

Toxicological Tipping Points & Cell Stress

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Discriminating between adaptation & adversity
SOT 2016, New Orleans, LA

March 15, 2016

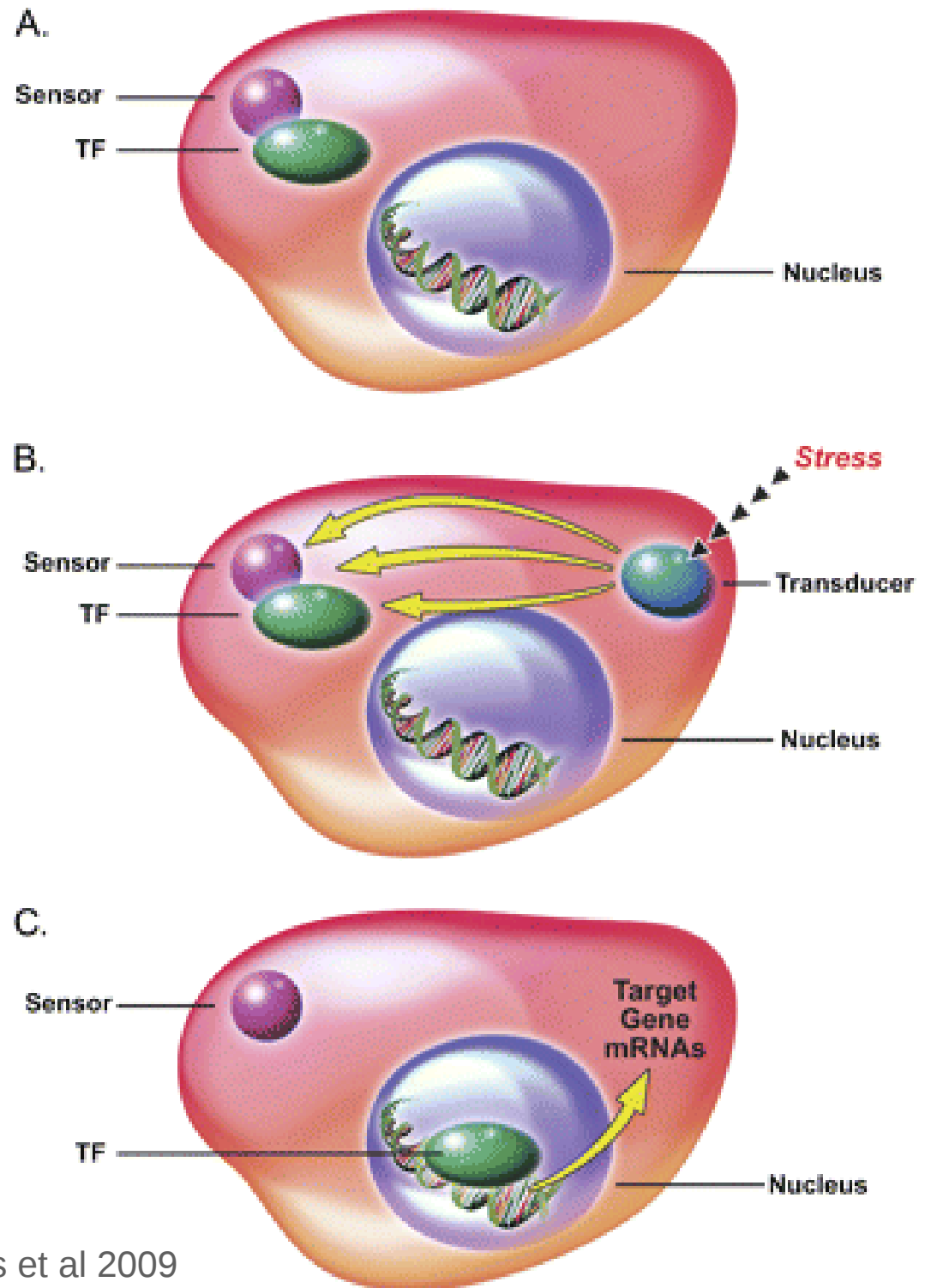
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Toxicological Tipping Points ...

- ... threshold between adaptation and adversity
- ... **may** be useful as PoD for risk assessment
- ... **may** involve stress response pathways
- ... **could** be found from **some** *in vitro* data
- ... needs new computational approaches

➤ **Case-Study: Use high-content data to find tipping points**

- ❑ Cellular stress response pathways
 - ❑ Oxidative stress
 - ❑ Hypoxia
 - ❑ DNA damage
 - ❑ ER stress
 - ❑ Mitochondrial stress
- ❑ Induced by chemicals to overcome “perturbation” of normal state



HCI → Stress Responses?

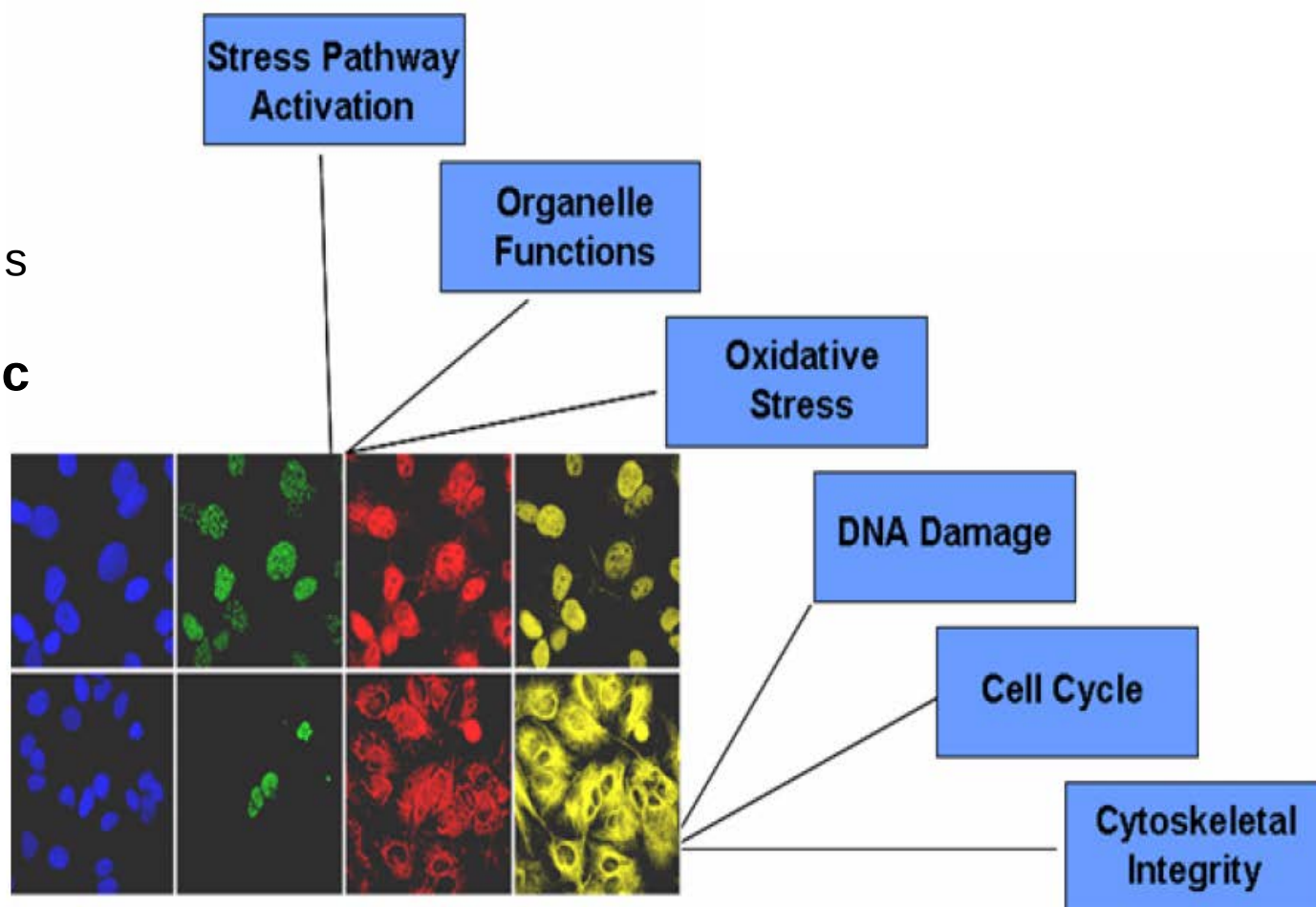
□ High content imaging

- HepG2 cell culture
- (ToxCast I, II)
- 10 conc, 3 times

□ HCI Assays

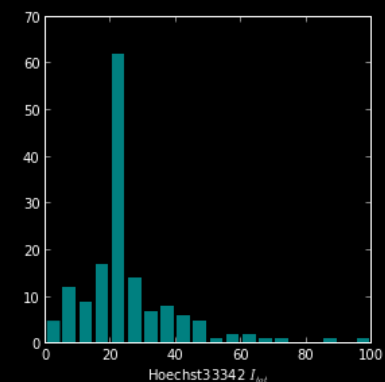
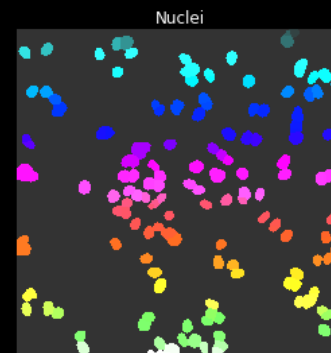
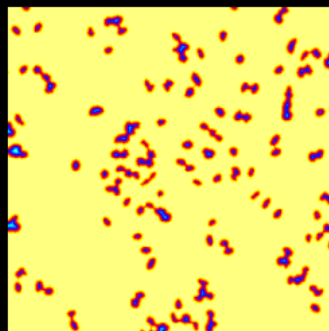
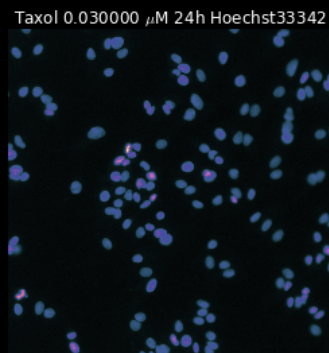
- System perturbation
- Cell health
- Some stress pathways

