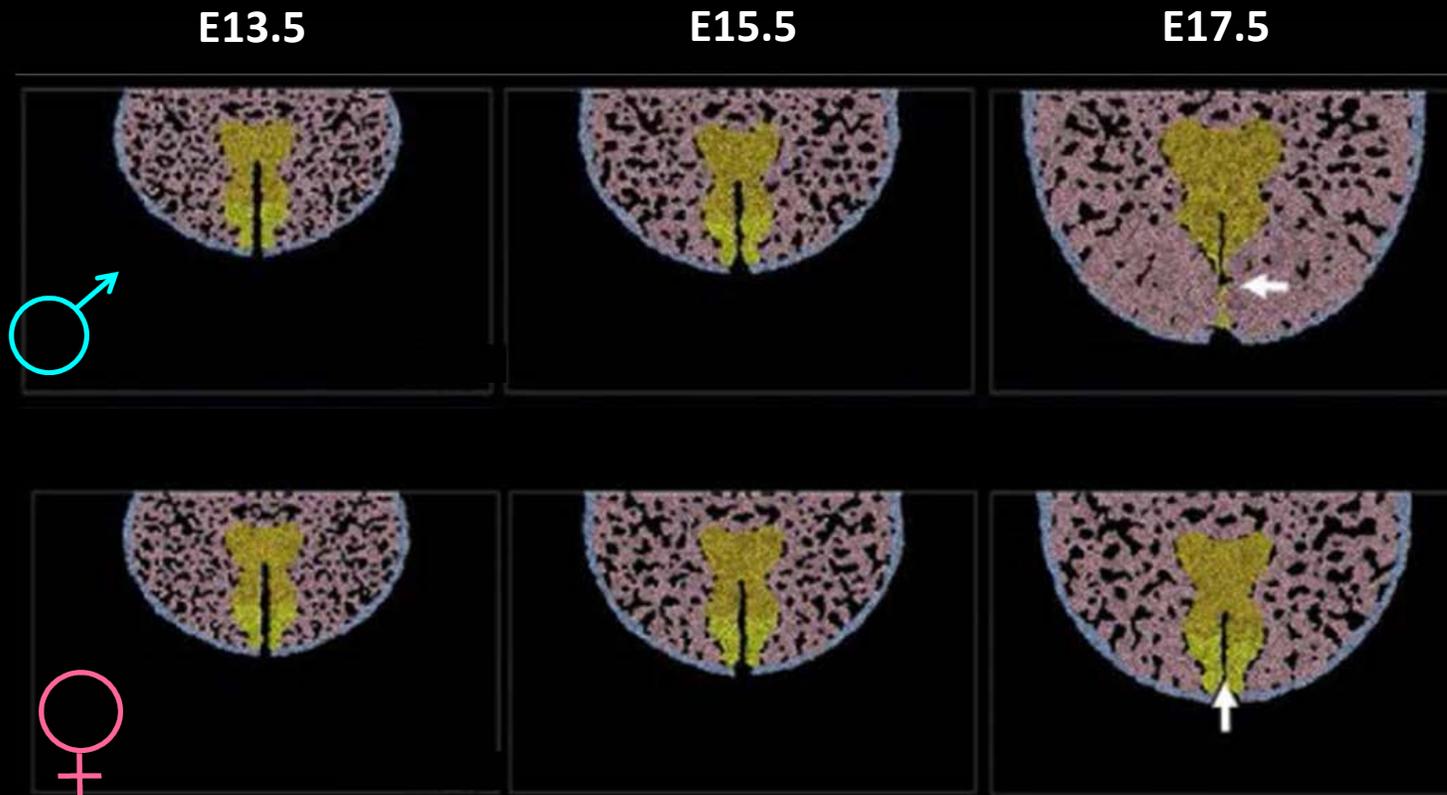
Note – model assumes androgen delivery by vascular perfusion



