

Finding Toxicological Tipping Points from High- Content Imaging Data

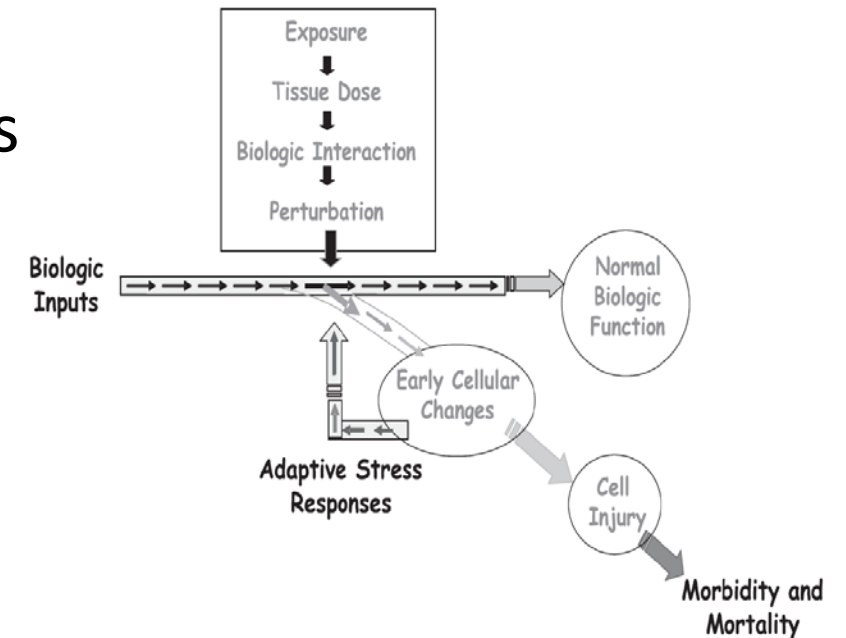
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Toxicological Tipping Points

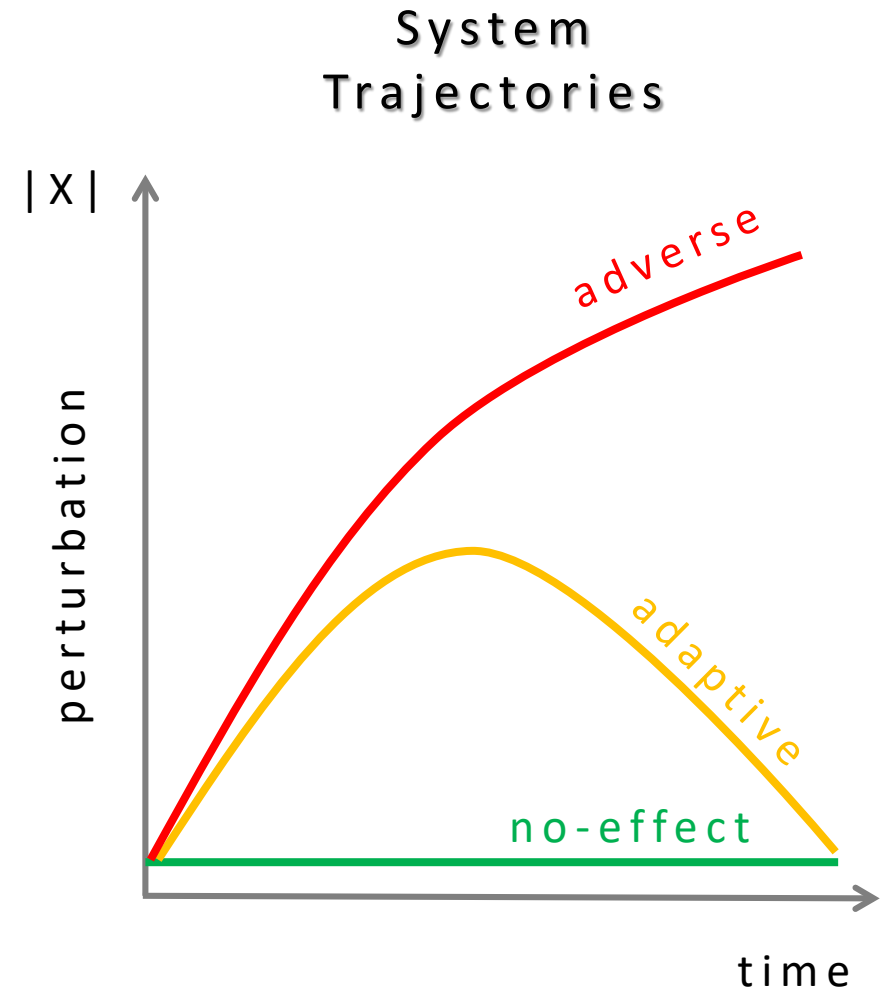
- ❑ Biological systems are resilient and adapt to environmental perturbations
- ❑ Can we find toxicological tipping points?
- ❑ Can a tipping point define a point of departure (PoD) ?



