



Computational Embryology:

Agent-Based Modeling and Simulation of External Genital Development

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"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.*

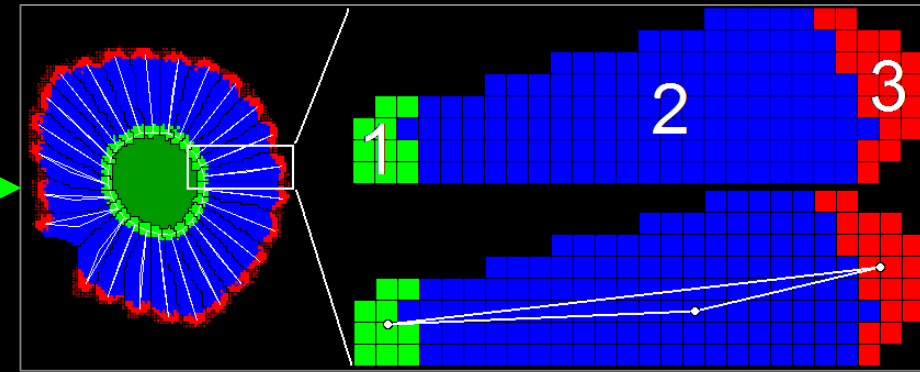
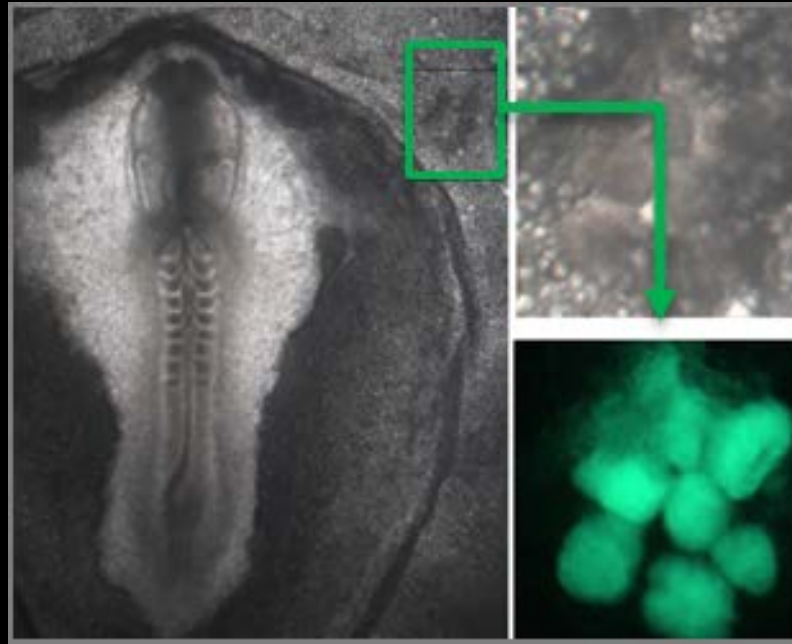
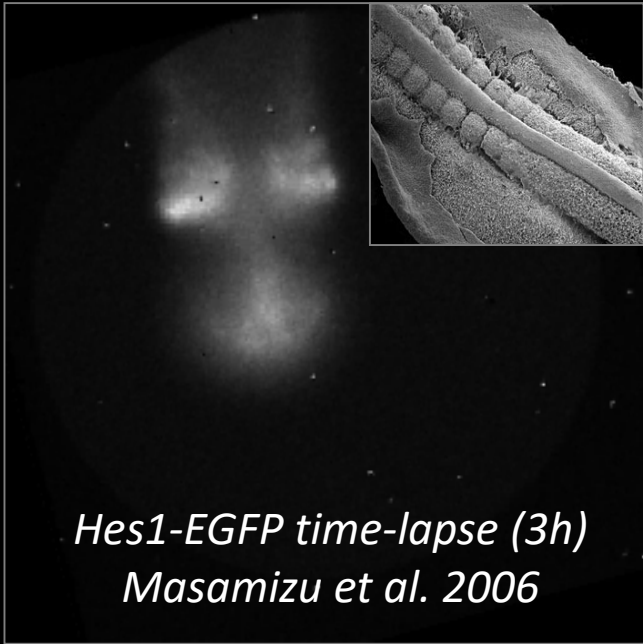
Agent-Based Models (ABMs):

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*)

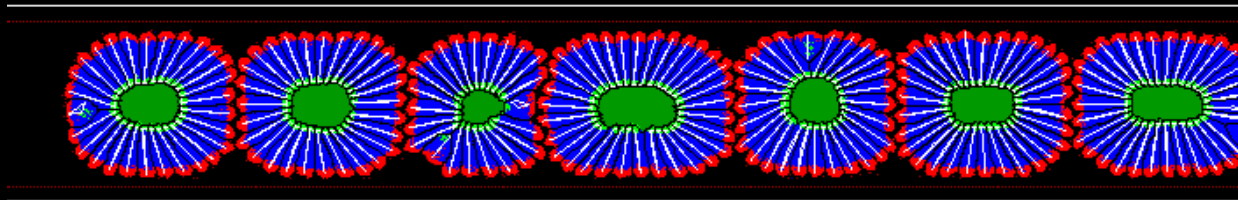


Modeling somite development



Differential cell adhesion

- clock genes do not oscillate
- somites form simultaneously

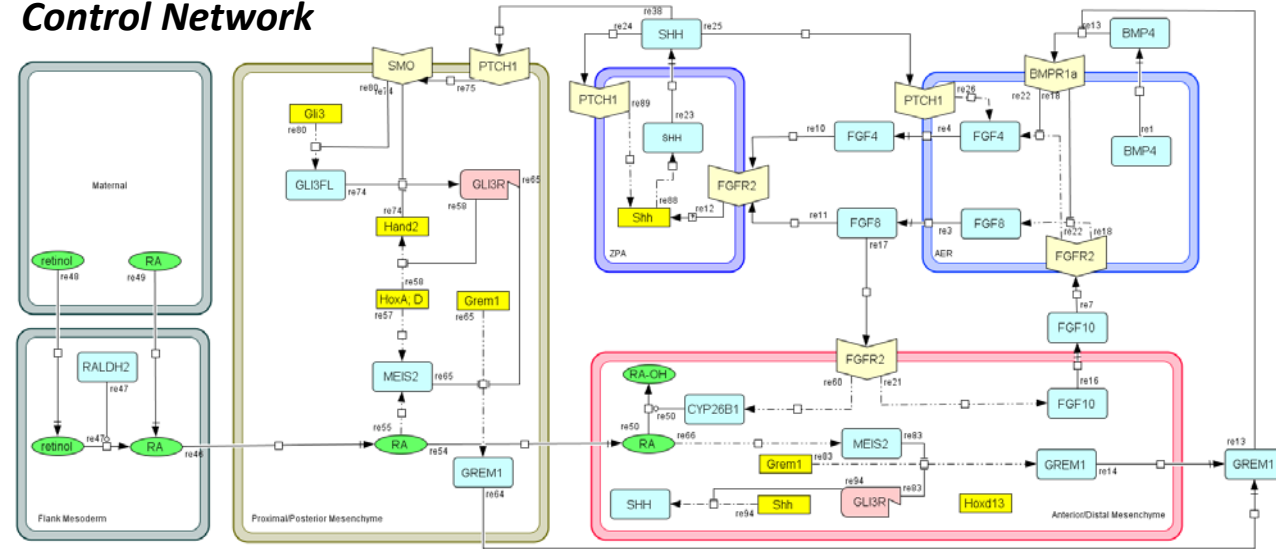


- *FGF8 wavefront restores sequentiality*
- *oscillatory clock improves regularity*