□ Dynamic phenotypic response of cells to chemicals

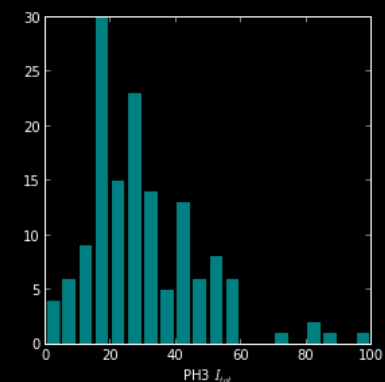
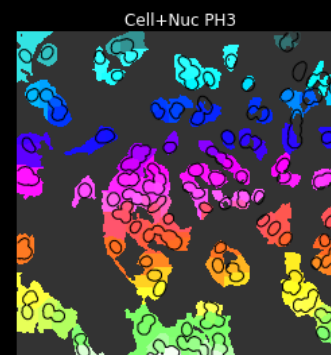
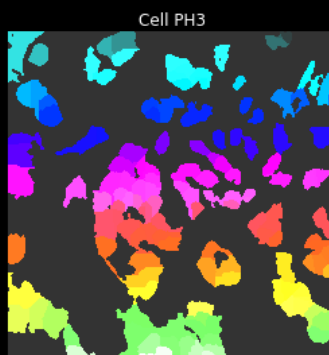
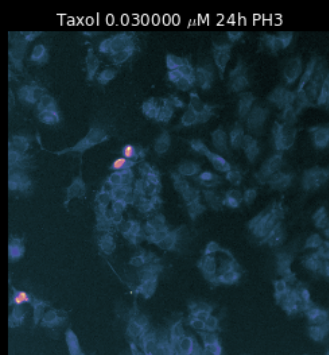


Taxol 0.03uM

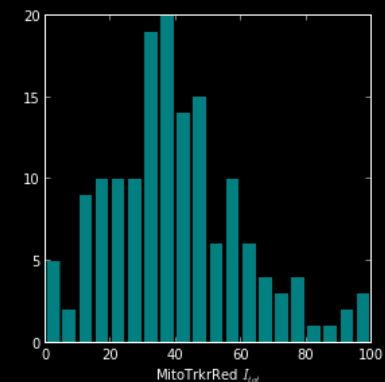
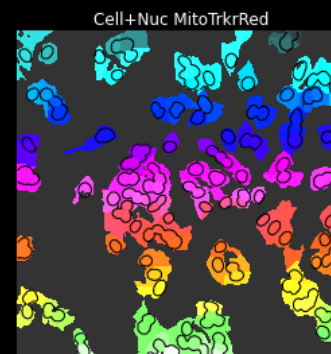
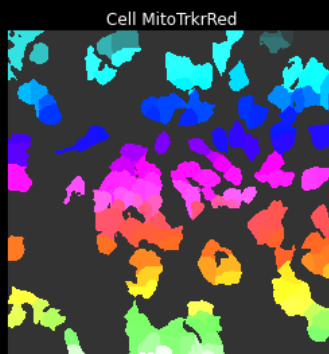
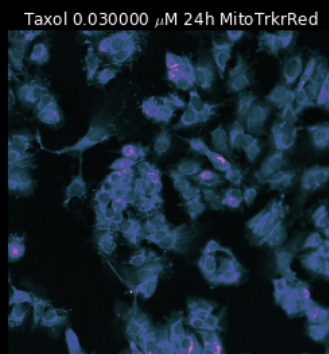
Hoechst33342



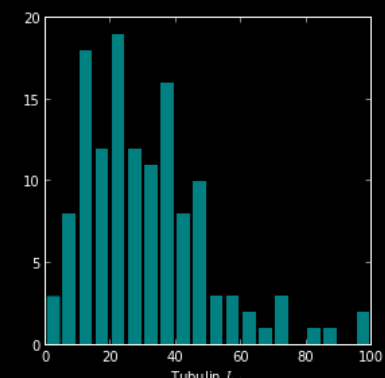
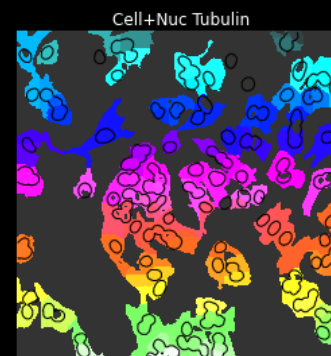
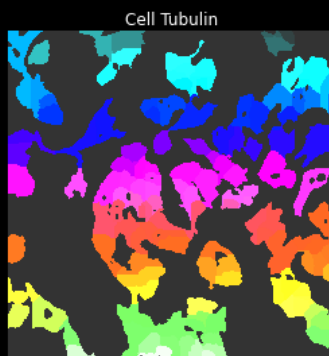
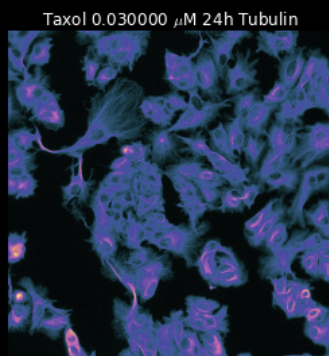
Phospho-Histone3



MitoTracker Red

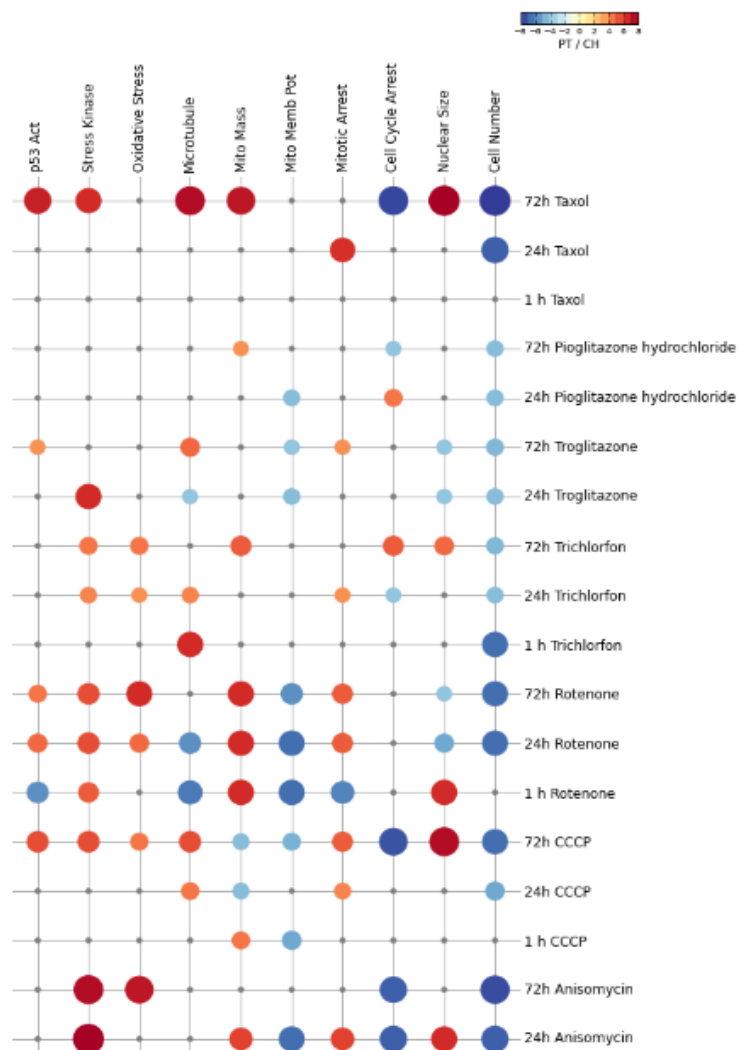


Phospho-Tubulin

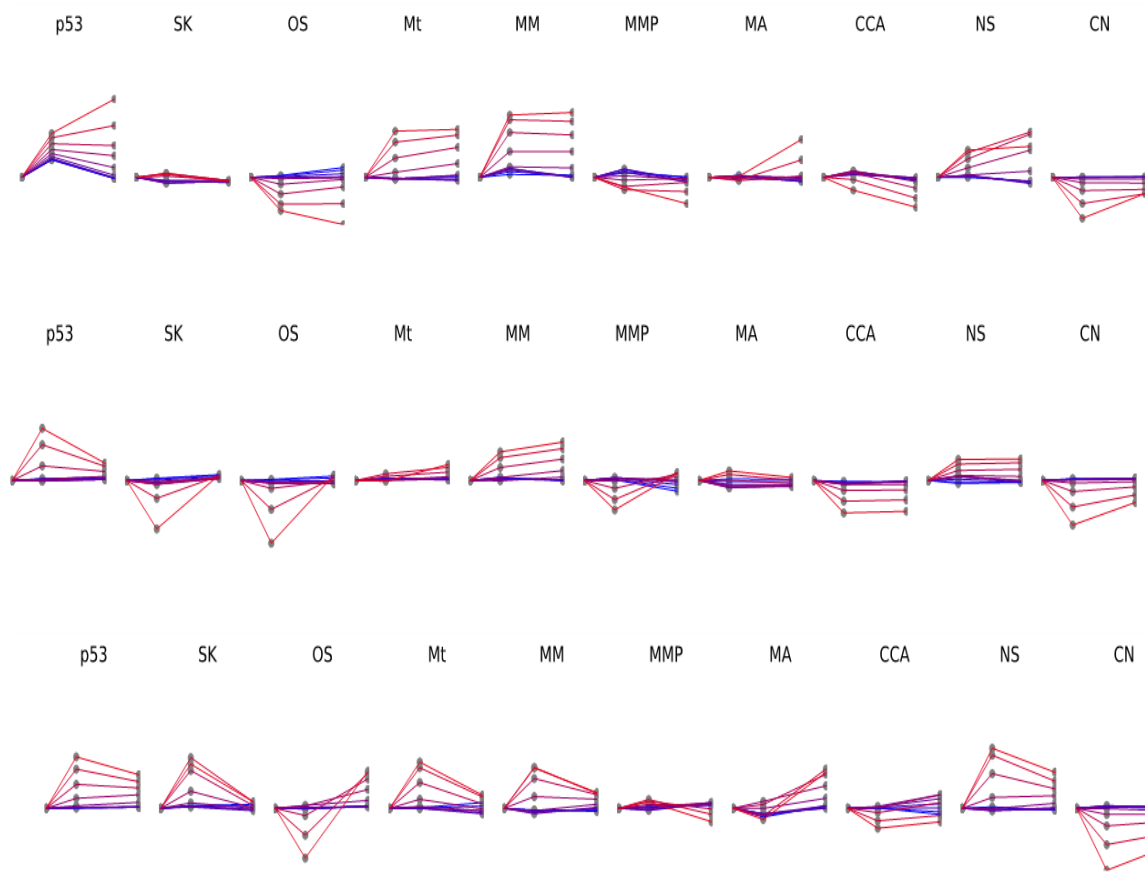


HCI → Multidimensional Data

“Discrete Responses”