Krewski, Daniel, Daniel Acosta Jr, Melvin Andersen, Henry Anderson, John C Bailar 3rd, Kim Boekelheide, Robert Brent, et al. "Toxicity Testing in the 21st Century: a Vision and a Strategy." *Journal of Toxicology and Environmental Health. Part B, Critical Reviews* 13, no. 2-4 (February 2010): 51-138.

Tipping Points, *in vitro*

- ❑ Used high-content imaging to model system trajectories: no-effect, adaptive and adverse
- ❑ Tipping point: critical point between adaptive and adverse trajectories
- ❑ How can we use network modeling to analyze tipping points?



High Content Imaging (HCI)

□ Study

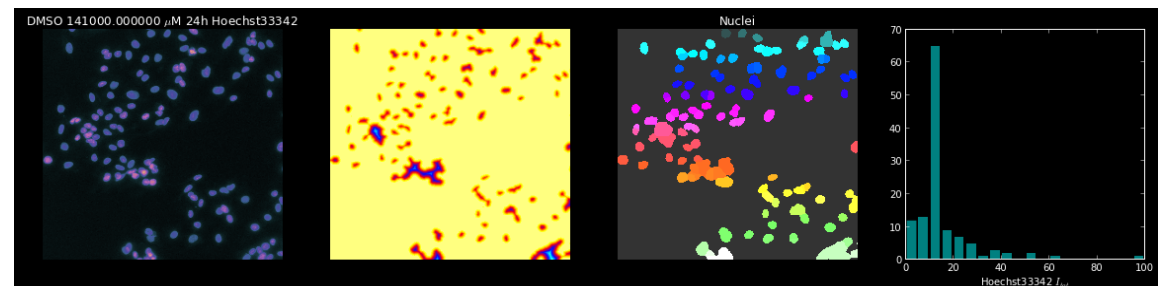
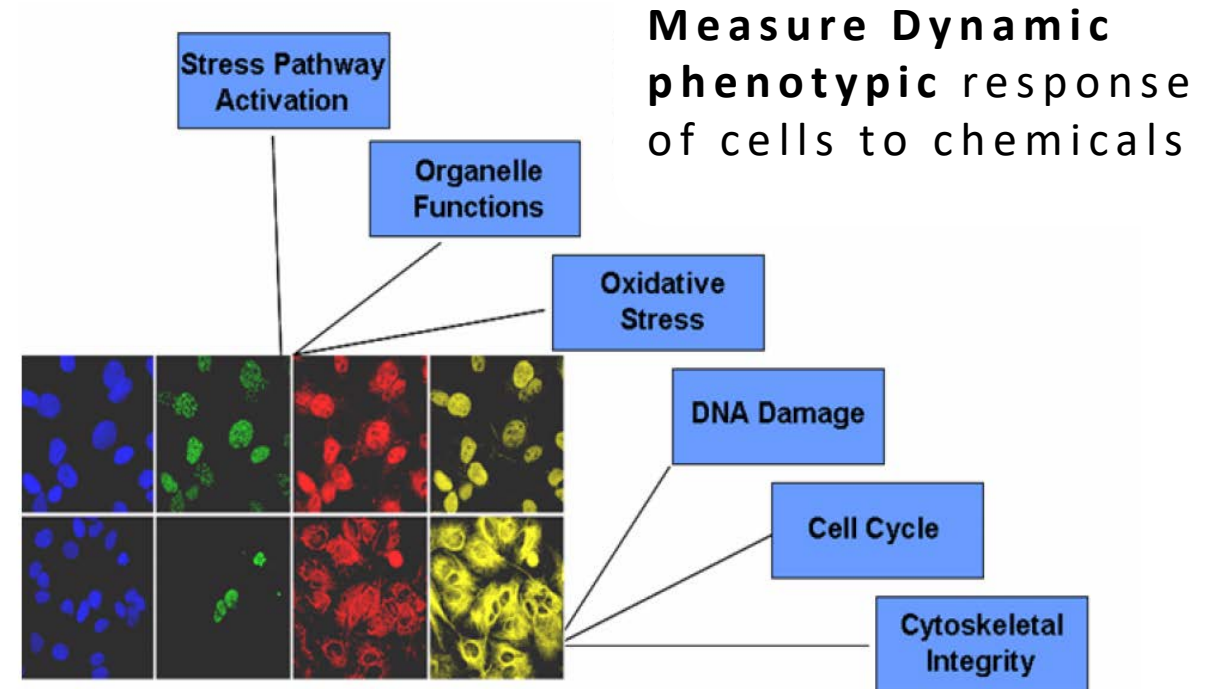
- HepG2 cell culture
- 967 chemicals (ToxCast)
- 10 conc

□ HCI Assays

- Health
- Stress
- Cell cycle

□ Large-scale data

- ~400 plates
- ~100,000 wells
- ~2,400,000 images
- ~30,000 chemical-conc-time-response points