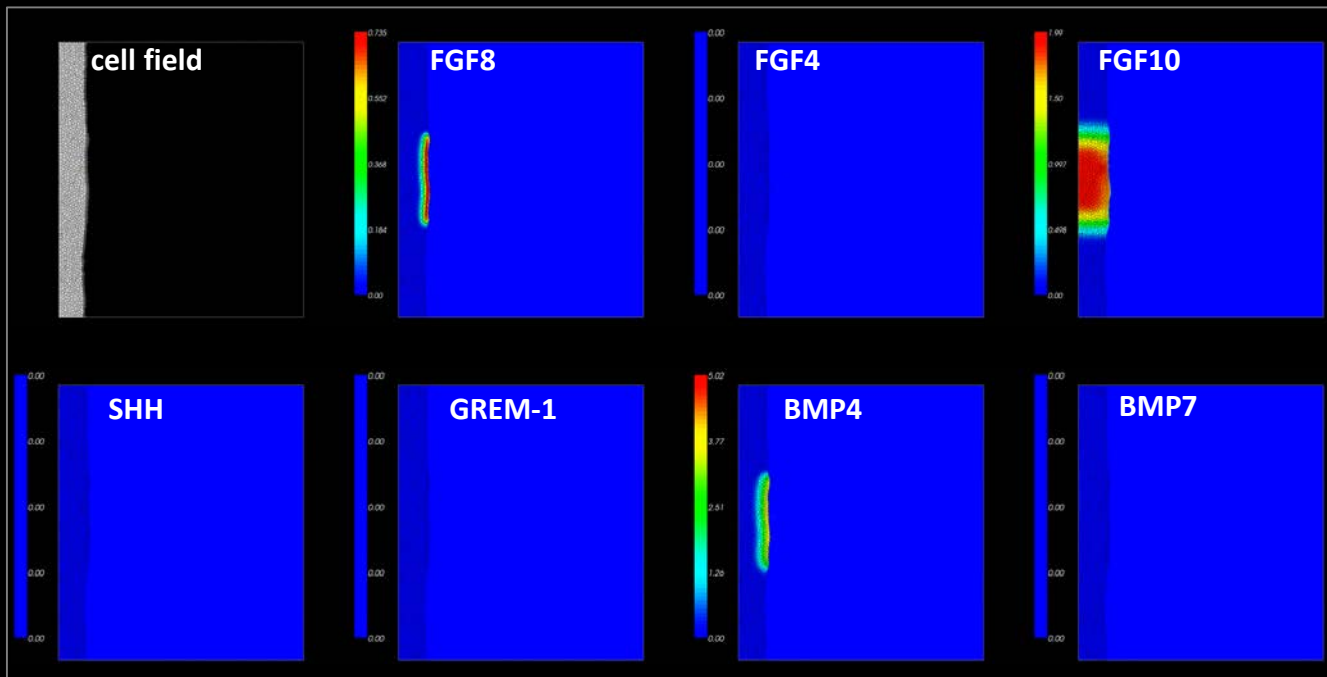
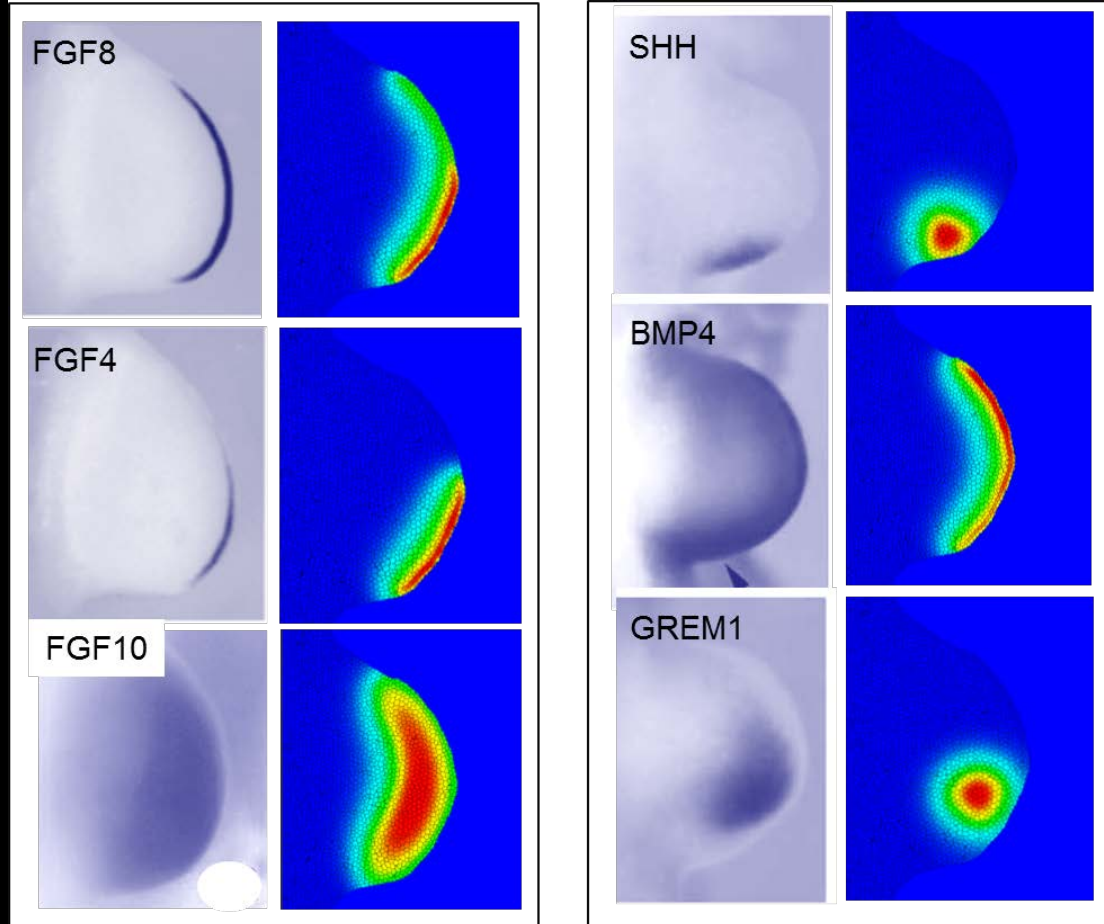
SOURCE: Dias et al. (2014) Science