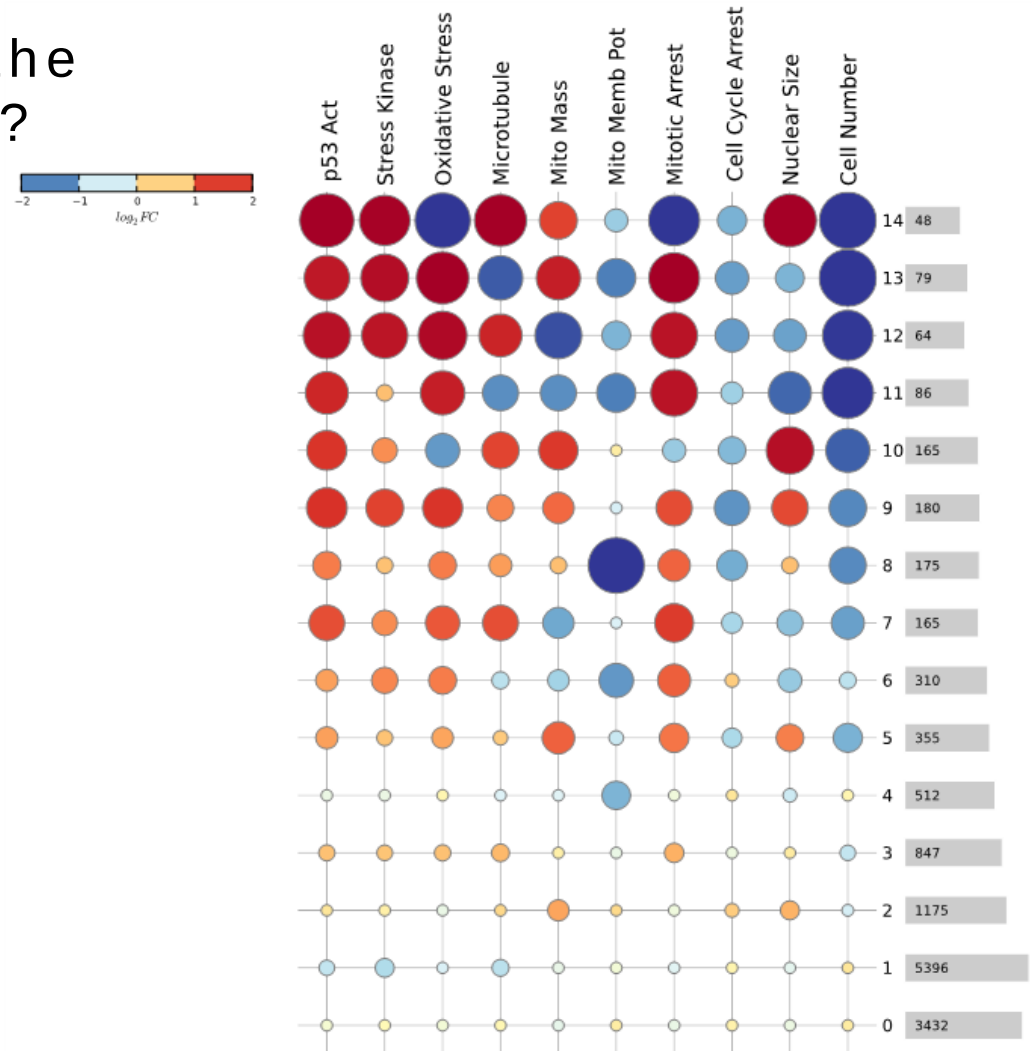


“Continuous Responses”



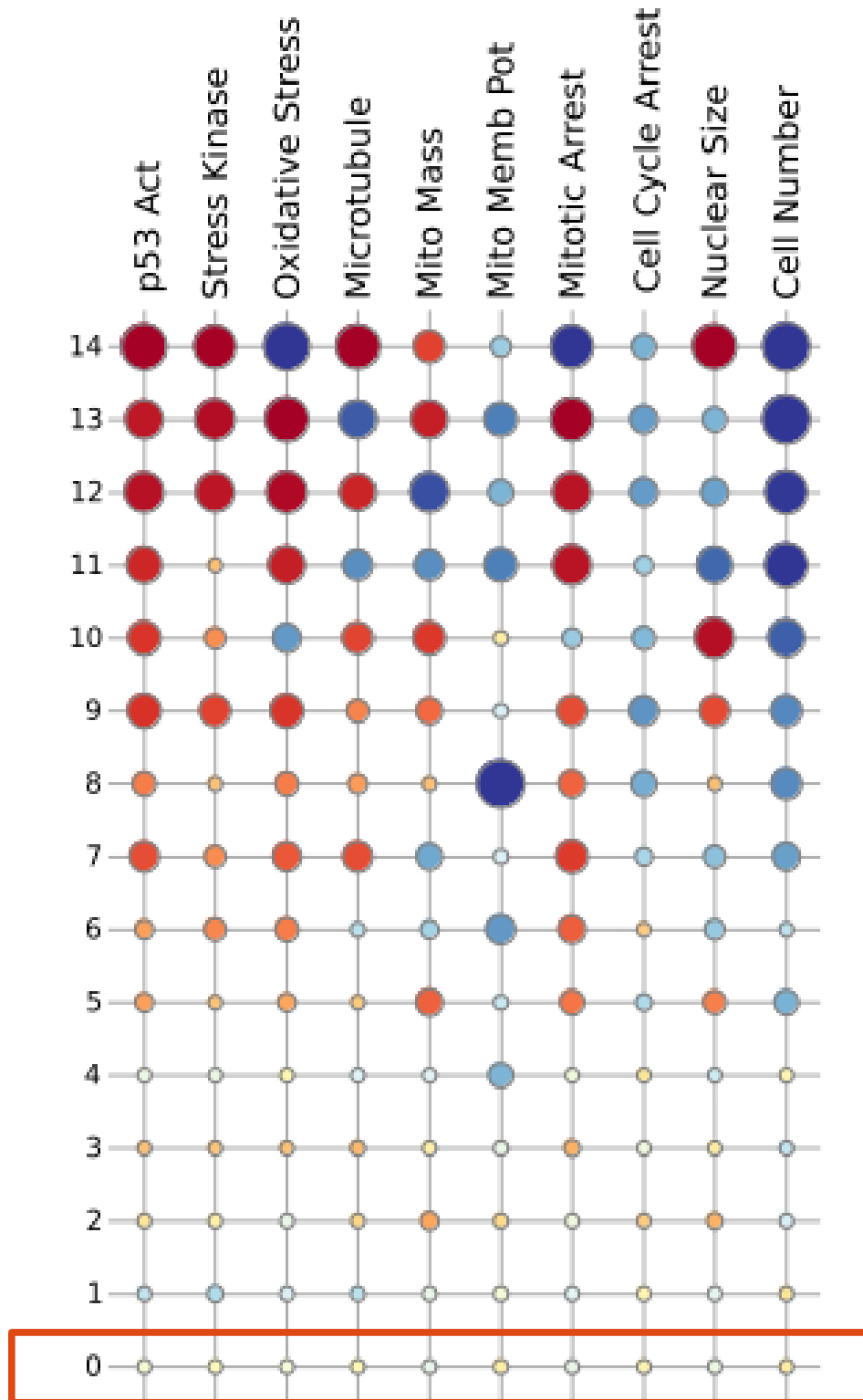
Discrete States – Phenotypes?

- How can we understand the behavior of HepG2 cells ?
- Cluster all data to derive phenotypic “states” of HepG2 system
- Analyze system trajectories
- Can we define normal adaptive and adverse responses ?



“Normal” State

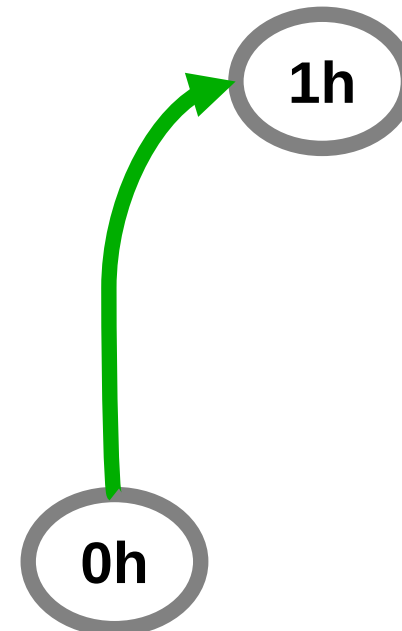
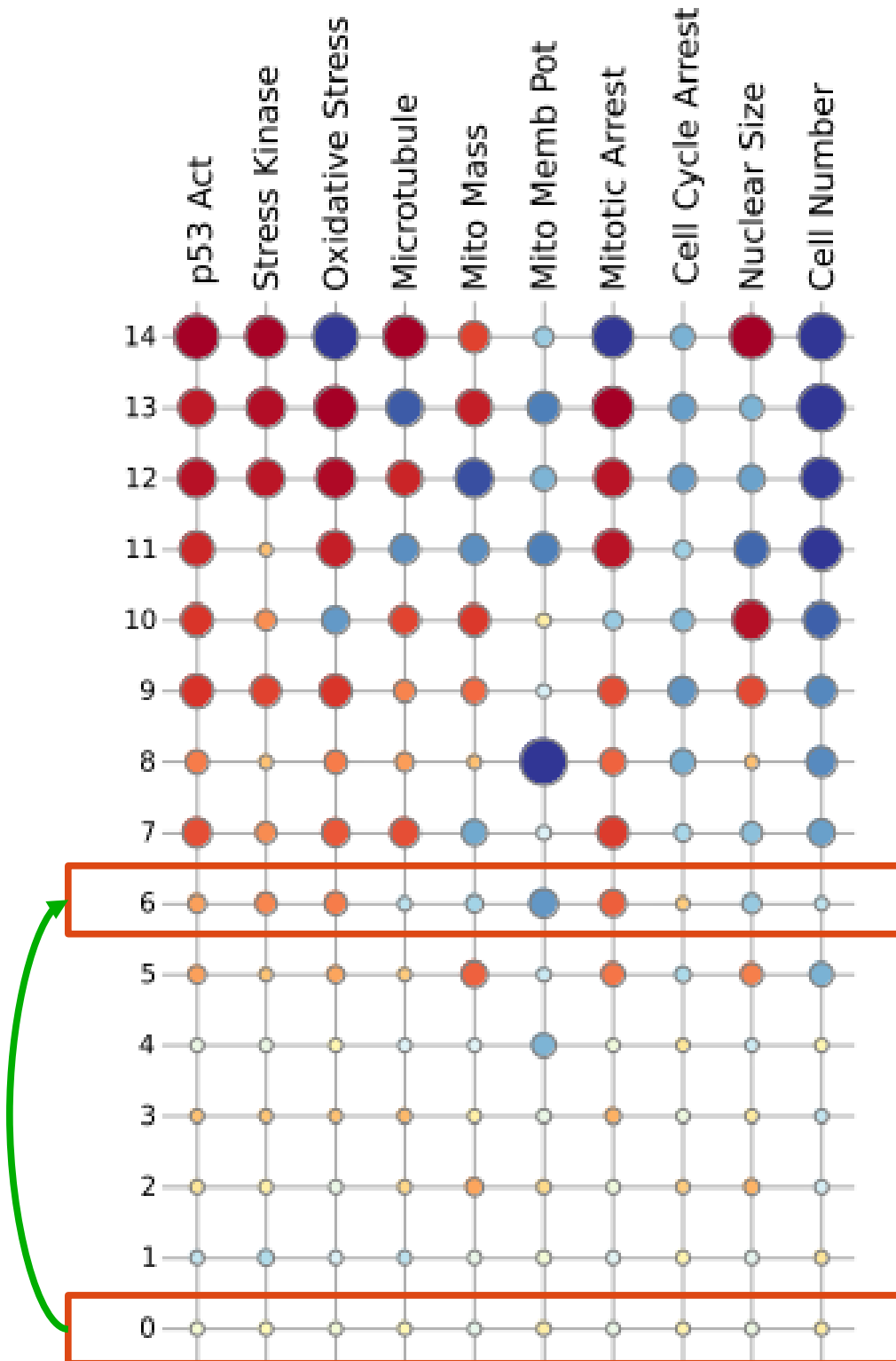
Fluazinam 0.78 μM