Androgenization: executed @ 2000 MCS as a continuous field → AR activation



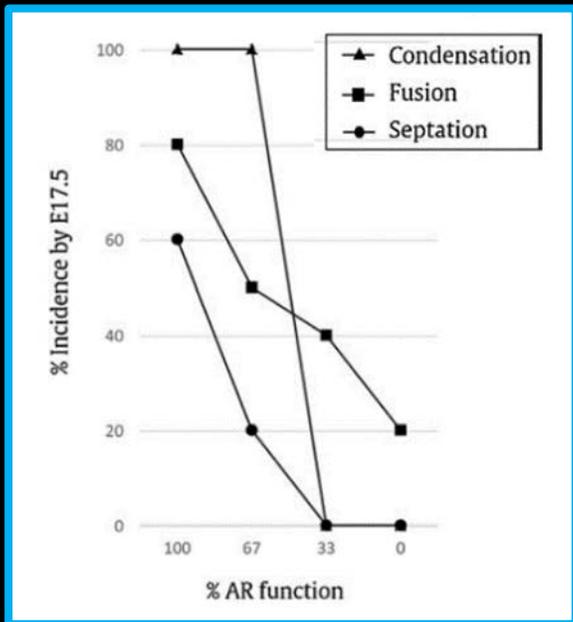
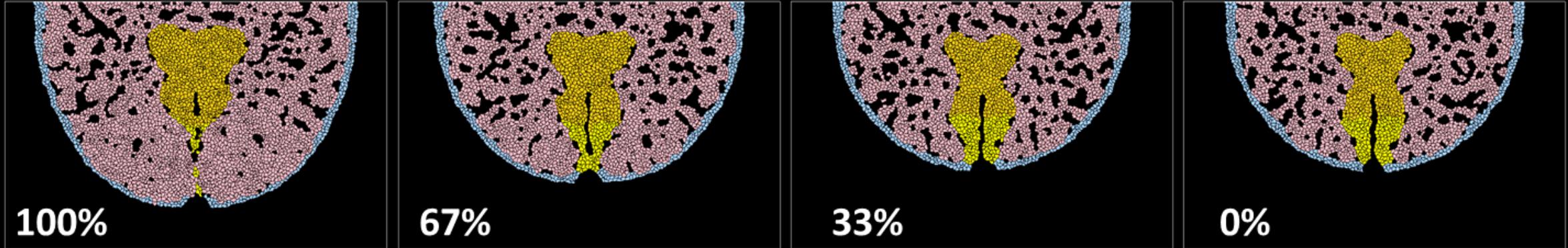
Sexual dimorphism



Urethral closure is an emergent property, driven by:

- urethral plate endoderm (apposition, contact, fusion and centralization)*
- preputial mesenchyme (proliferation, condensation, migration)*
- androgen-dependent effects mediated by FGFR2-IIIb signaling*

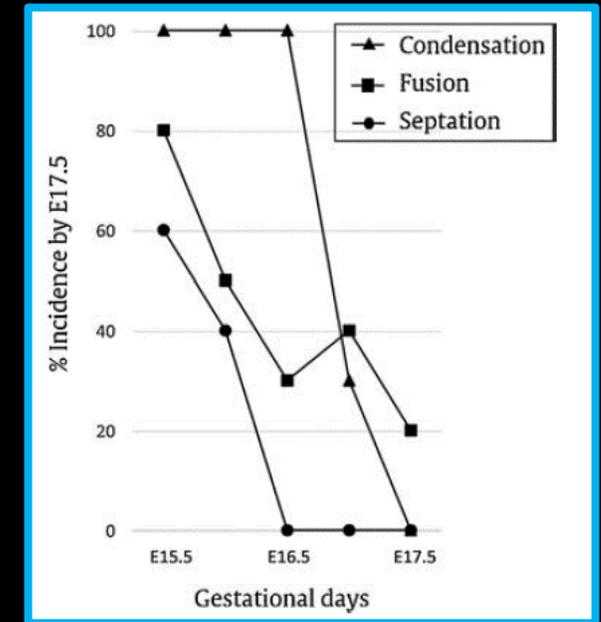
Androgen disruption: closure rates @4000 MCS \int androgen supply