Control Network

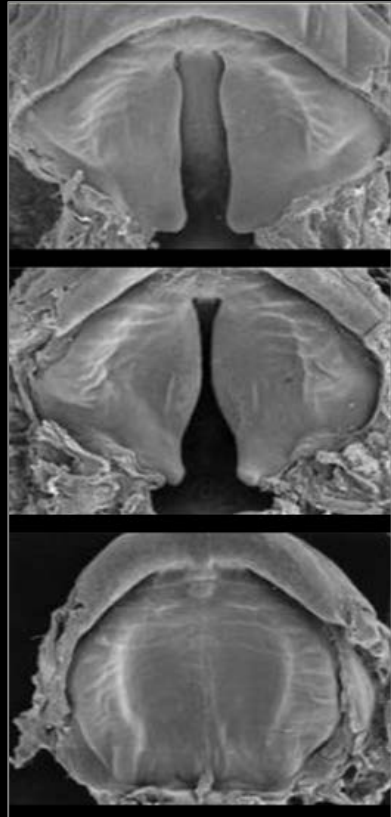


Limb-bud outgrowth



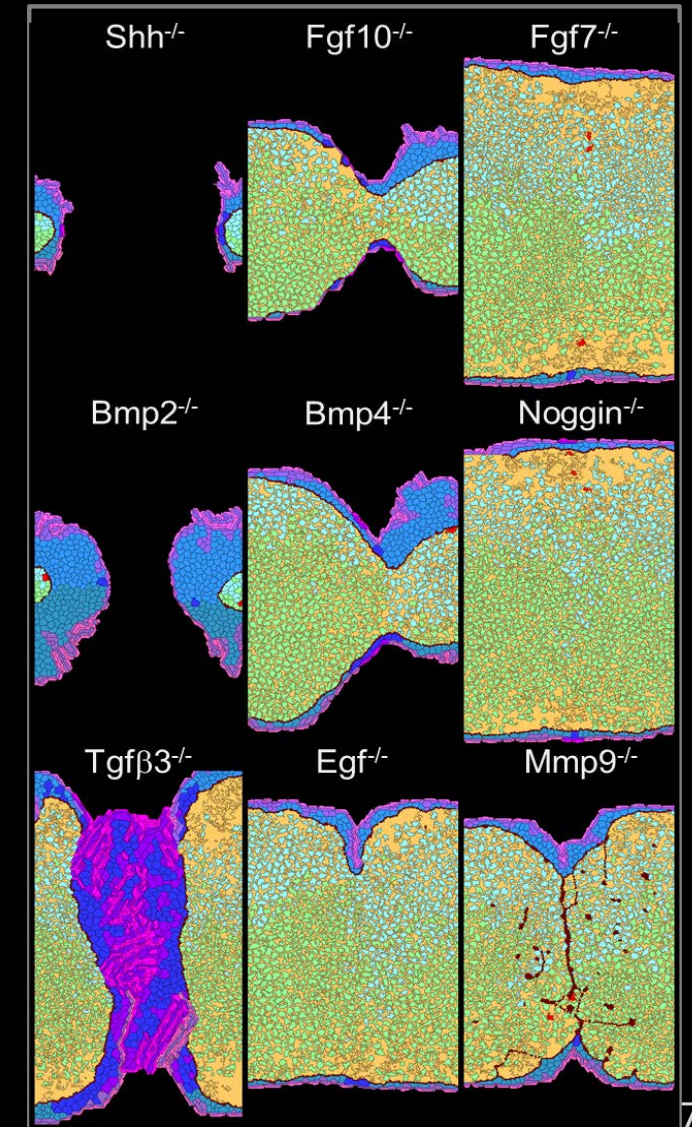
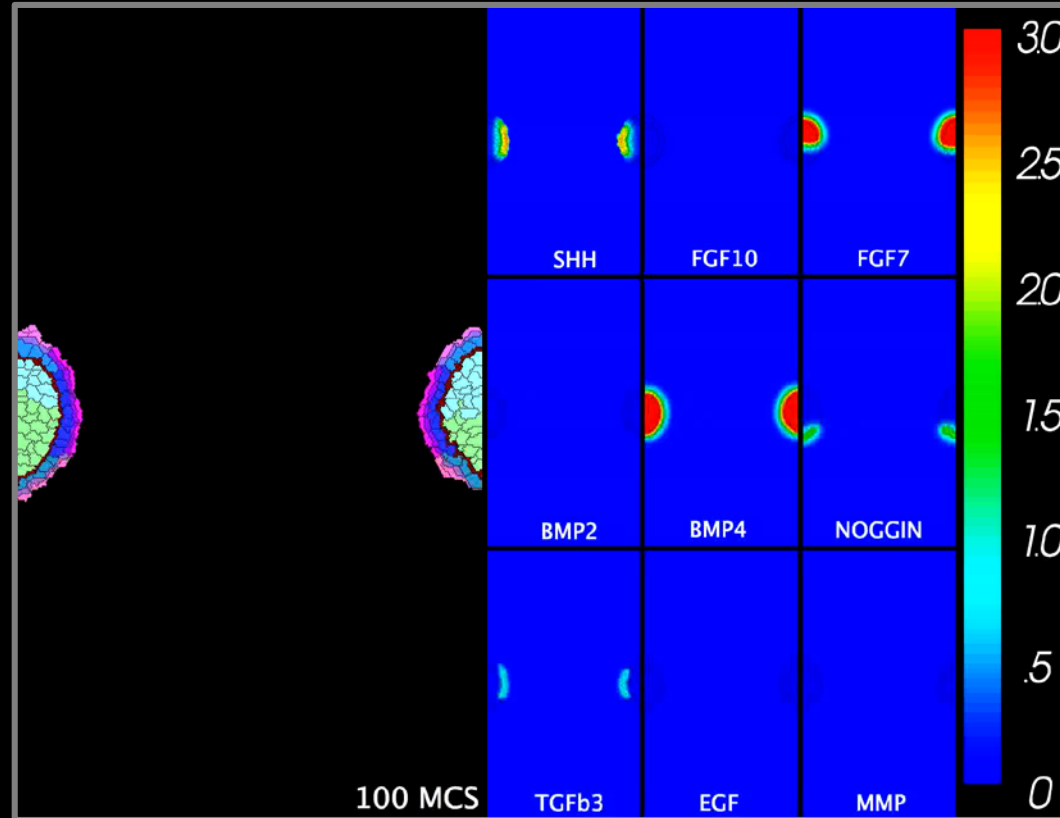
Palatal closure: driven by medial edge epithelium (MEE) seam breakdown

Palatal fusion *in vivo*



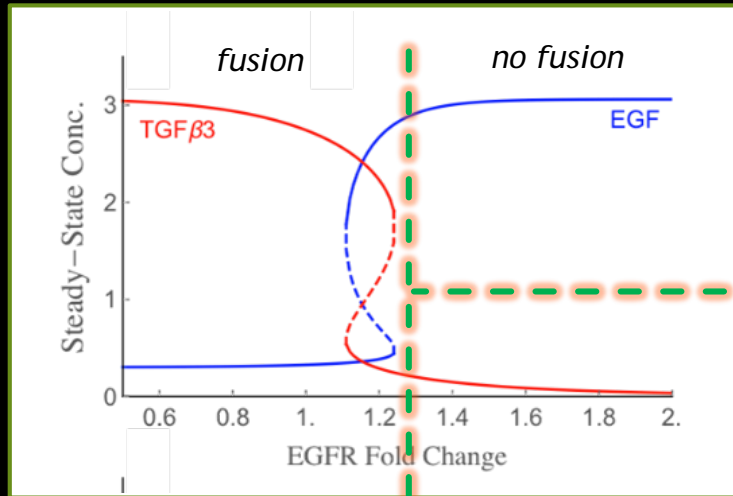
SEMs of human palate by K Sulik, UNC

Palatal fusion *in silico*



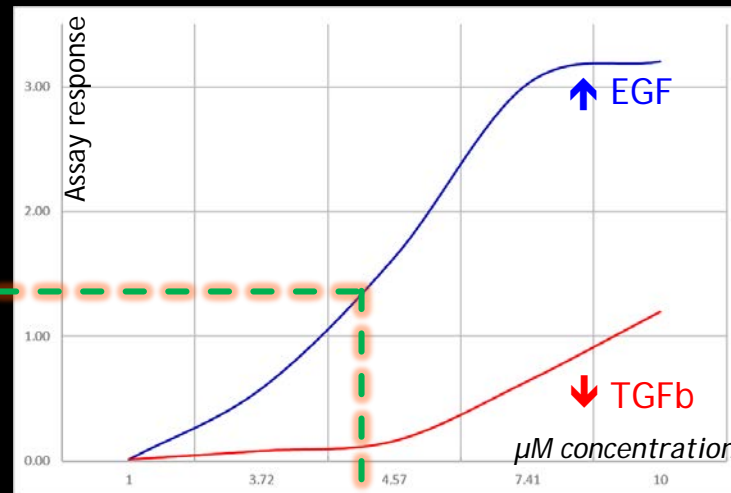
TGF-beta/EGF latch switch: *controls MEE breakdown*

INPUT: switch dynamics



tipping point predicted by
computational dynamics
(hysteresis switch)