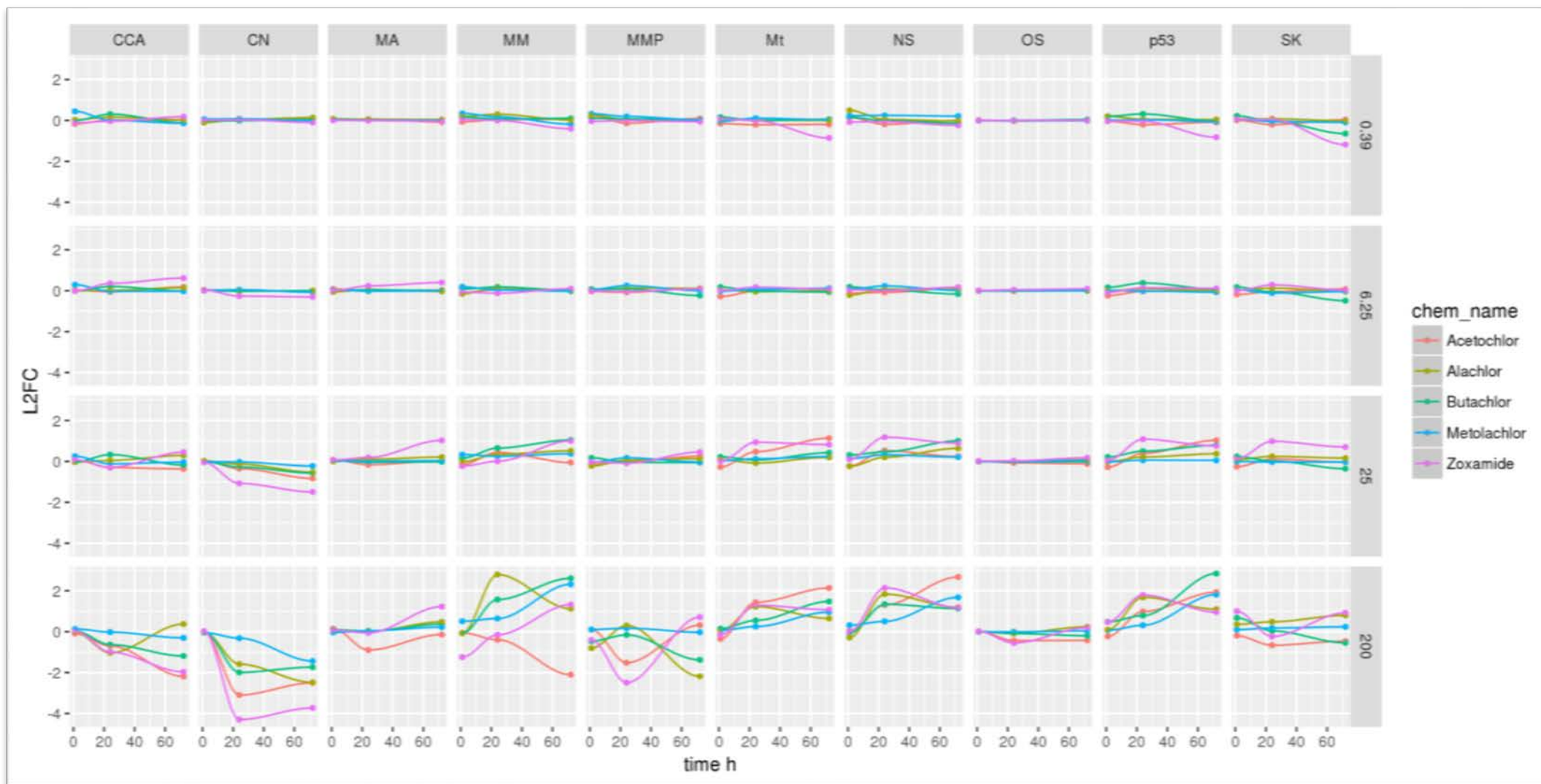


Raw Image
(Hoechst)