0h

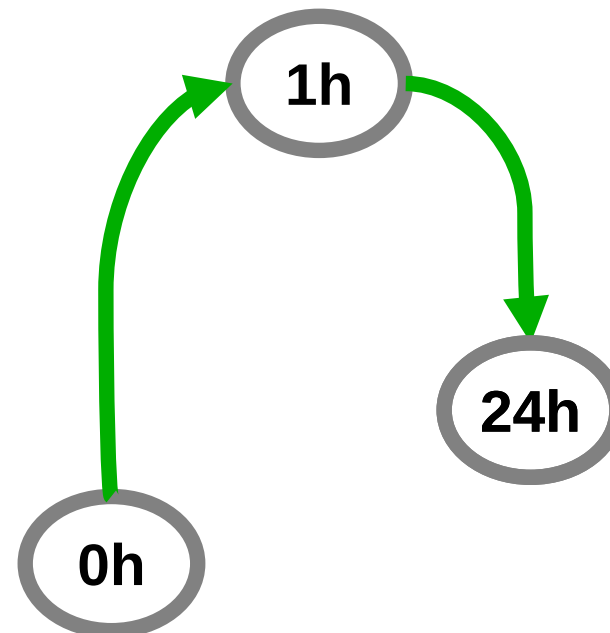
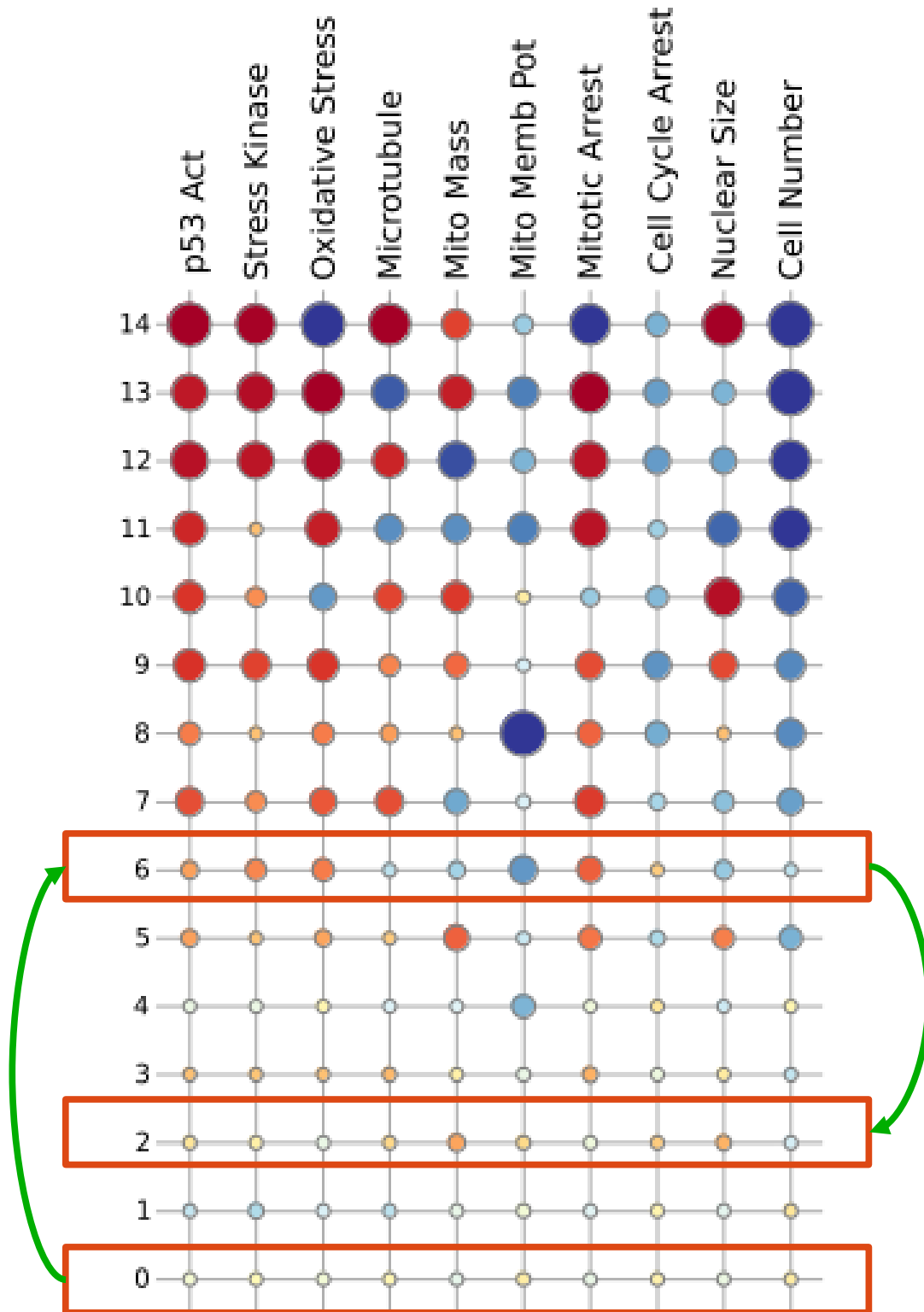
State Transition

Fluazinam 0.78 μM



System trajectory

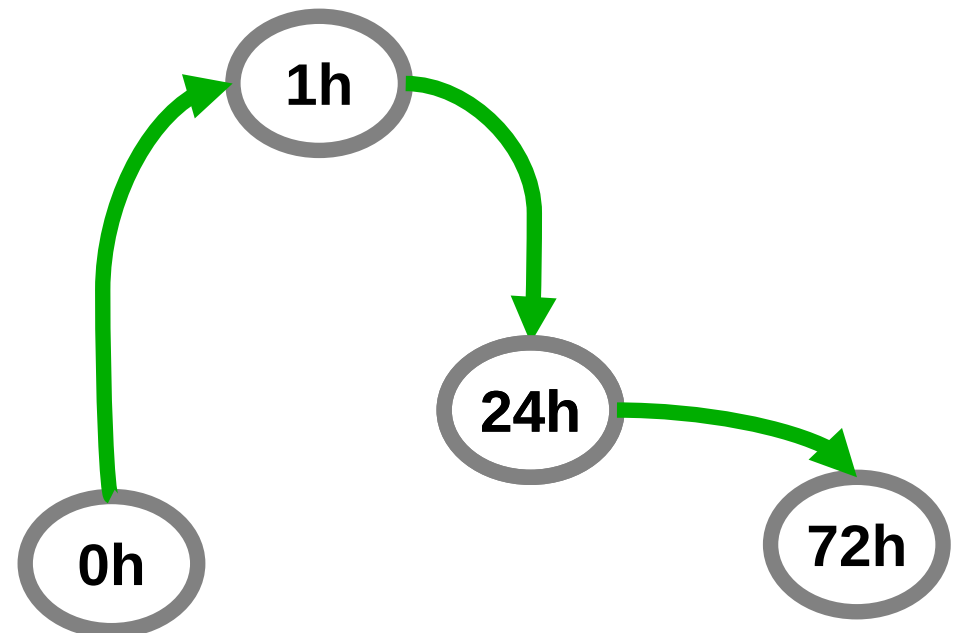
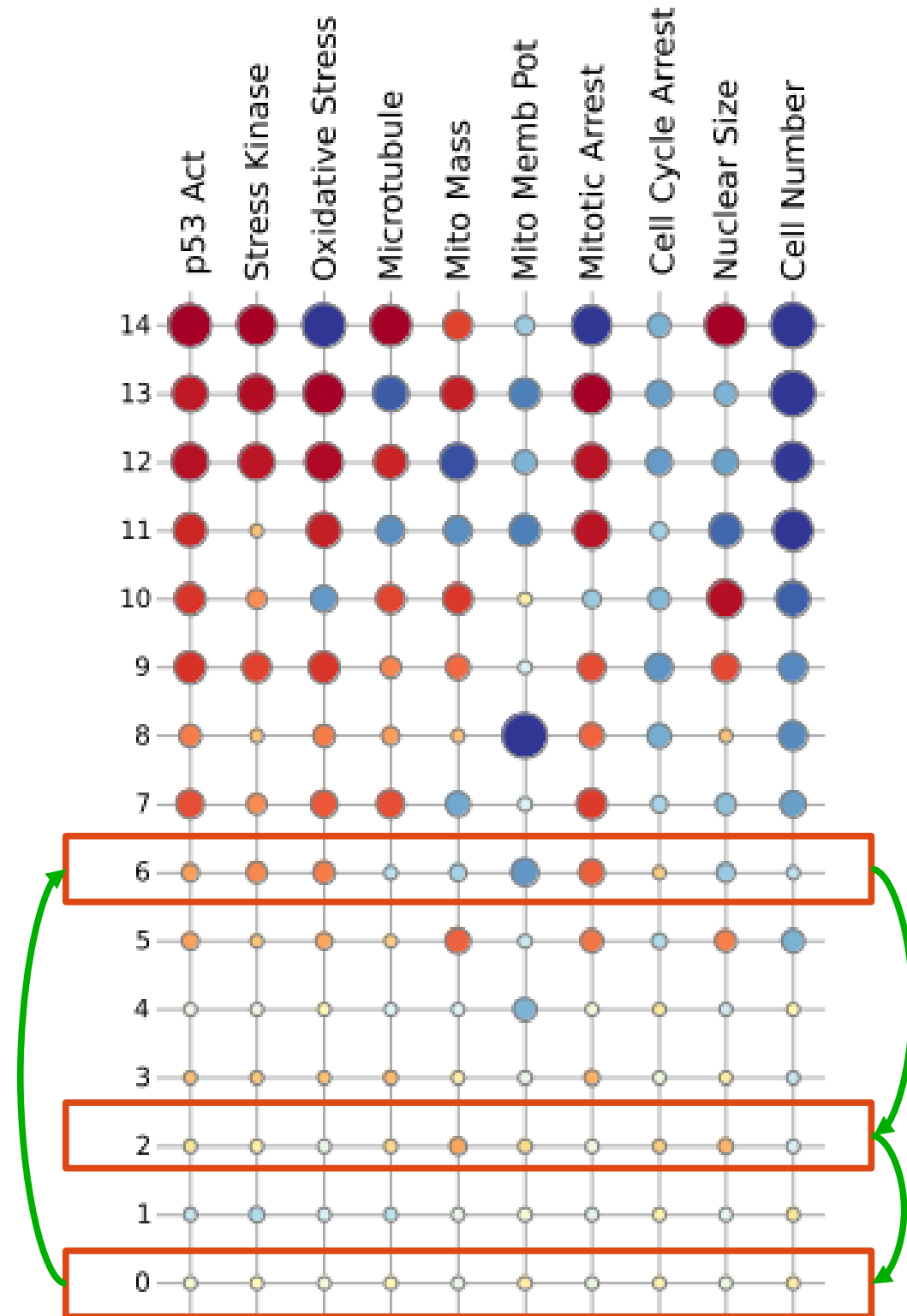
Fluazinam 0.78 μM