Captan in ToxCast



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

Captan in ToxRefDB

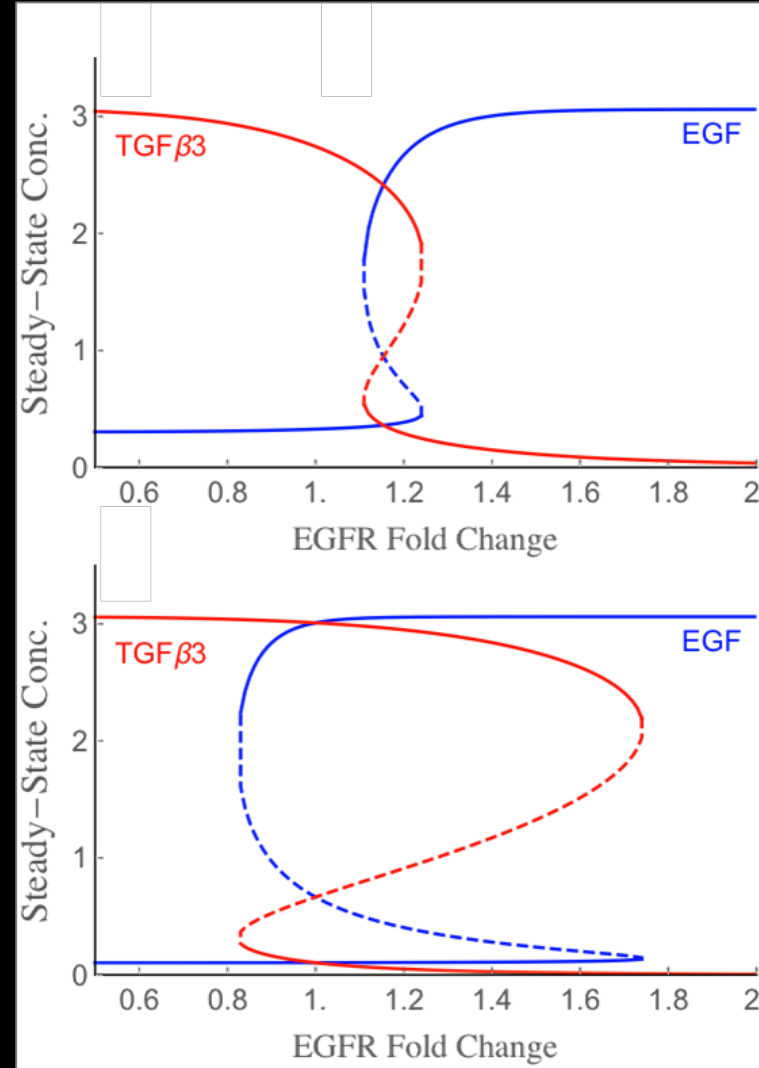
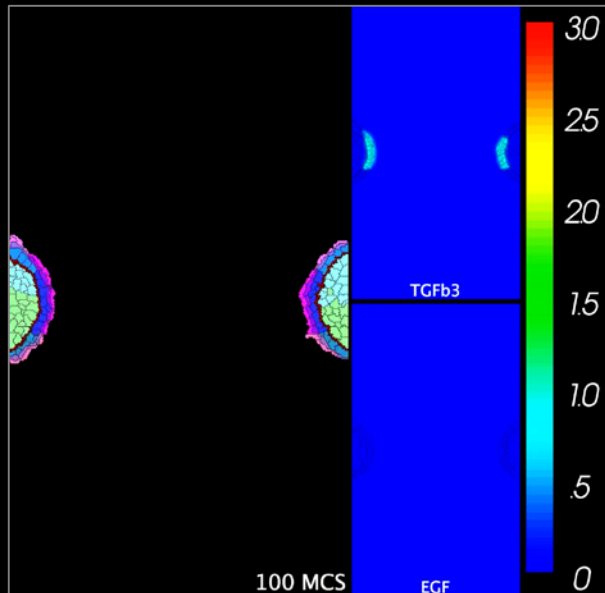
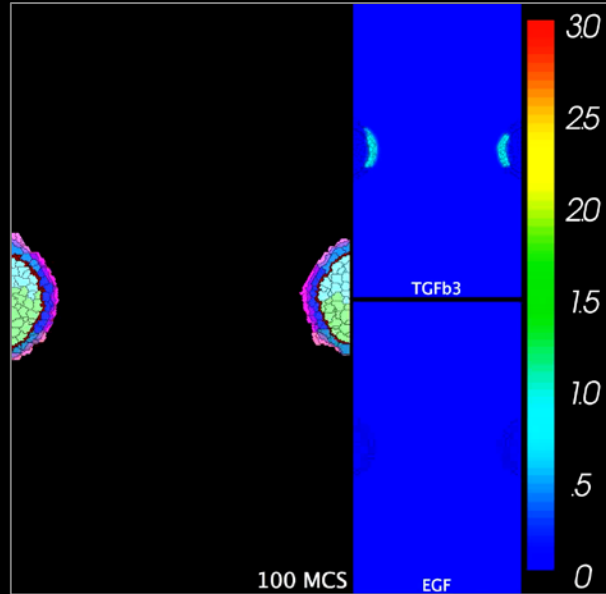
NOAEL = 10 mg/kg/day

LOAEL = 30 mg/kg/day

human HTTK model

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

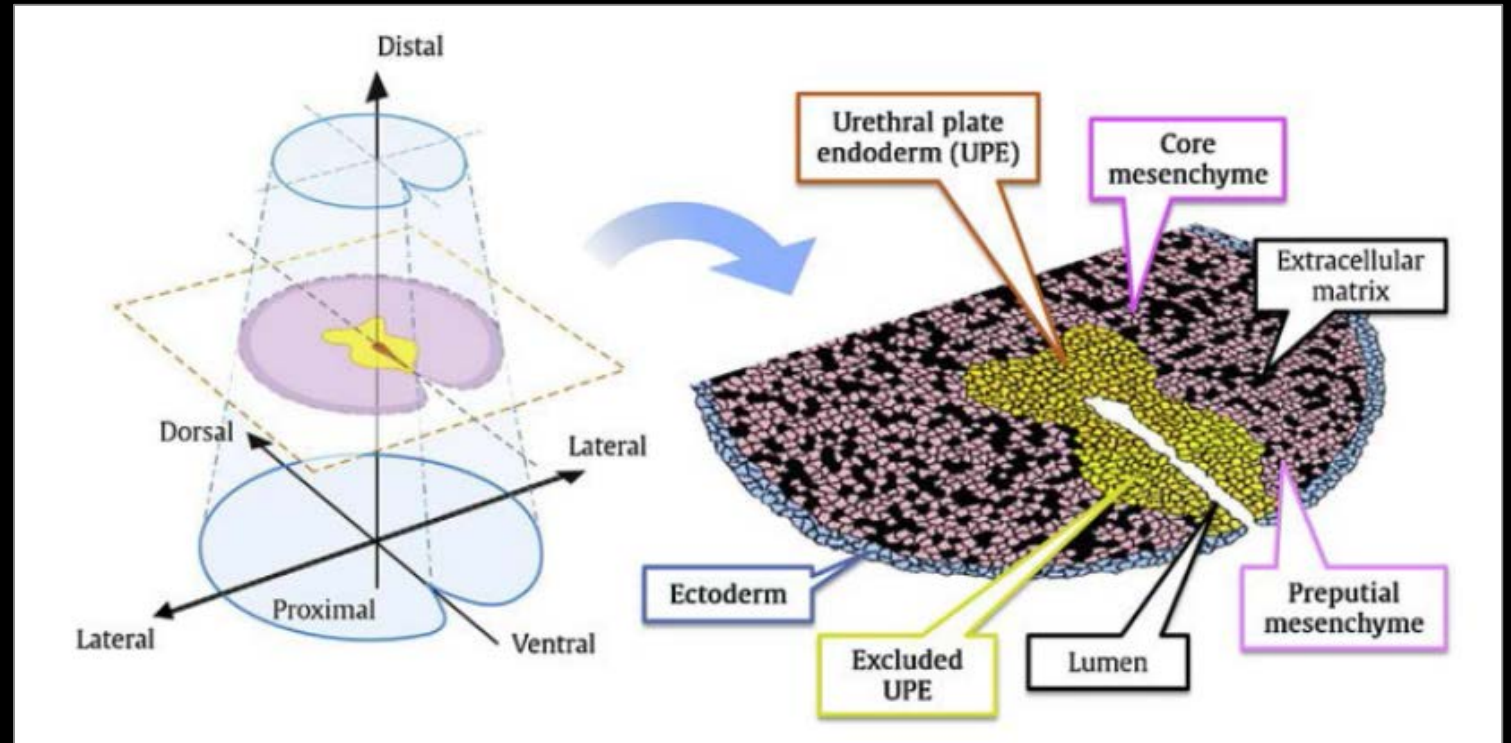
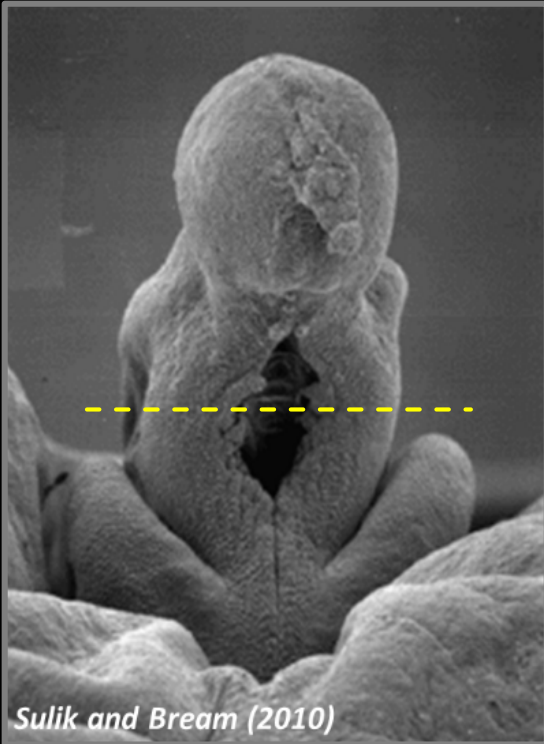
Messin' with the switch: *two scenarios for bistable dynamics*