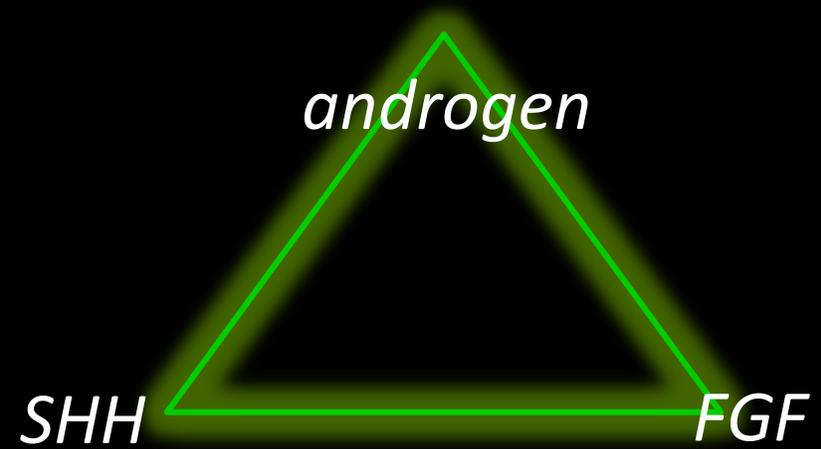
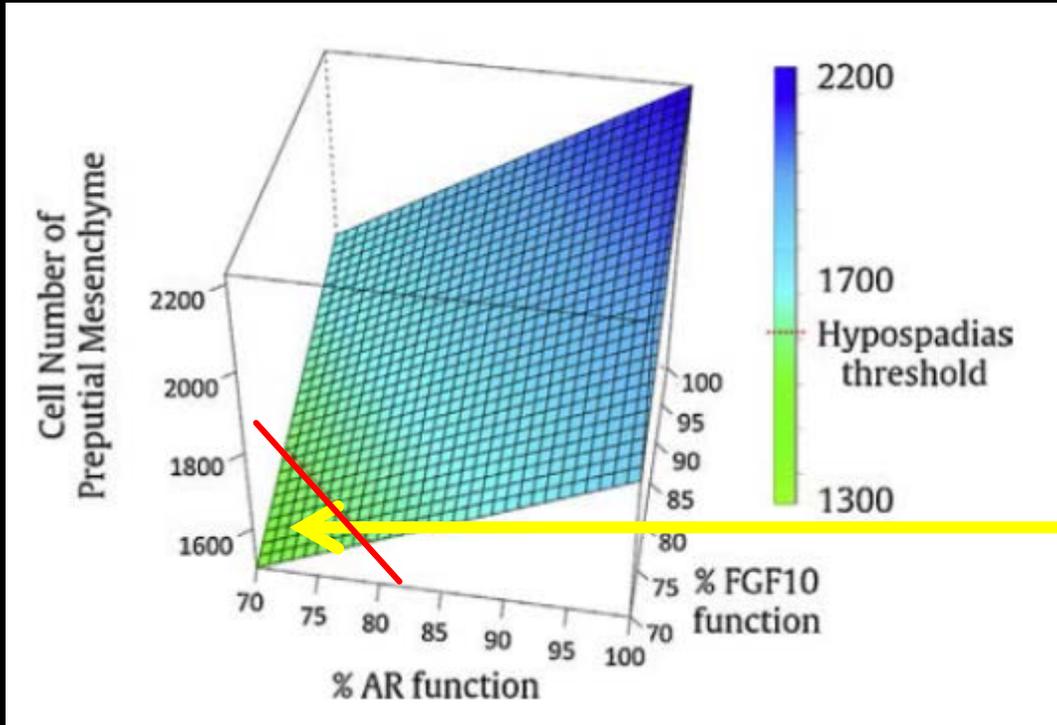
Closure indices (simulated, n=10)

LEFT: androgen insufficiency

RIGHT: delayed virulization



Gene-environment interaction (*sensitivity analysis*)