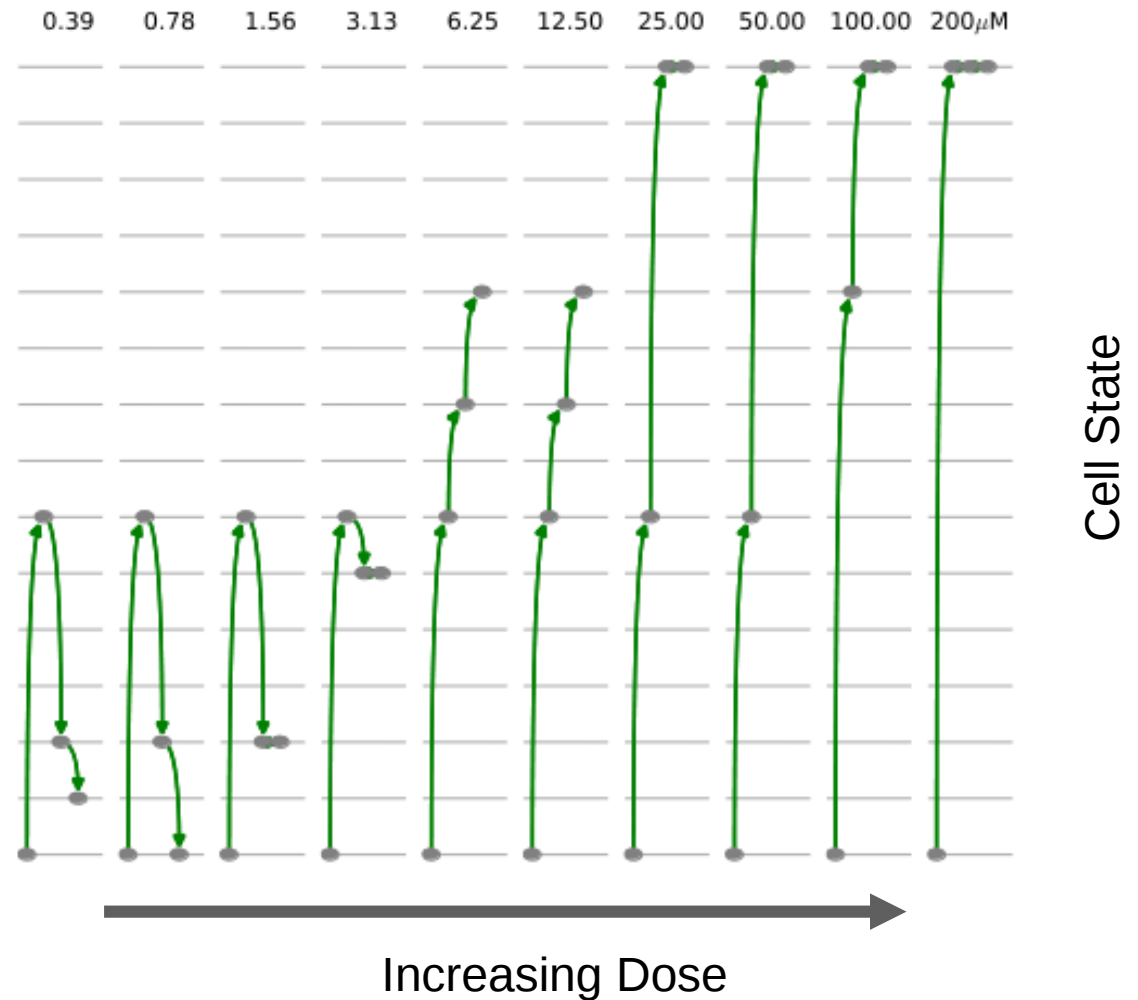
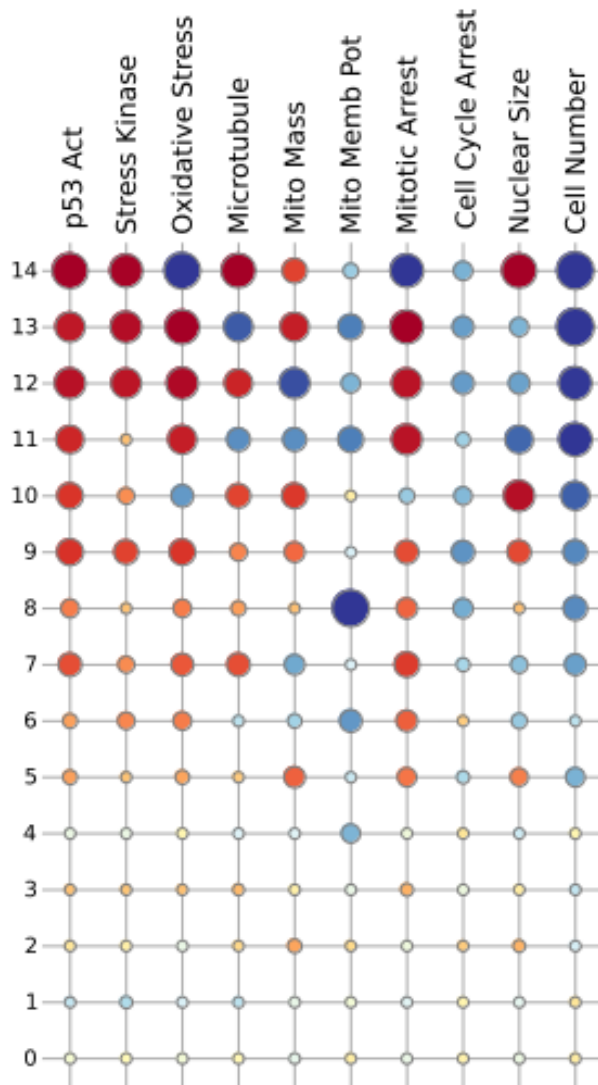
System trajectory

Fluazinam 0.78 μ M

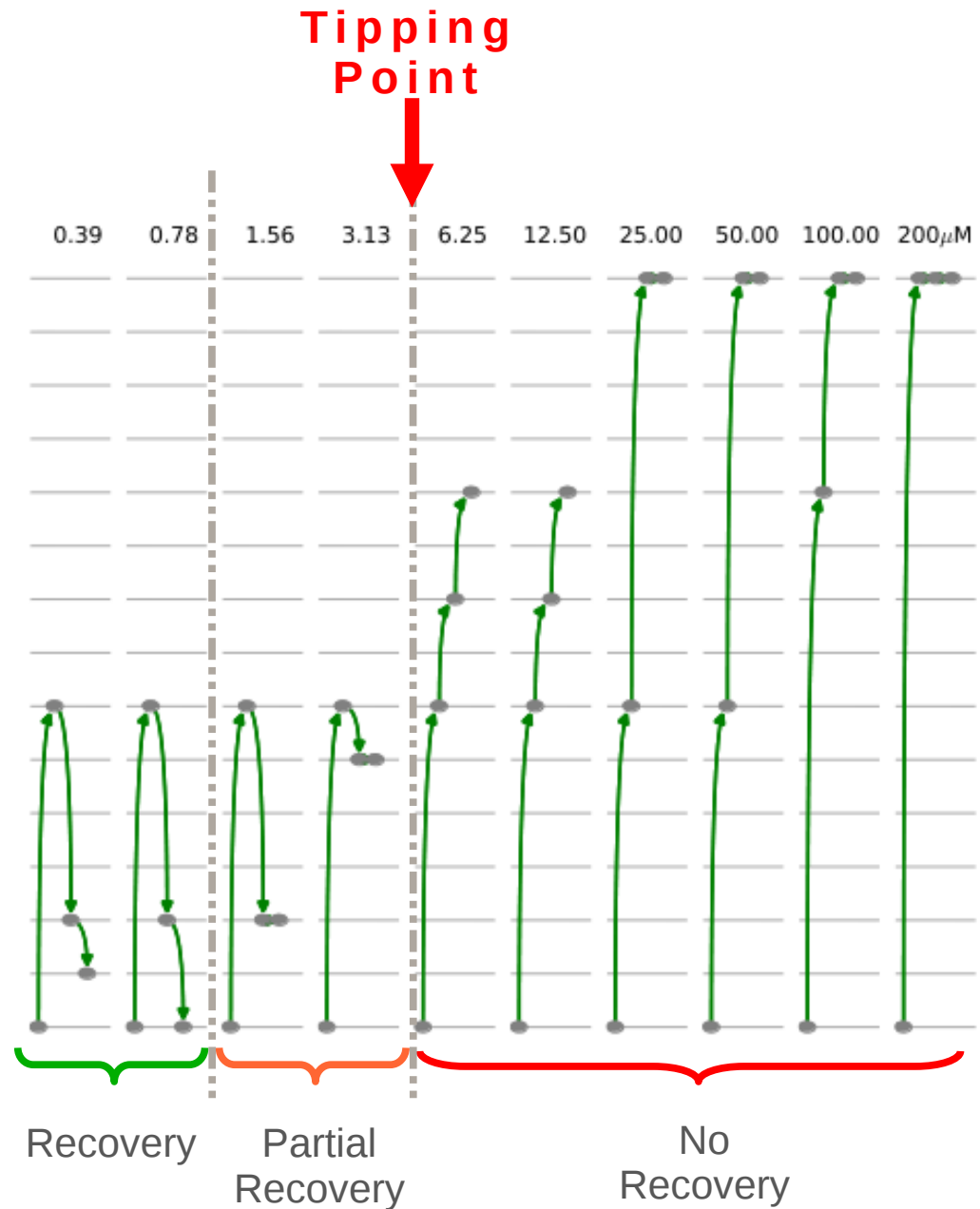
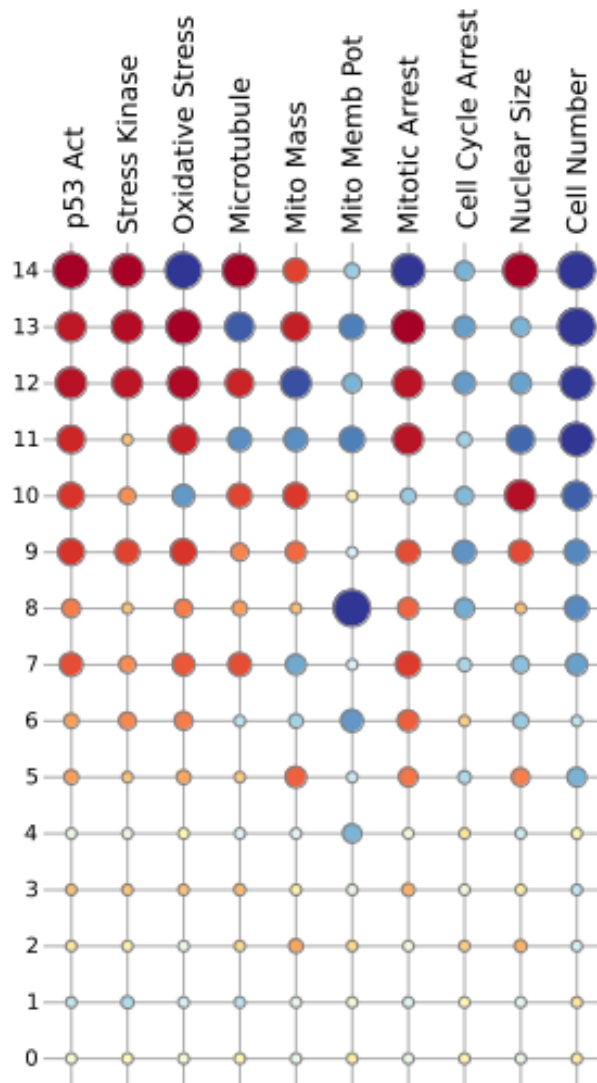
Trajectory = Sequence of states