Narrow hysteresis:
*less resilient
but reversible*

Broad hysteresis:
*more resilient
but irreversible*

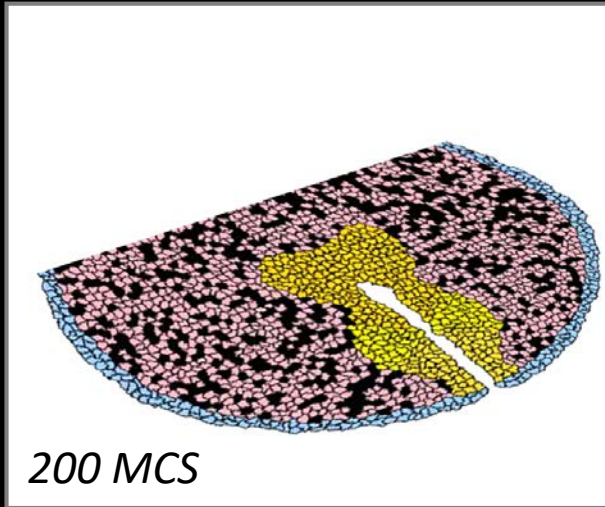
Genital tubercle: abstraction for ABM-simulated urethral closure



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



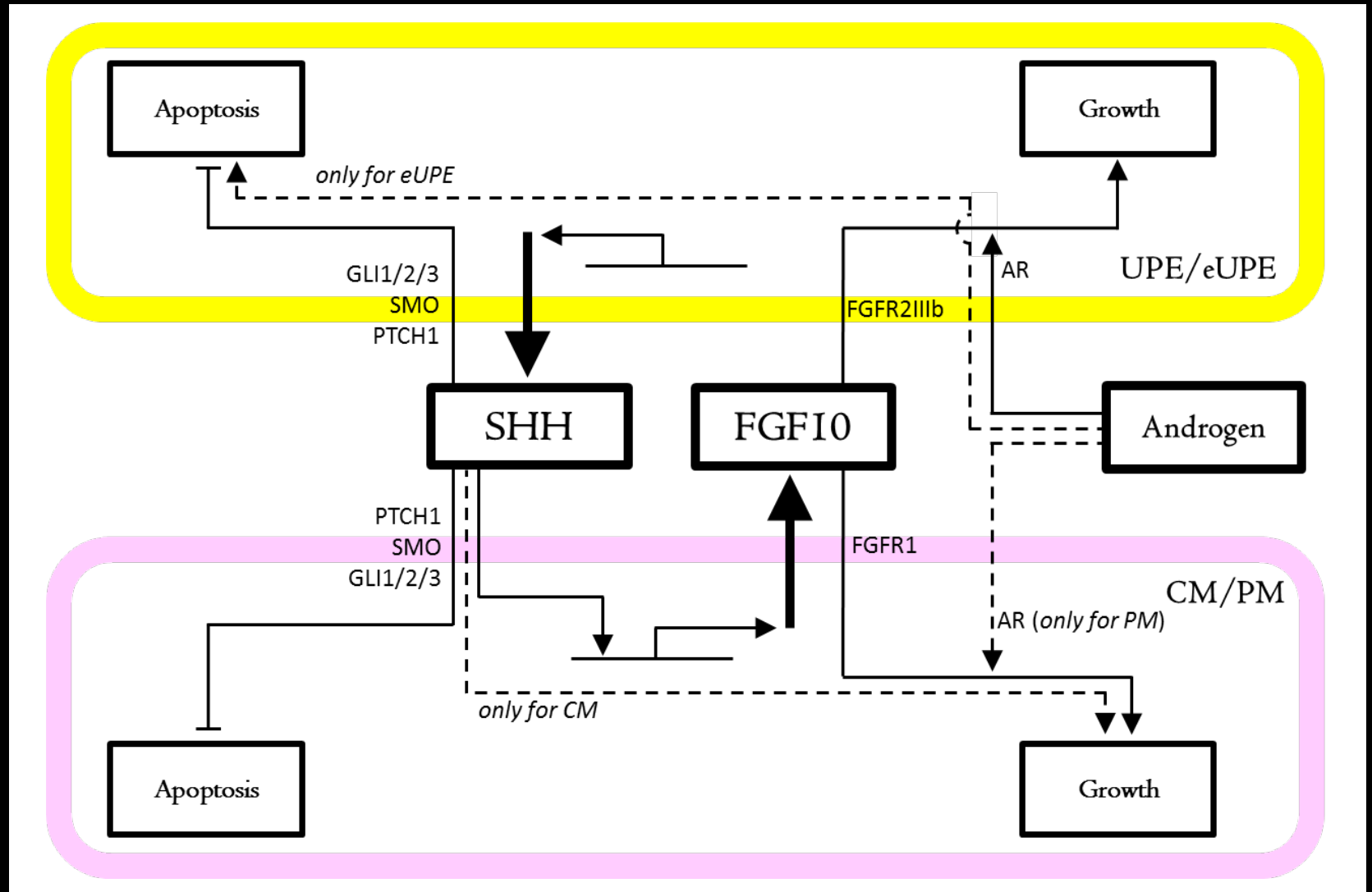
Biokinetic gradients:

FGF10 – simple diffusion

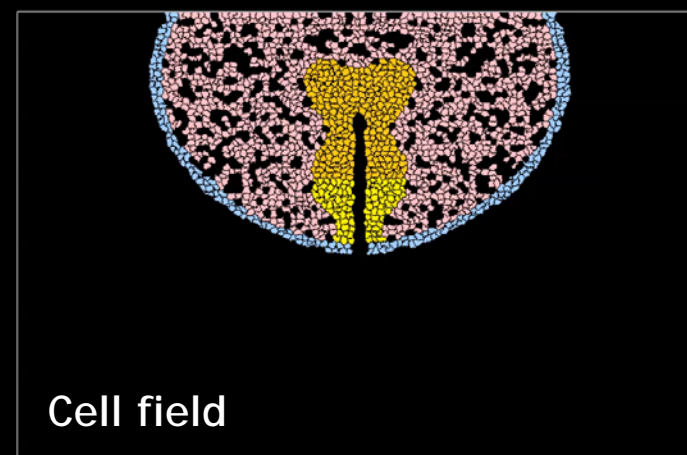
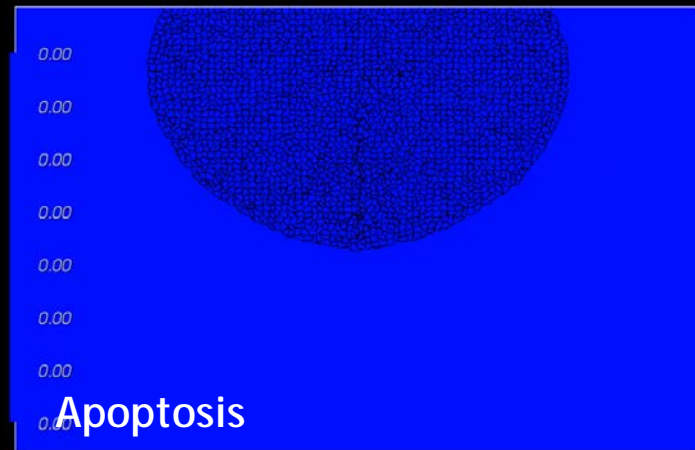
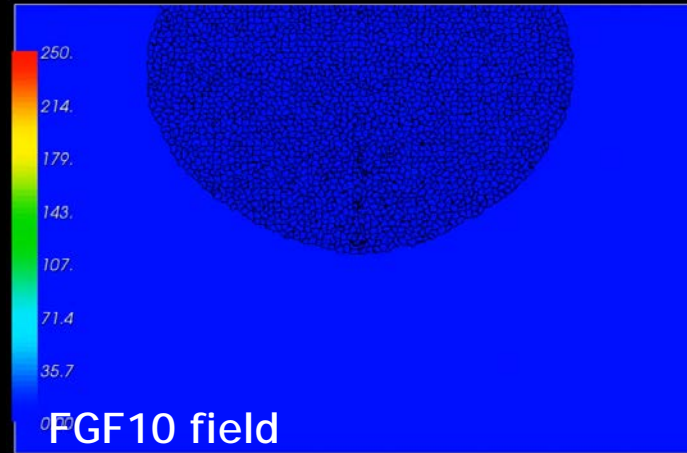
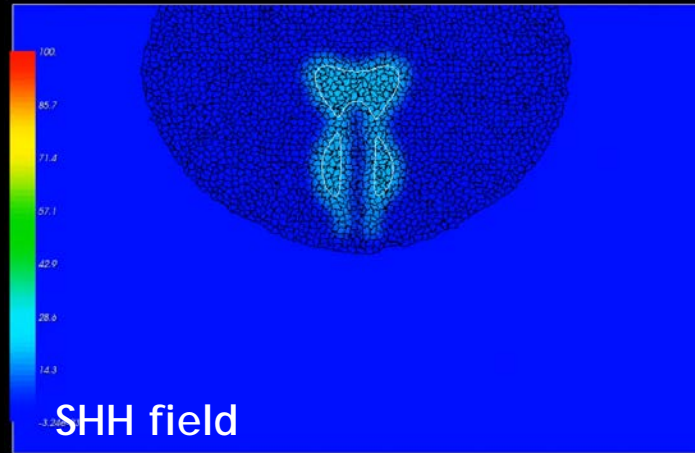
SHH – transcytosis (1/8 rate)

An – latched @2000 MCS (E15.5)

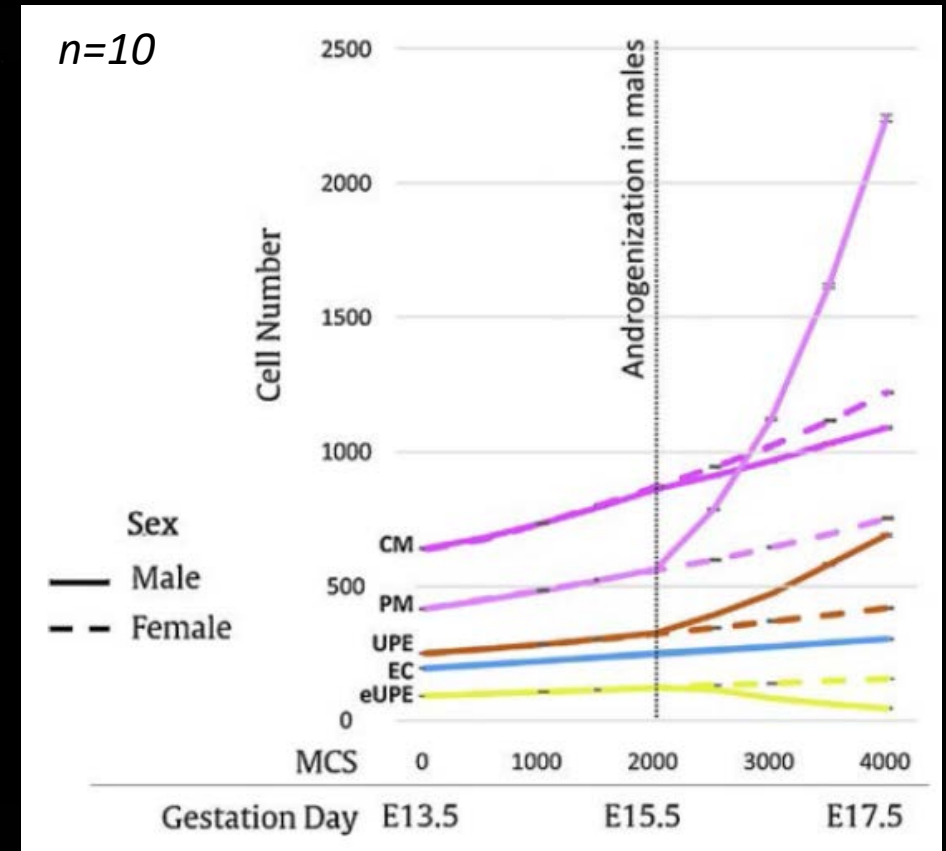
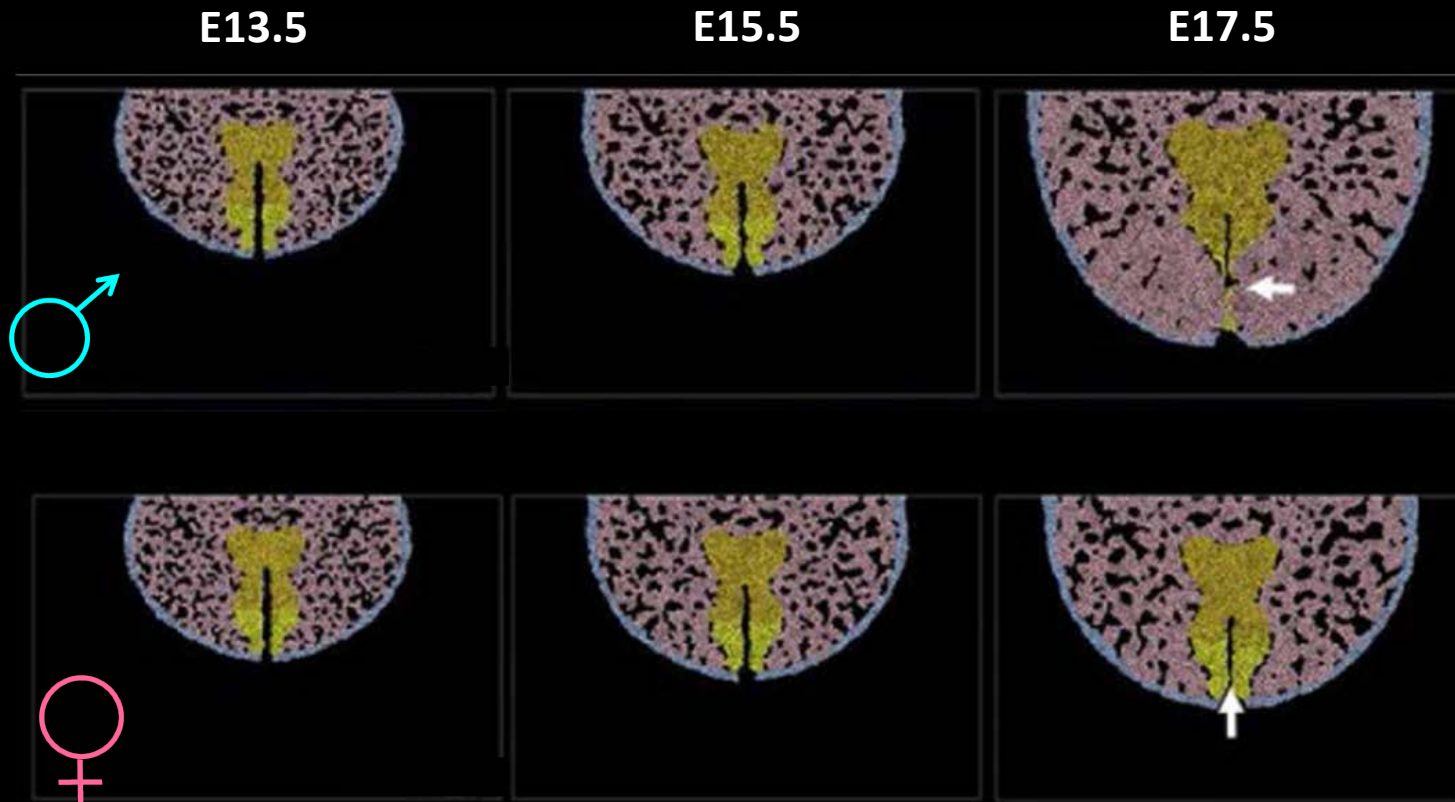
Note – model assumes androgen delivery by vascular perfusion