Intensity
Analysis

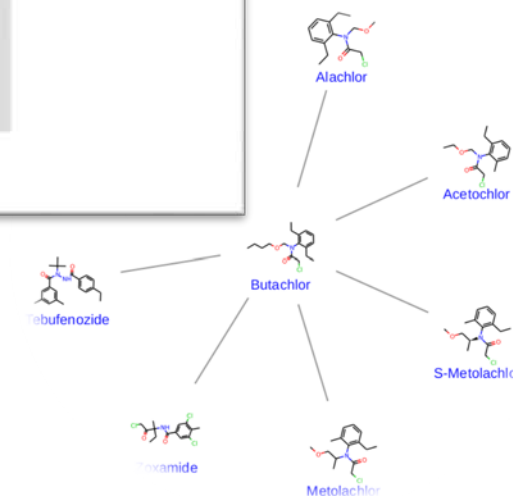
Object
Identification

Nuclear intensity
distribution

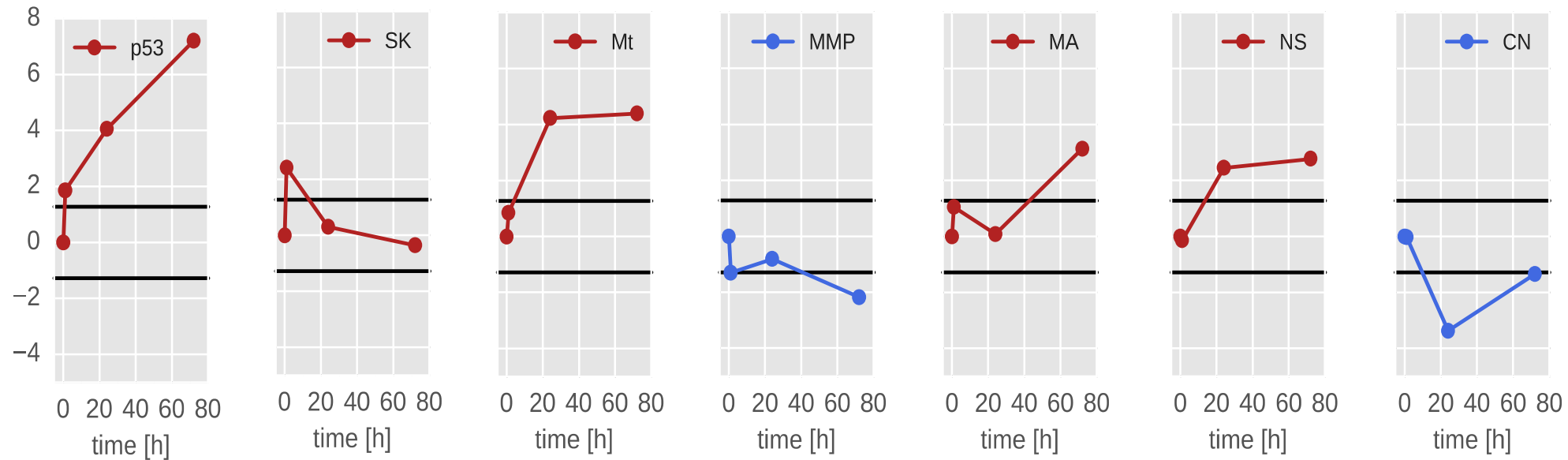


- P53: p53 activity (p53)
- c-Jun: stress kinase (SK)
- Tubulin: microtubule organisation (Mt)
- MitoTracker Red: Mitochondrial memb. pot. (MMP)

- PH3: mitotic arrest (MA)
- Hoechst33342: nuclear size (NS), cell number (CN)
- MitoTrackerRed: Mitochondrial memb. pot. (MMP)