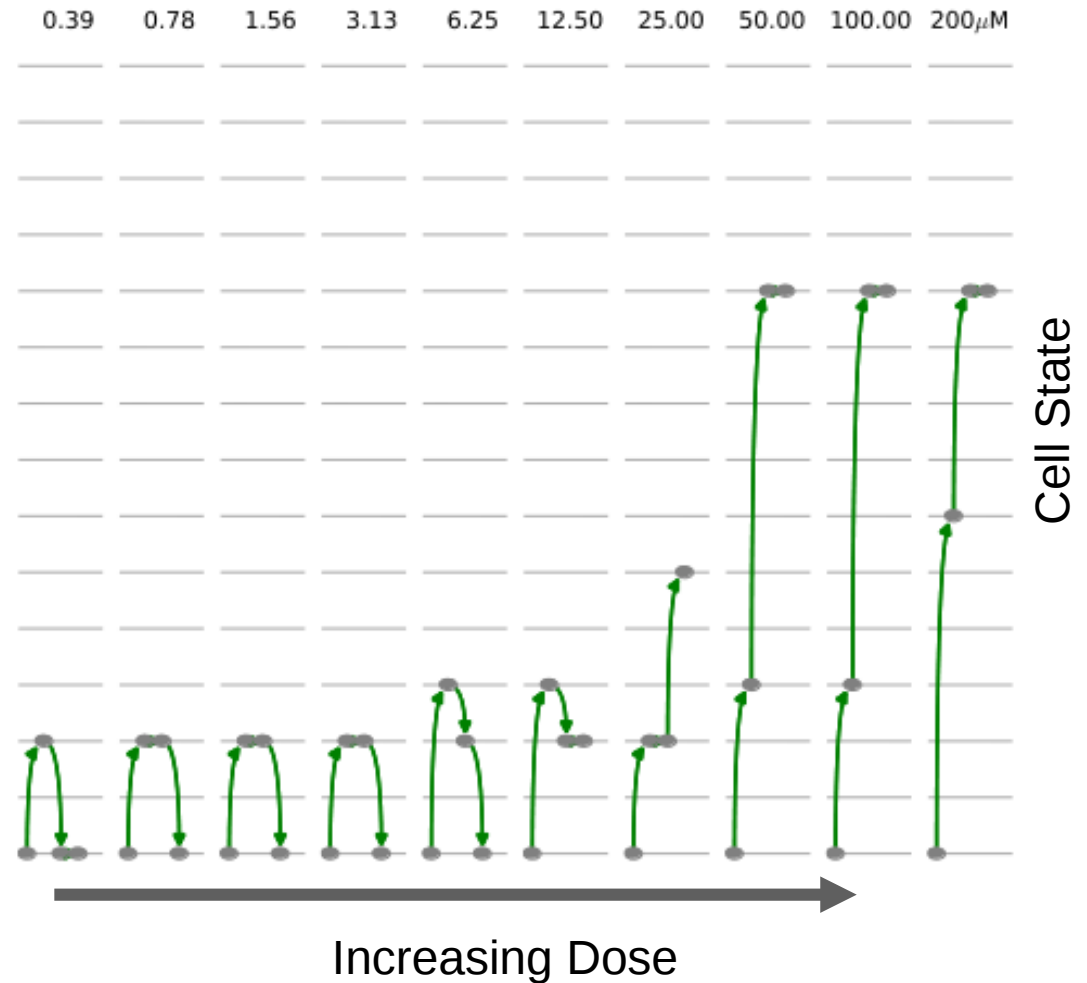
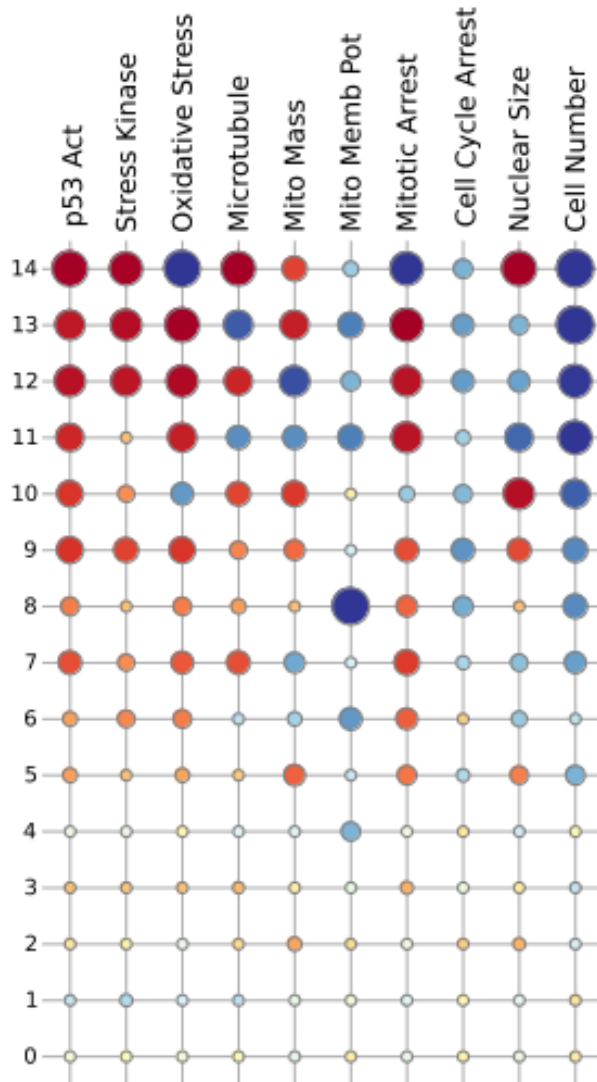
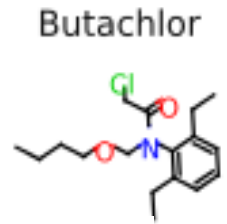
Fluazinam “Trajectories”



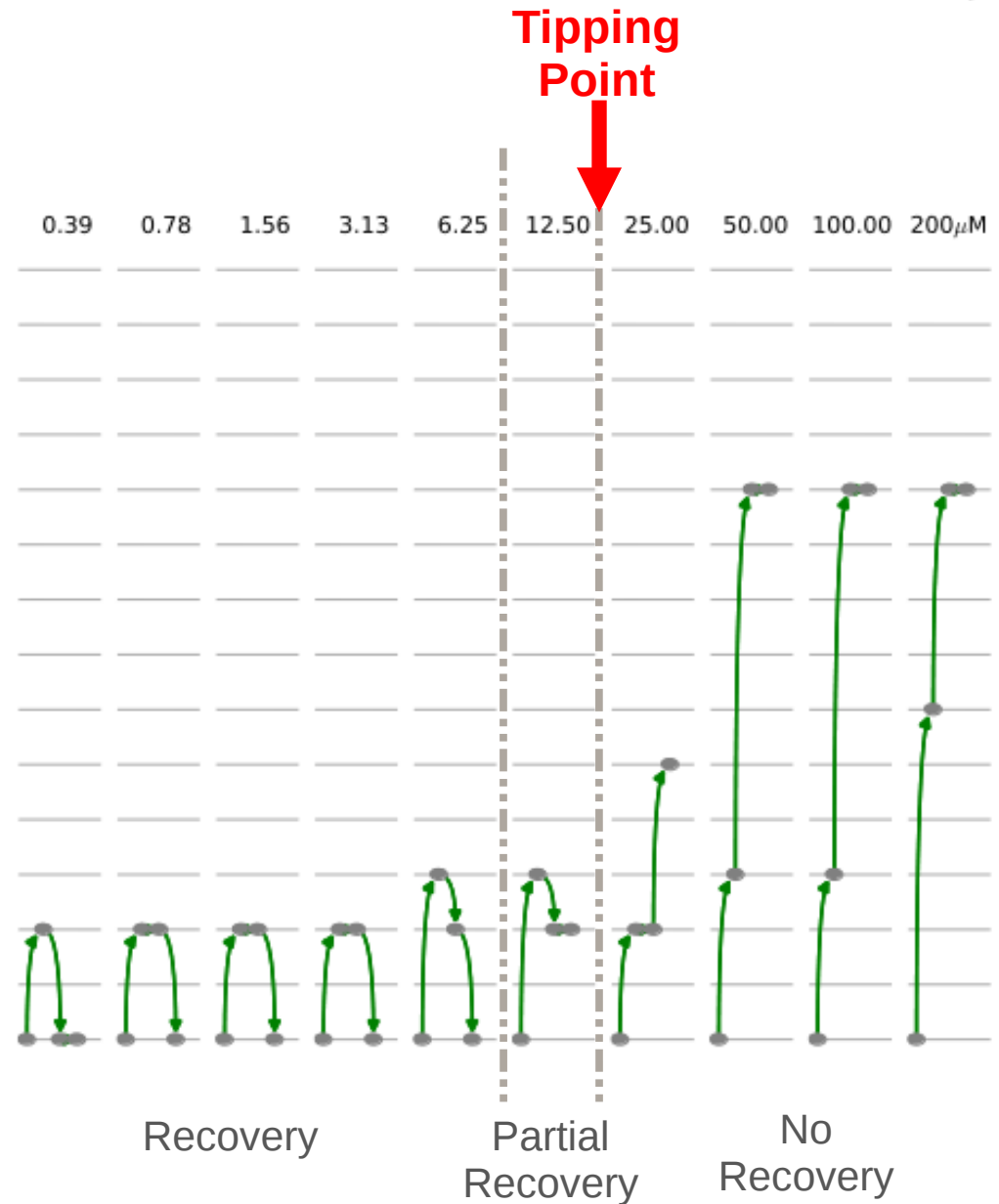
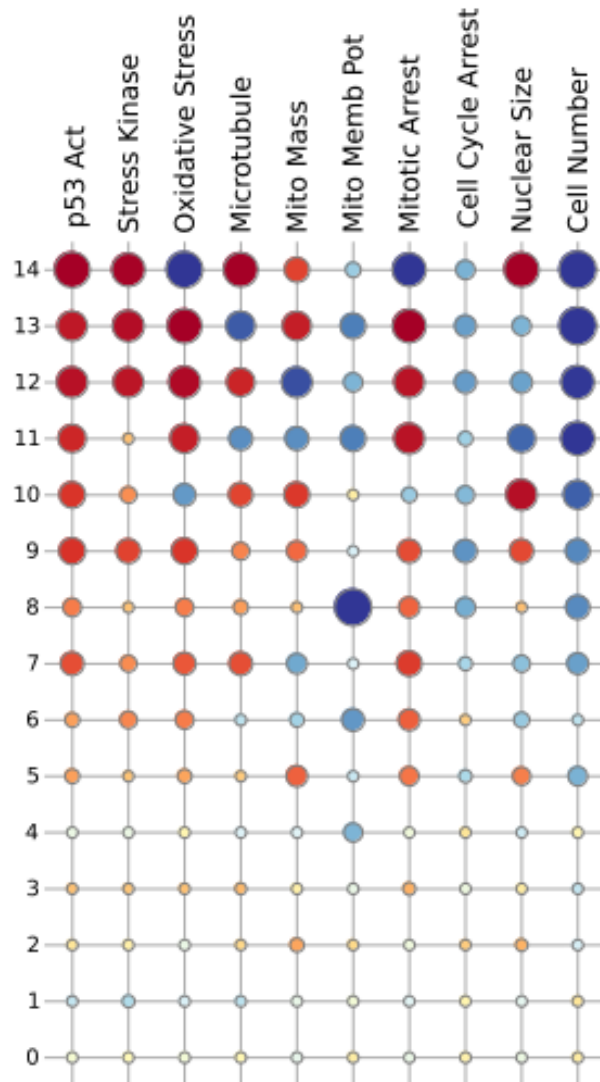
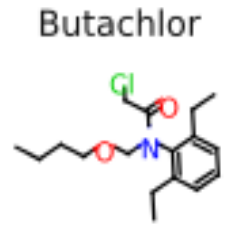
Fluazinam “Trajectories”