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



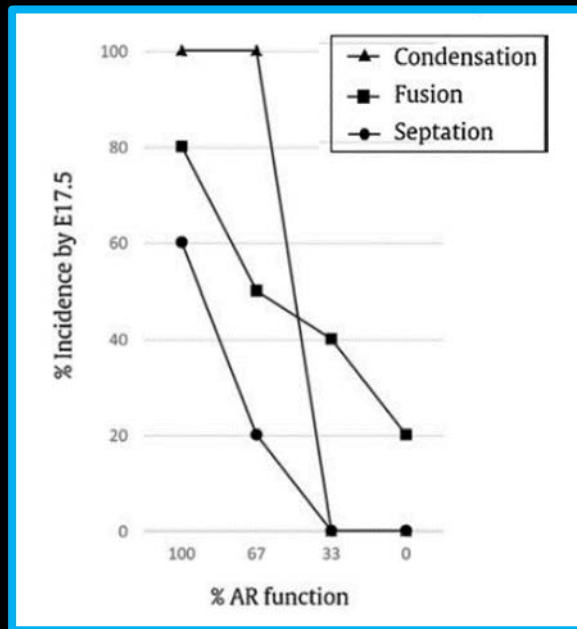
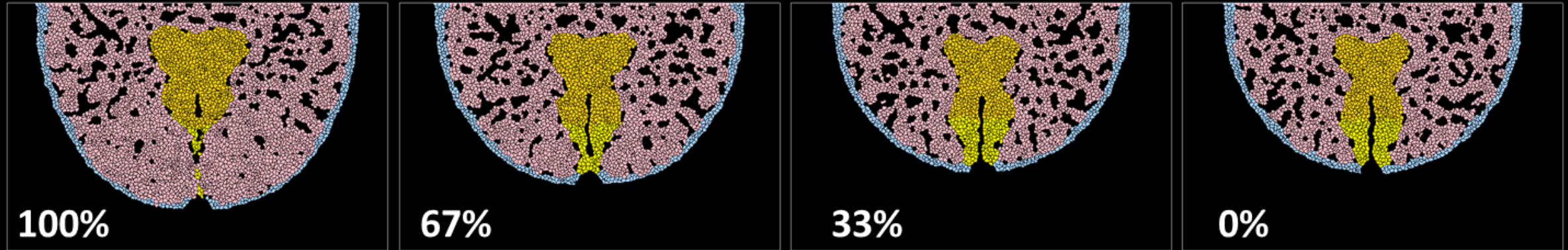
Sexual dimorphism



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*

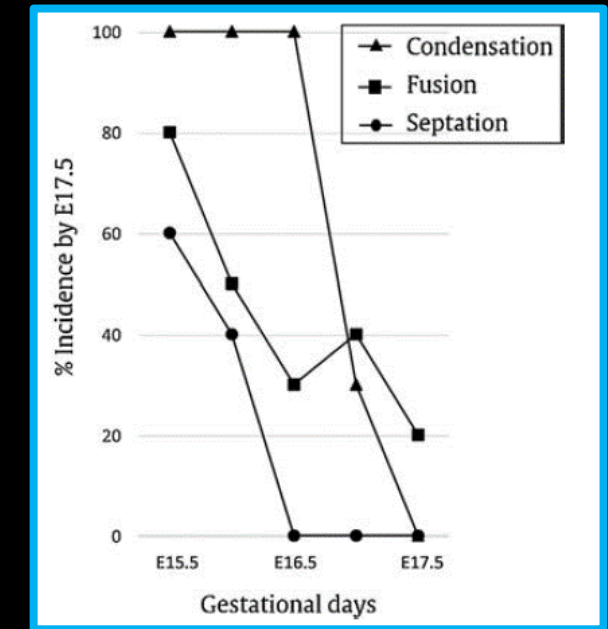
Androgen disruption: closure rates @4000 MCS \int androgen supply