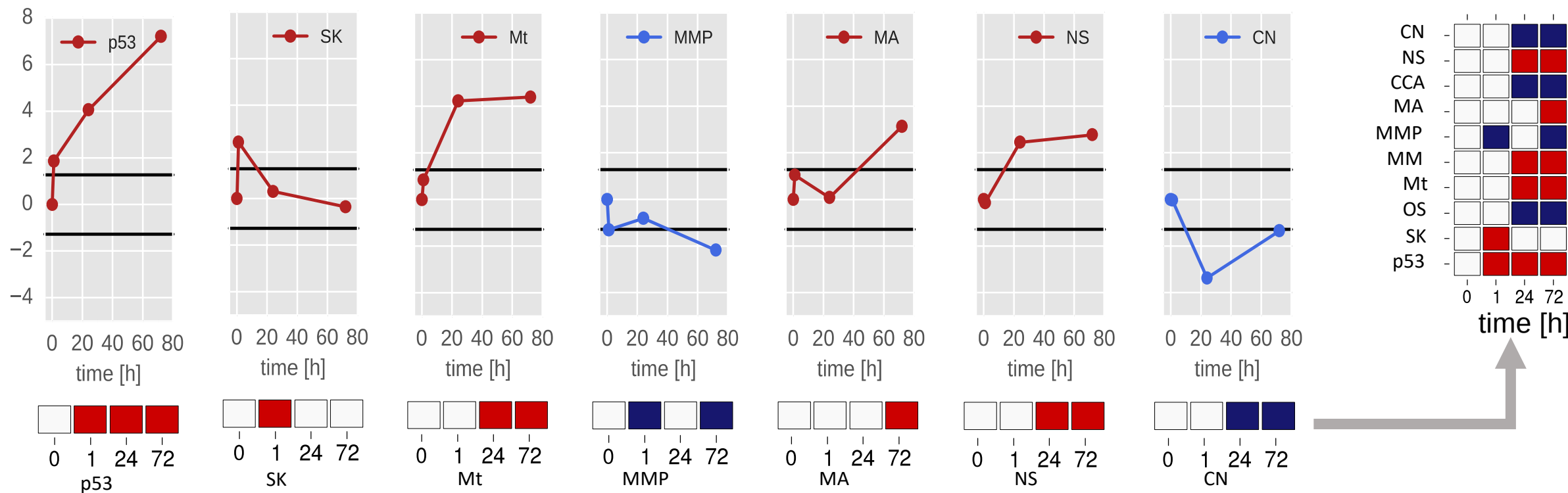
Butachlor 200 μ M Trajectory



- P53: p53 activity (p53)
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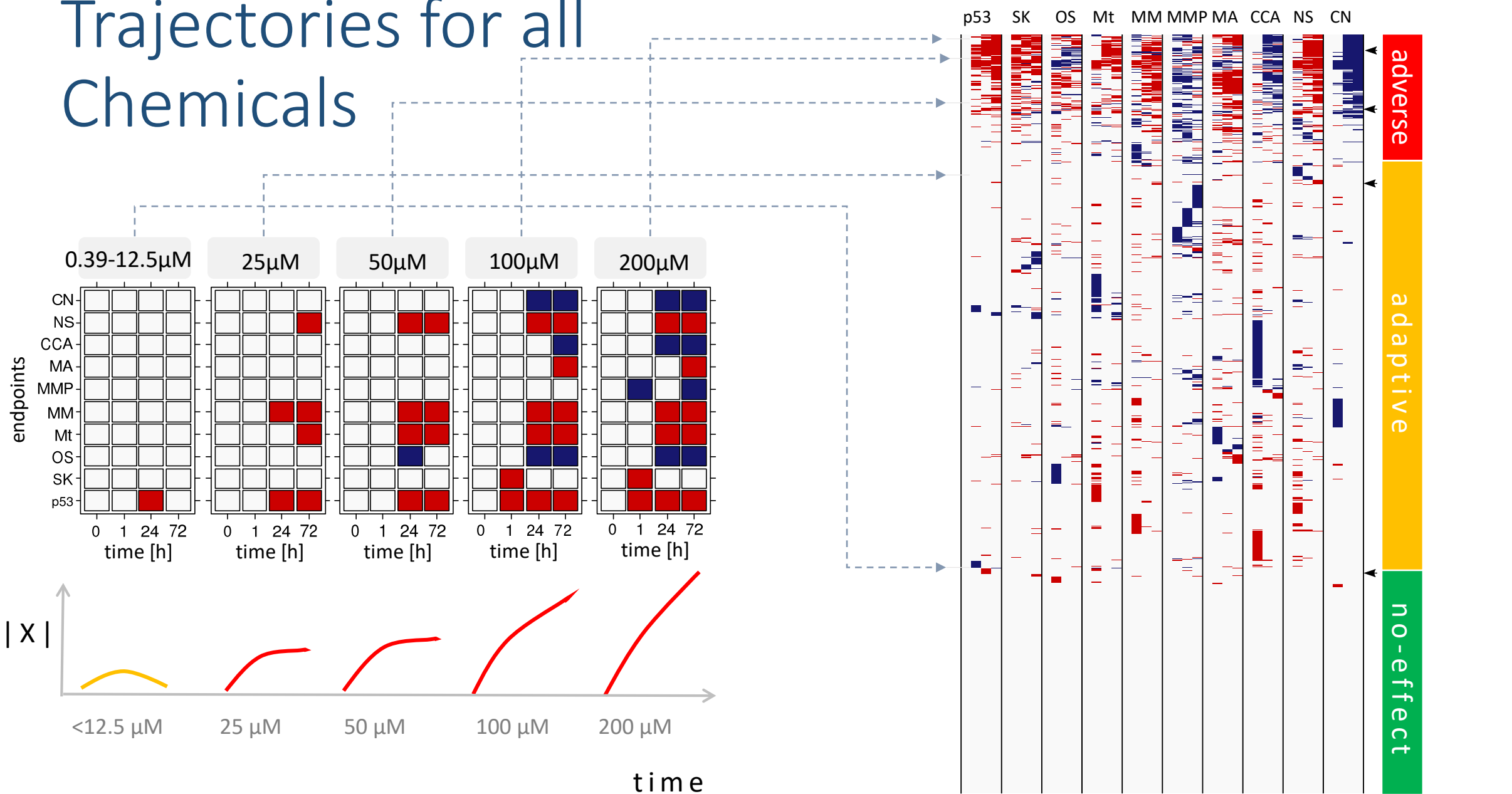
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Discretizing Data