Butachlor Trajectories



Butachlor Trajectories



A Systems View...

- ❑ HCI data → System State (X)
- ❑ Chemical → System perturbation
- ❑ Perturbation → Dynamic response
- ❑ **Dynamics → Trajectory**
- ❑ **Trajectory → Adaptation/Adversity**

Biologic endpoint,

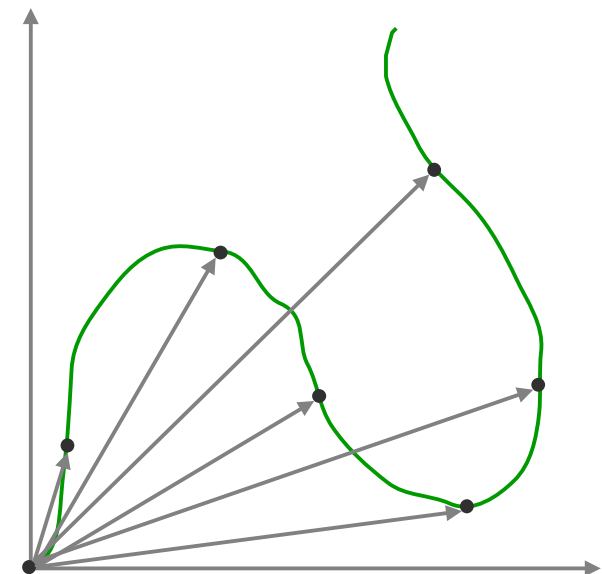
Biologic alteration,

System state, $X^k = [x_1, x_2, x_3, x_4, \dots, x_N]$

Scalar perturbation, $X^k = |S^k|$

Trajectory, scalar

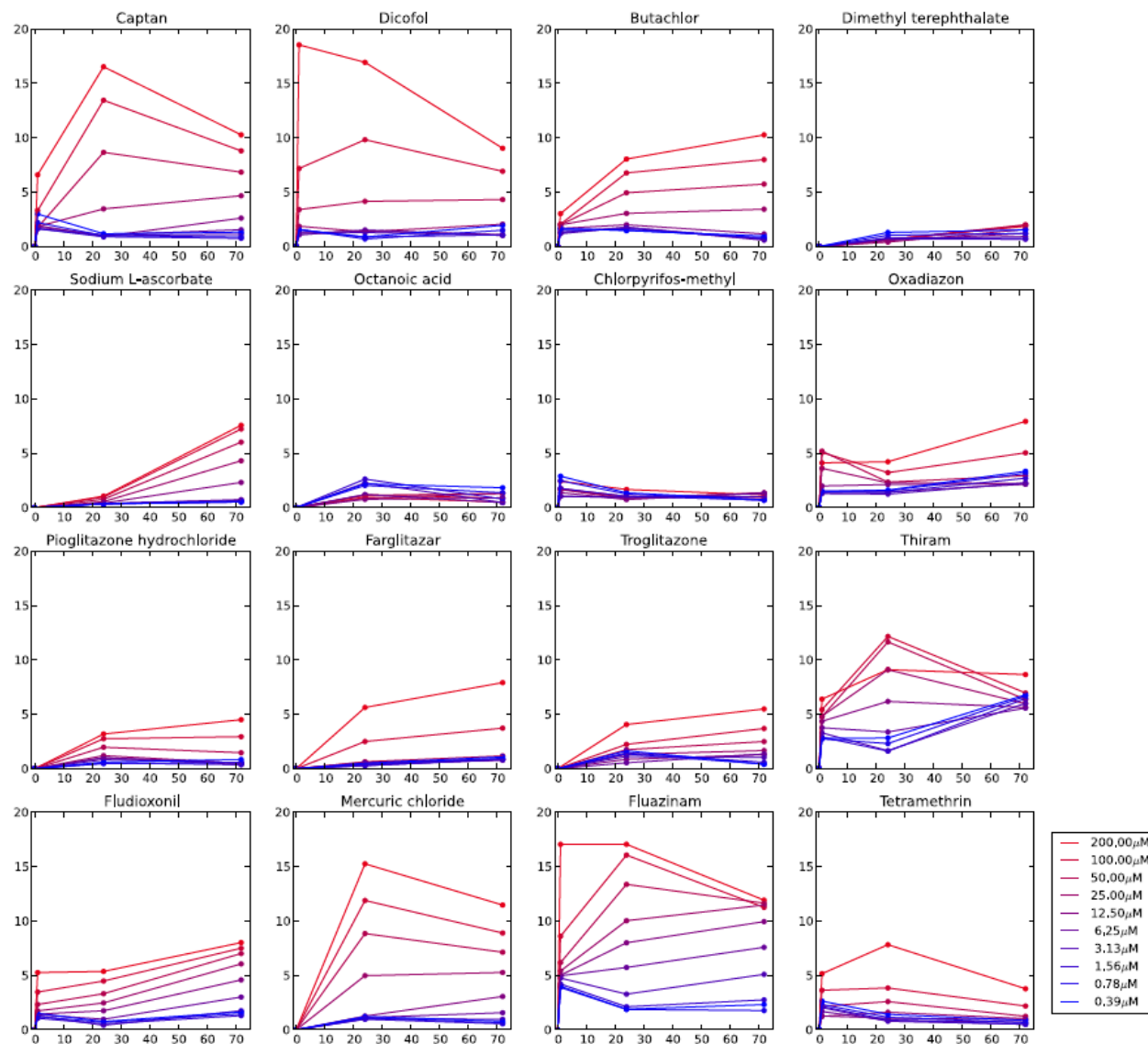
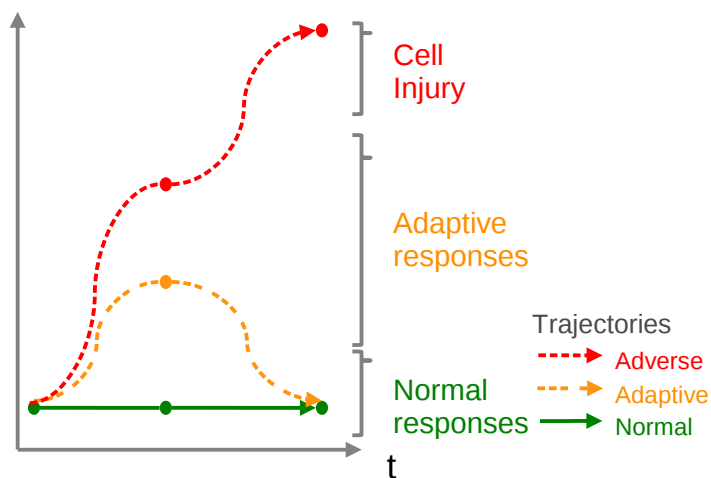
System Trajectory