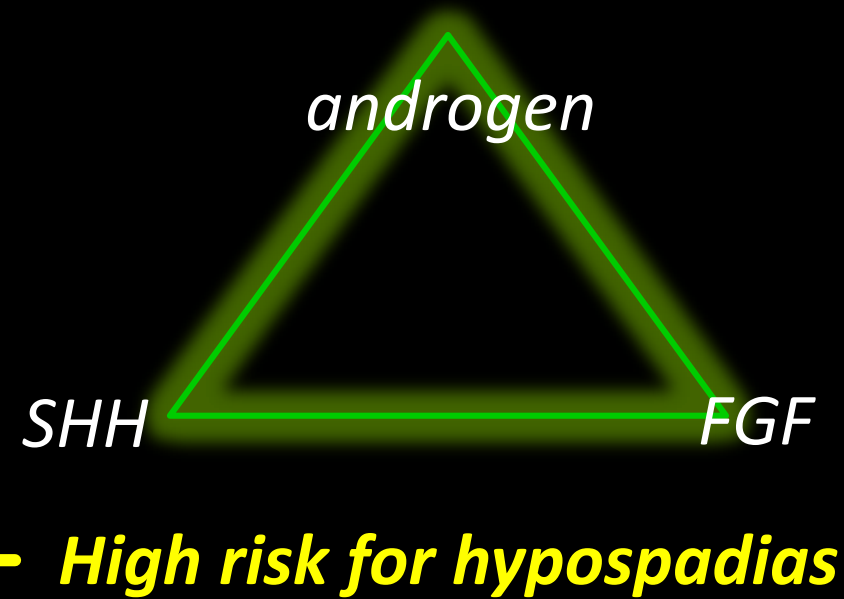
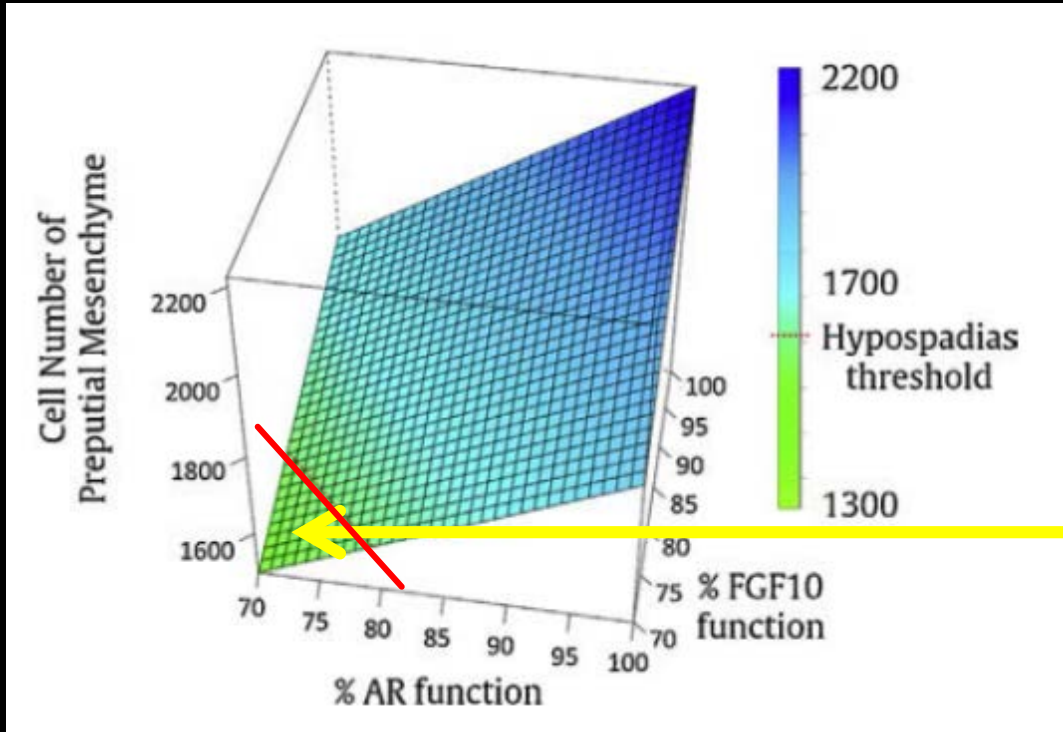
Closure indices (simulated, n=10)

LEFT: androgen insufficiency

RIGHT: delayed virulization



Gene-environment interaction (*sensitivity analysis*)

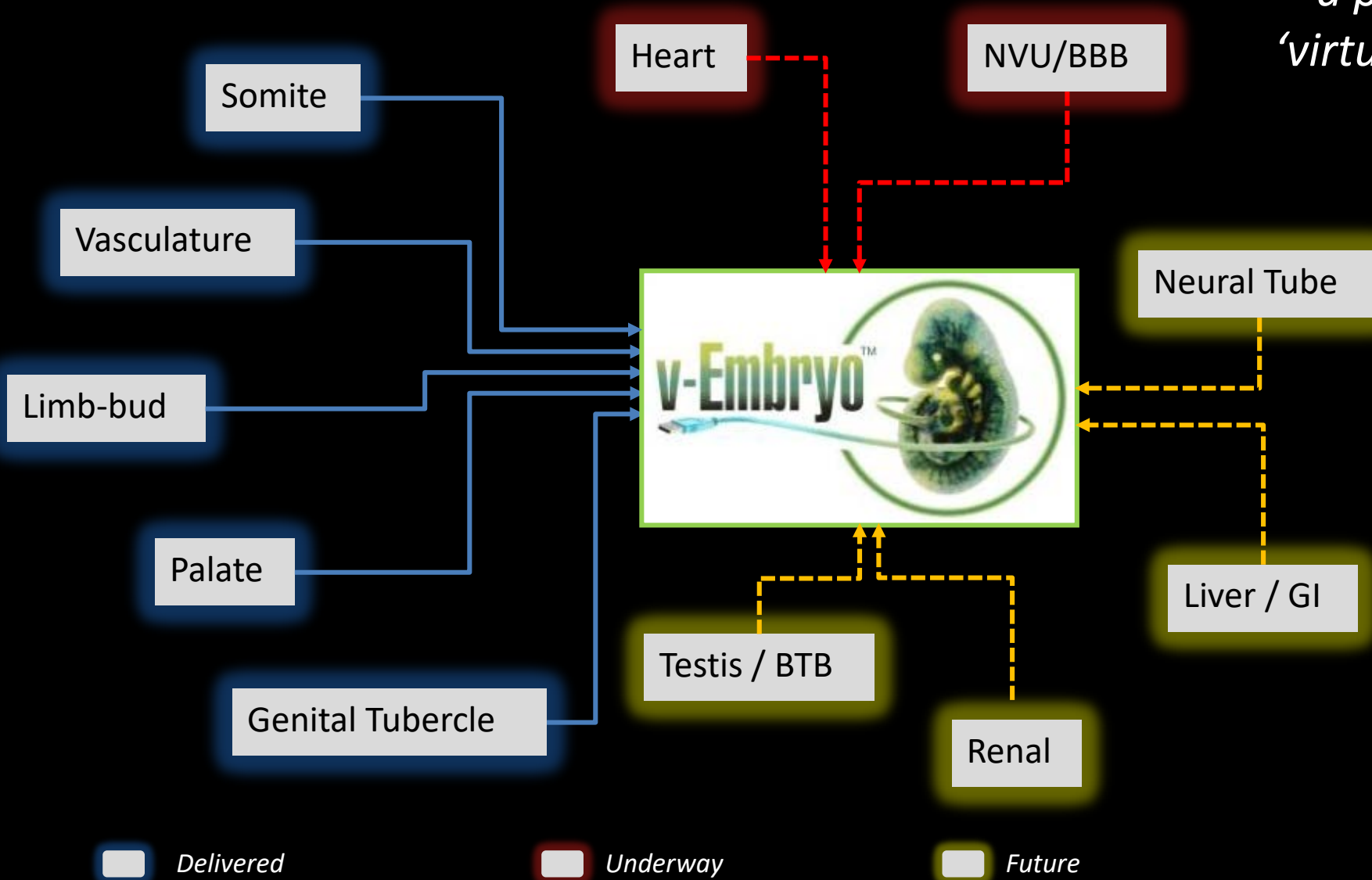


Multi-disturbance surface plot for individual risk factors:

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

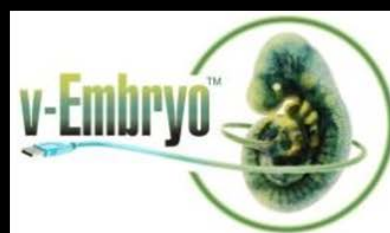
Grand Challenge:

*a predictive
'virtual embryo'*



Special Thanks

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- 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