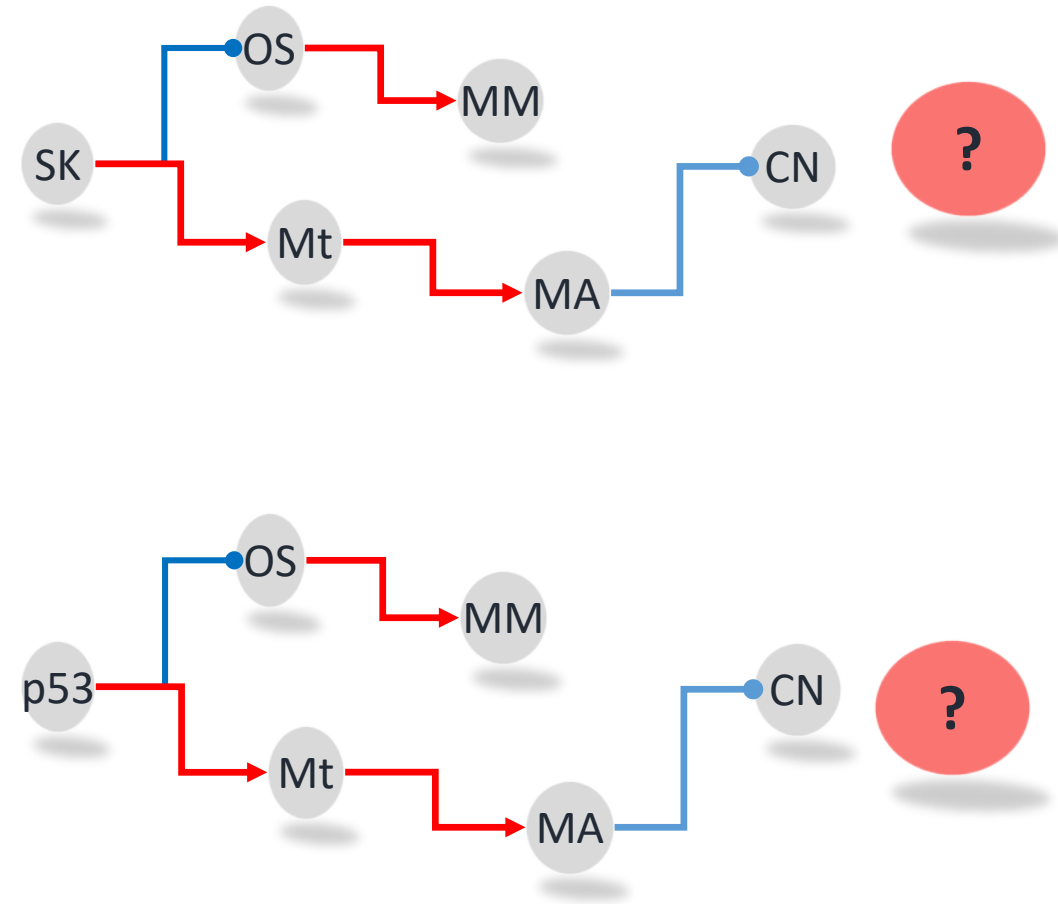
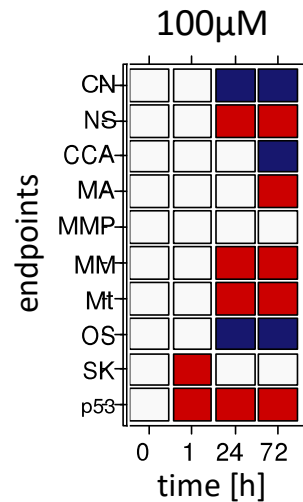
Discretization highlights key responses and reduces noise

Trajectories for all Chemicals

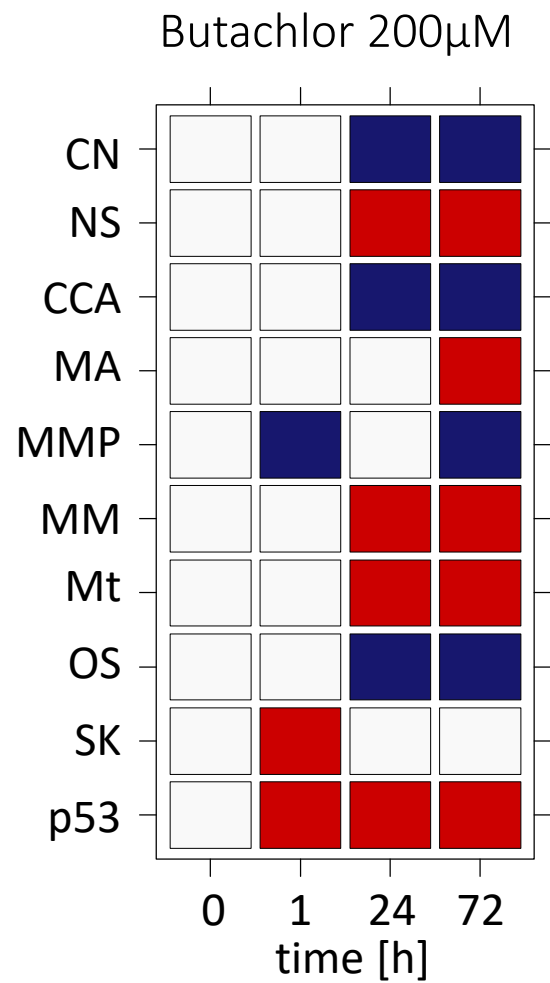


Learning Mechanisms from Data

- Analyse dependencies between endpoints across time
- Tedious to do by hand
- Many network inference approaches
- We used Boolean network (BN) inference using best-fit extension