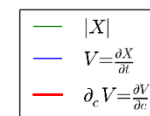
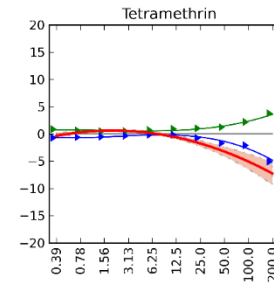
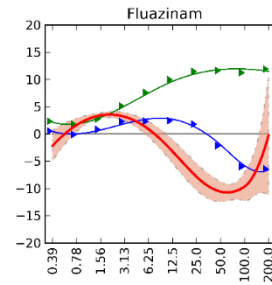
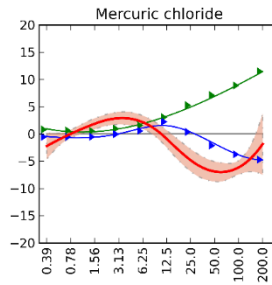
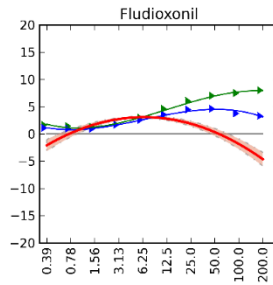
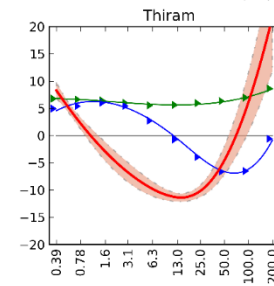
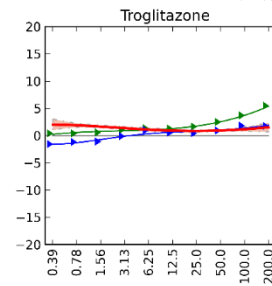
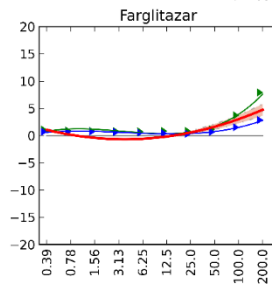
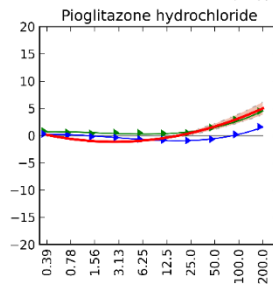
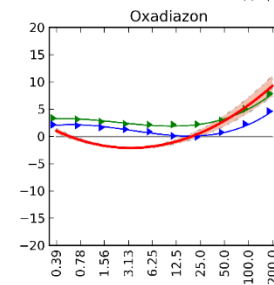
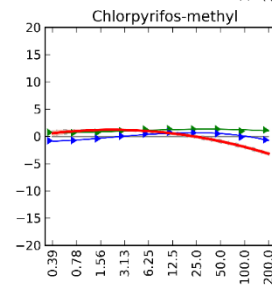
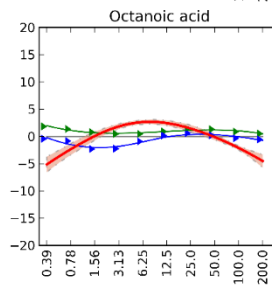
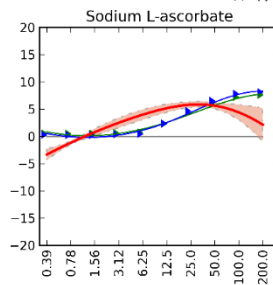
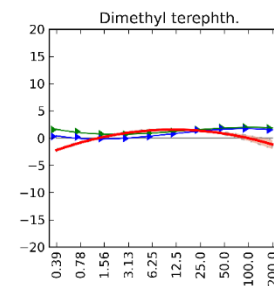
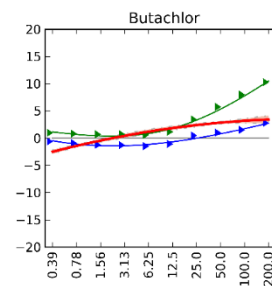
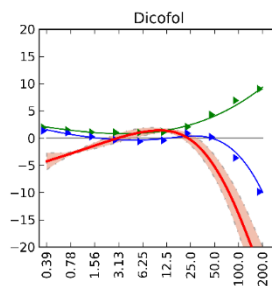
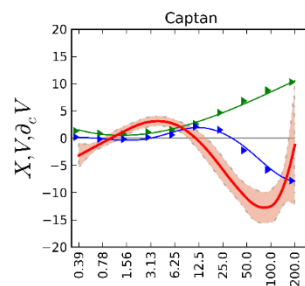
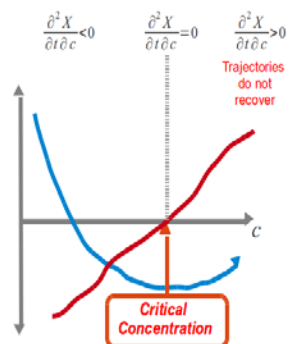
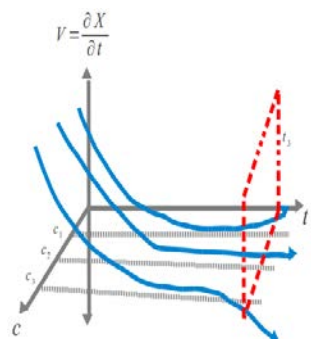
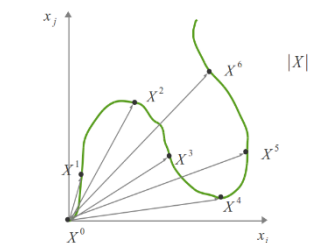
A 2-dimensional projection of trajectory along dimensions (x_i, x_j)

Shah et al 2015

Trajectories & Adaptation

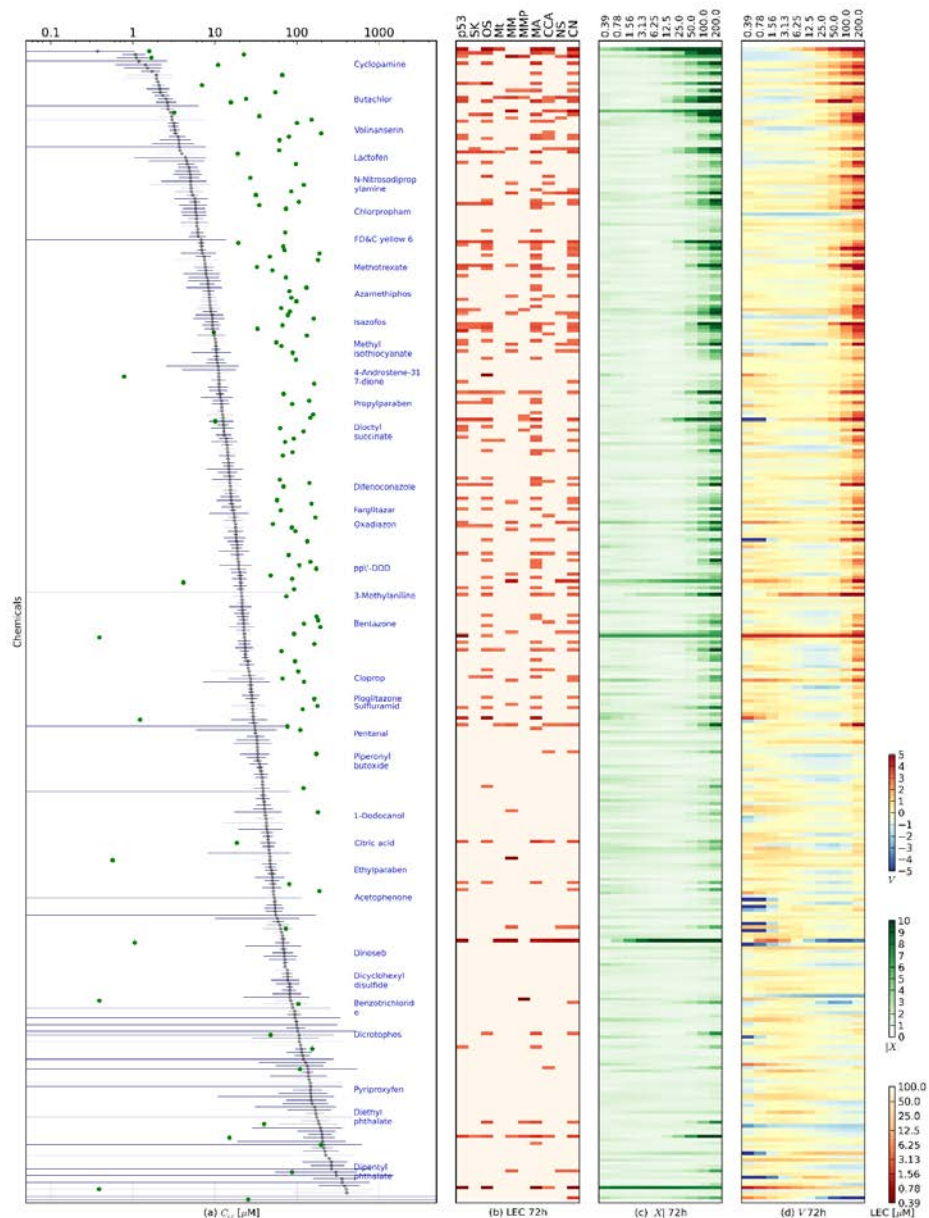


Trajectories & Tipping Points



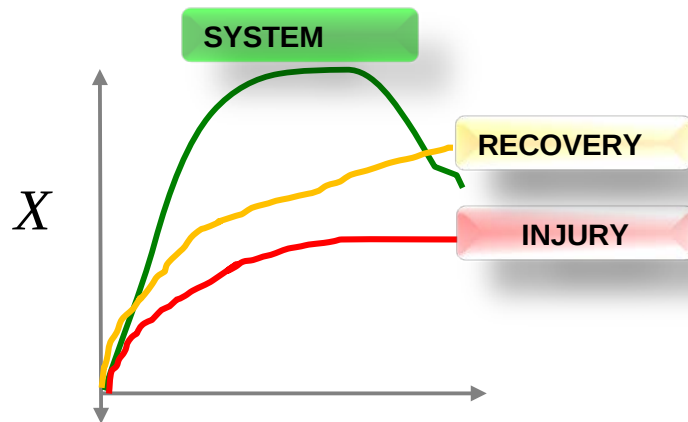
Trajectories & Tipping Points

- Can systematically identify critical concentrations-
“tipping points”
- Tipping points precede
concentration that
produces significant
cell loss

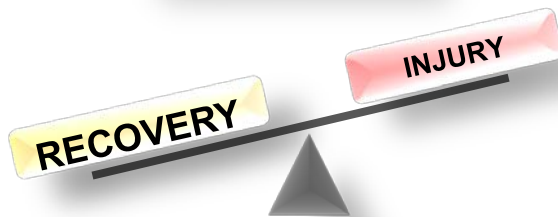


Toxicological Tipping Points

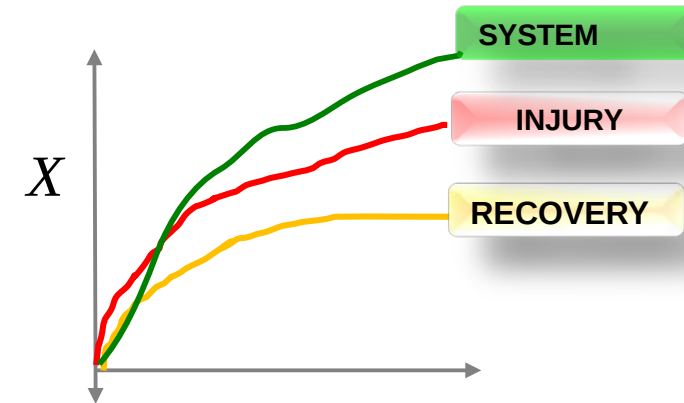
ADAPTATION



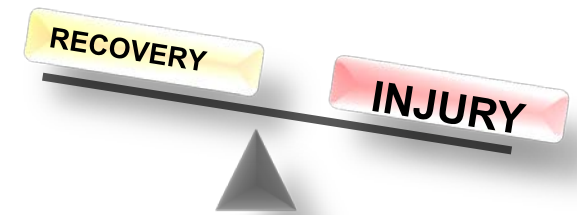
**SYSTEM
ADAPTATION**



ADVERSITY



**SYSTEM
ADVERSITY**



**SYSTEM
TIPPING POINT**



CHALLENGES...

- ❑ Study design
 - More time points !
 - More stress-response markers !
 - How to define critical window to identify tipping points ?
 - More physiologically-relevant cell model
- ❑ Tipping point evaluation
 - Is it just PK or is there real adaptation ?
 - Are they reproducible ?
 - Are they biologically relevant (ie. mechanisms)?
- ❑ Translation
 - How do *in vitro* tipping points compare *in vivo* PoD ?