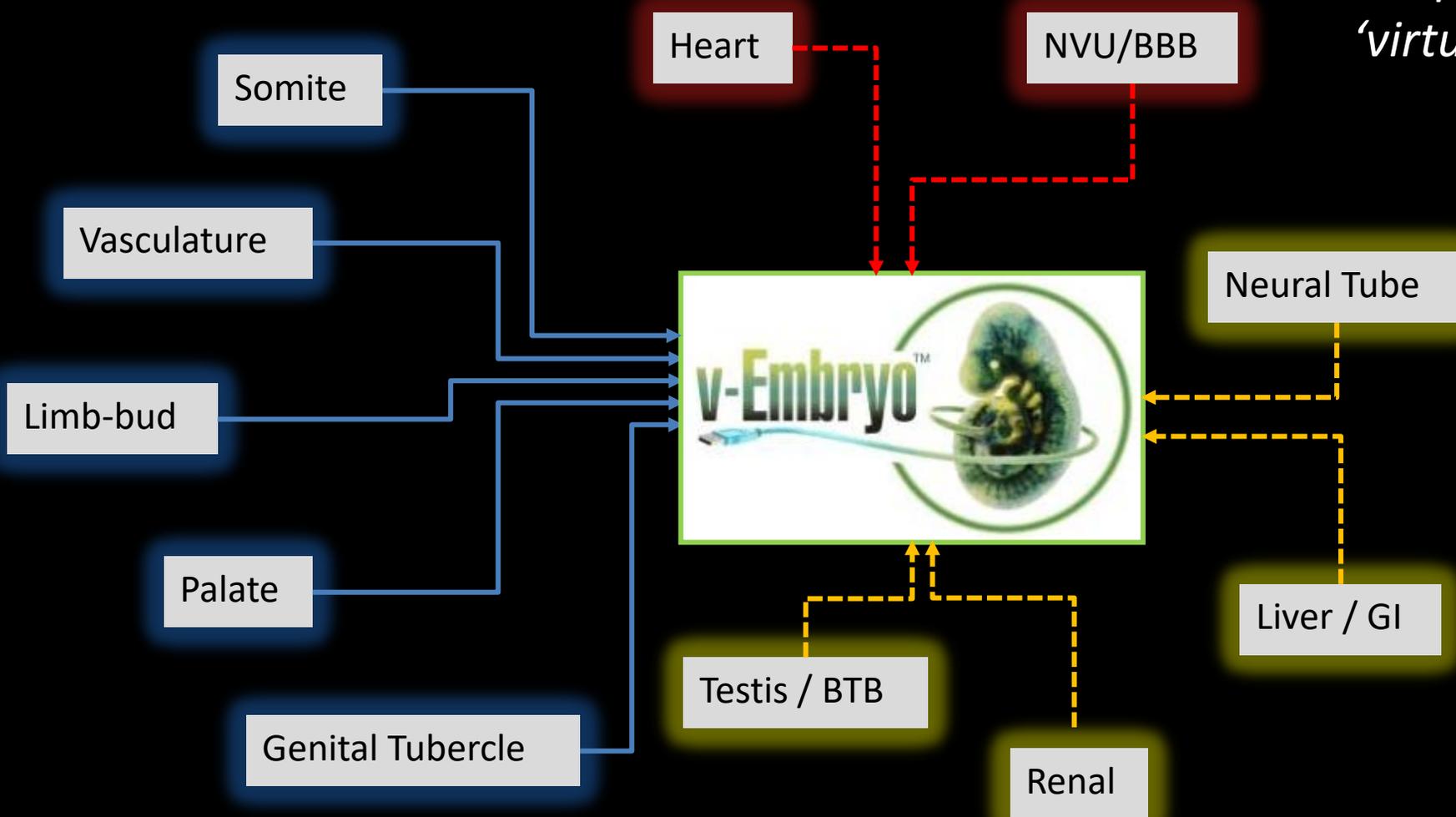
High risk for hypospadias

Multi-disturbance surface plot for individual risk factors:

- genetics (e.g., FGFR polymorphism)
- metabolism (e.g., SHH cholesterol-ification)
- environmental exposure (e.g., androgen disrupters)

Grand Challenge:

*a predictive
'virtual embryo'*



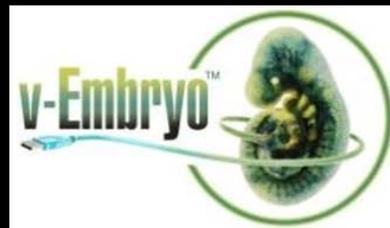
Delivered

Underway

Future

Special Thanks

- Max Leung – NCCT (now CalEPA)
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- Kyle Grode – NHEERL (now Nikon)
- Andrew Schwab – NHEERL/ISTD
- Barbara Abbott – NHEERL/TAD
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- Brian Johnson – U Wisconsin
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- Nicole Kleinstreuer – NCCT (now NTP)
- George Daston – Procter & Gamble Co.
- Ashley Seifert – U Kentucky
- Martin Cohn - (U Florida)



http://www2.epa.gov/sites/production/files/2015-08/documents/virtual_tissue_models_fact_sheet_final.pdf