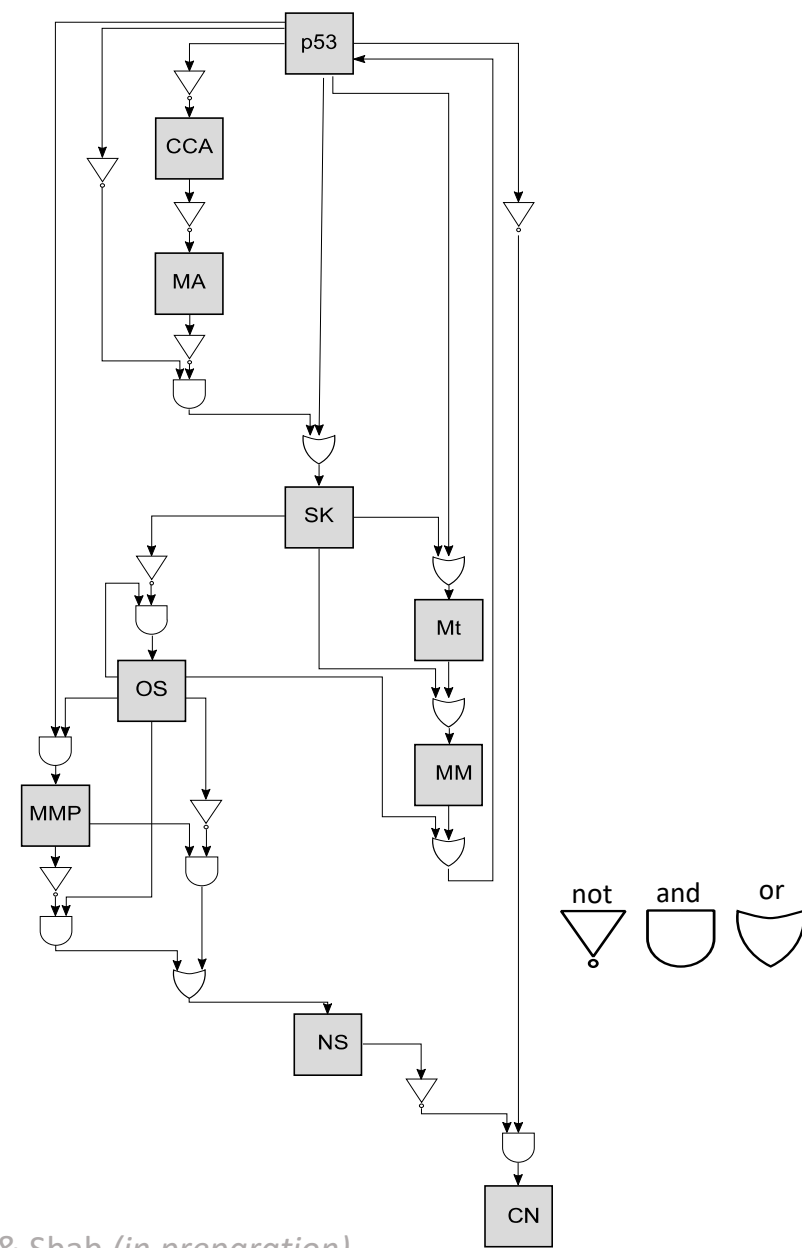


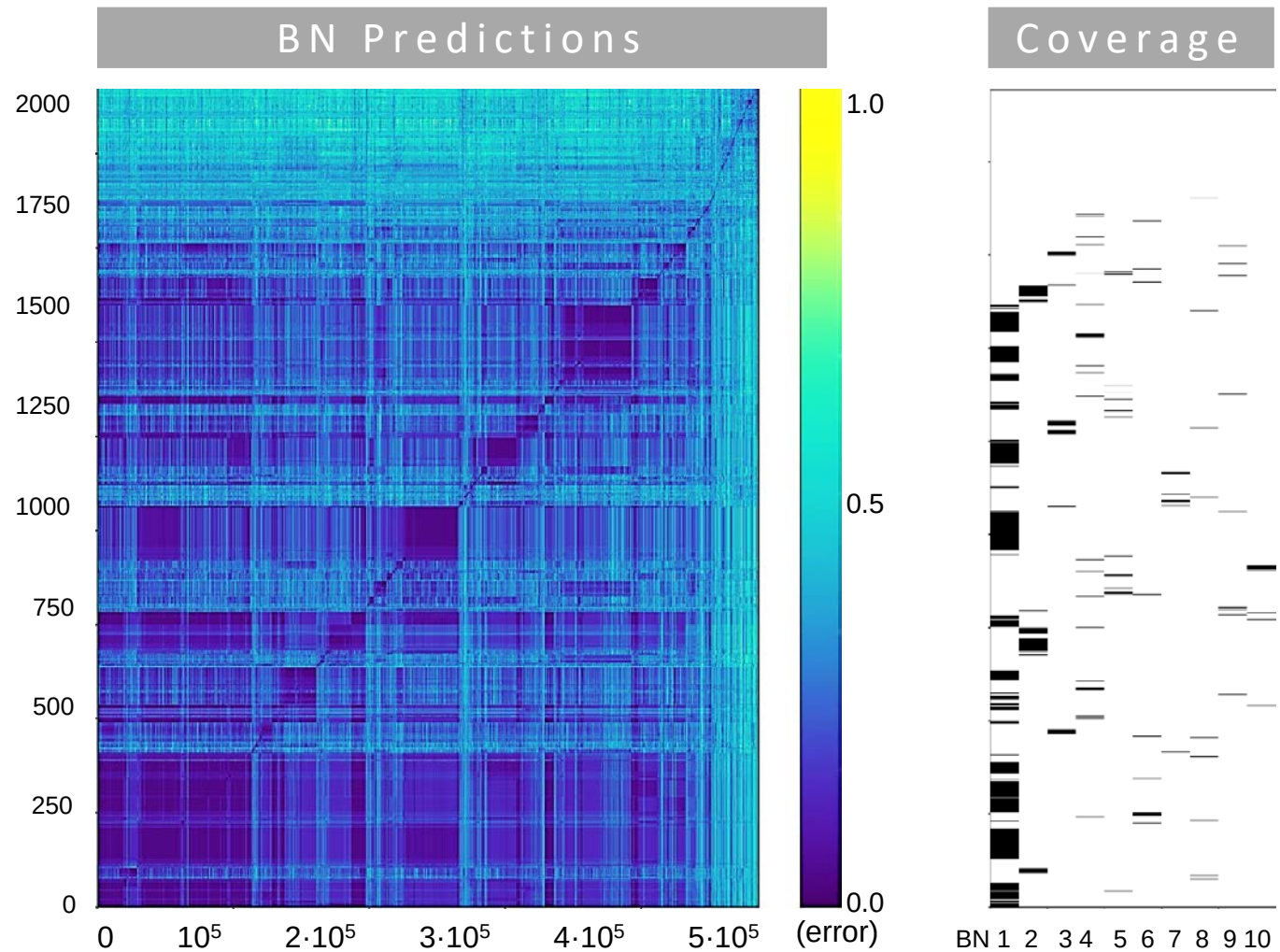
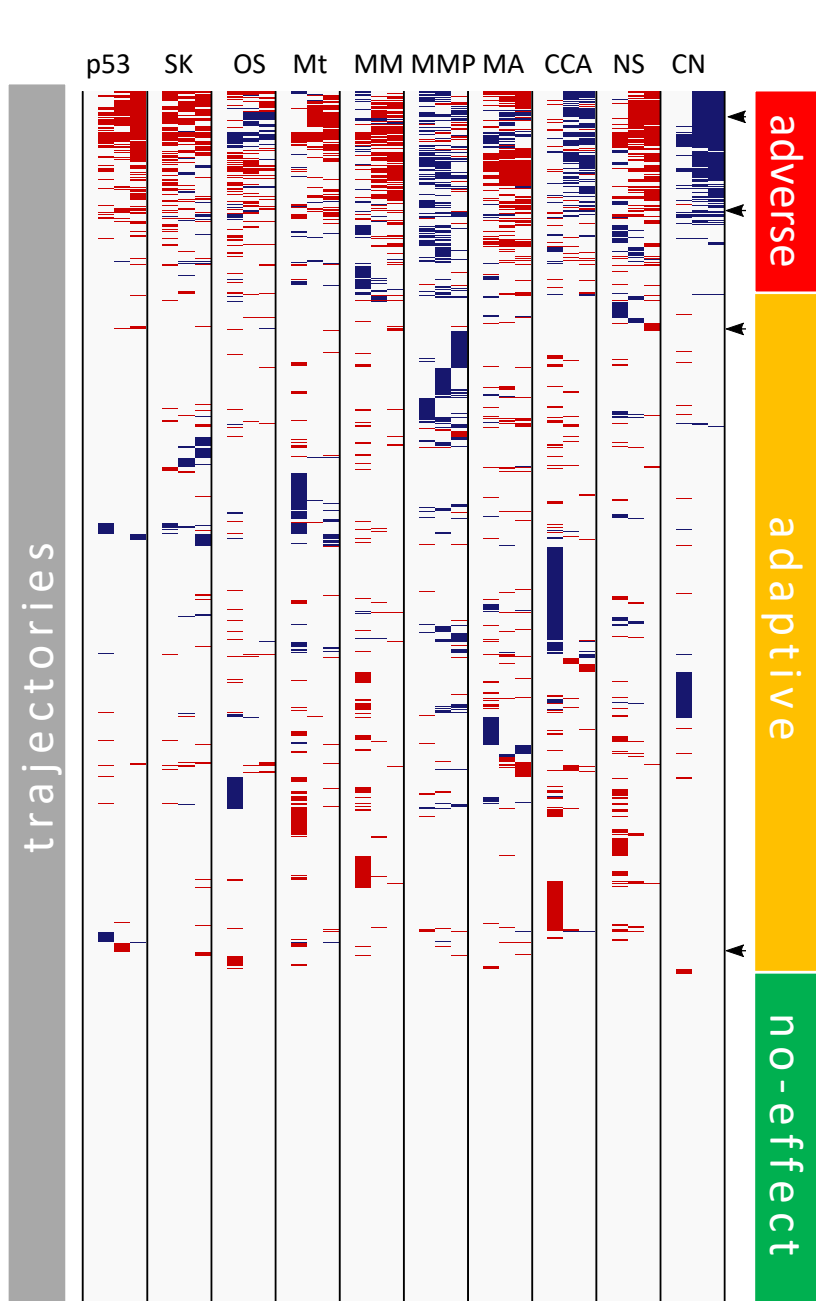
HCI Data



Boolean
Network
Inference

Predicted Mechanism





- Predicted 5×10^5 BN for each trajectory (10^9)
- Evaluated each BN against each trajectory
- Just 10 BN predict 80% of all trajectories
- Adverse trajectories

Summary

- ❑ Toxicological tipping points are critical points between adaptive and adverse trajectories
- ❑ Adaptive trajectories can be modeled by Boolean Networks
- ❑ Feasible to predict tipping points computationally (?)
- ❑ Need to understand chemical perturbations alter biological networks

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