

How to make an organ: Specifying cell fate during development

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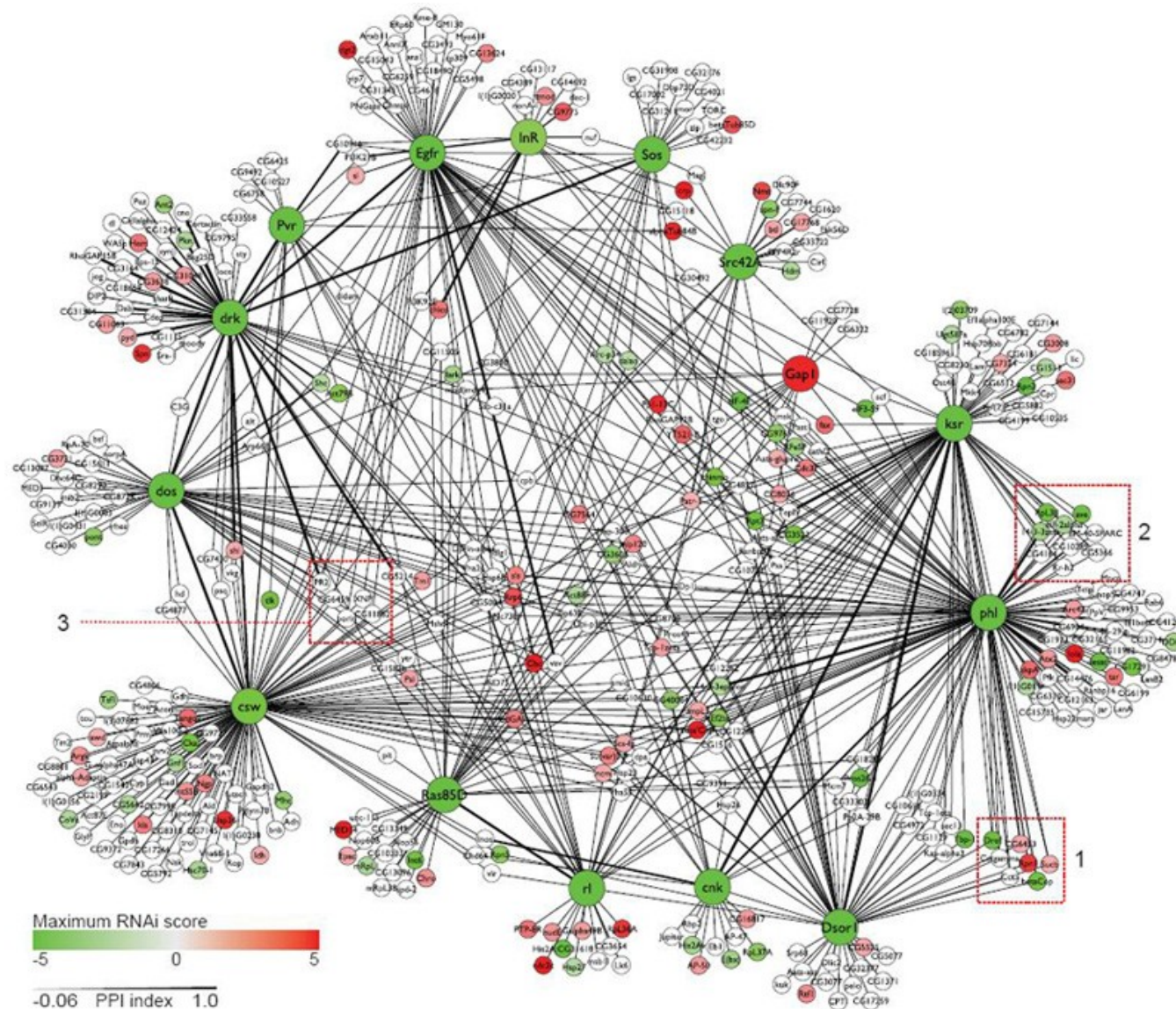


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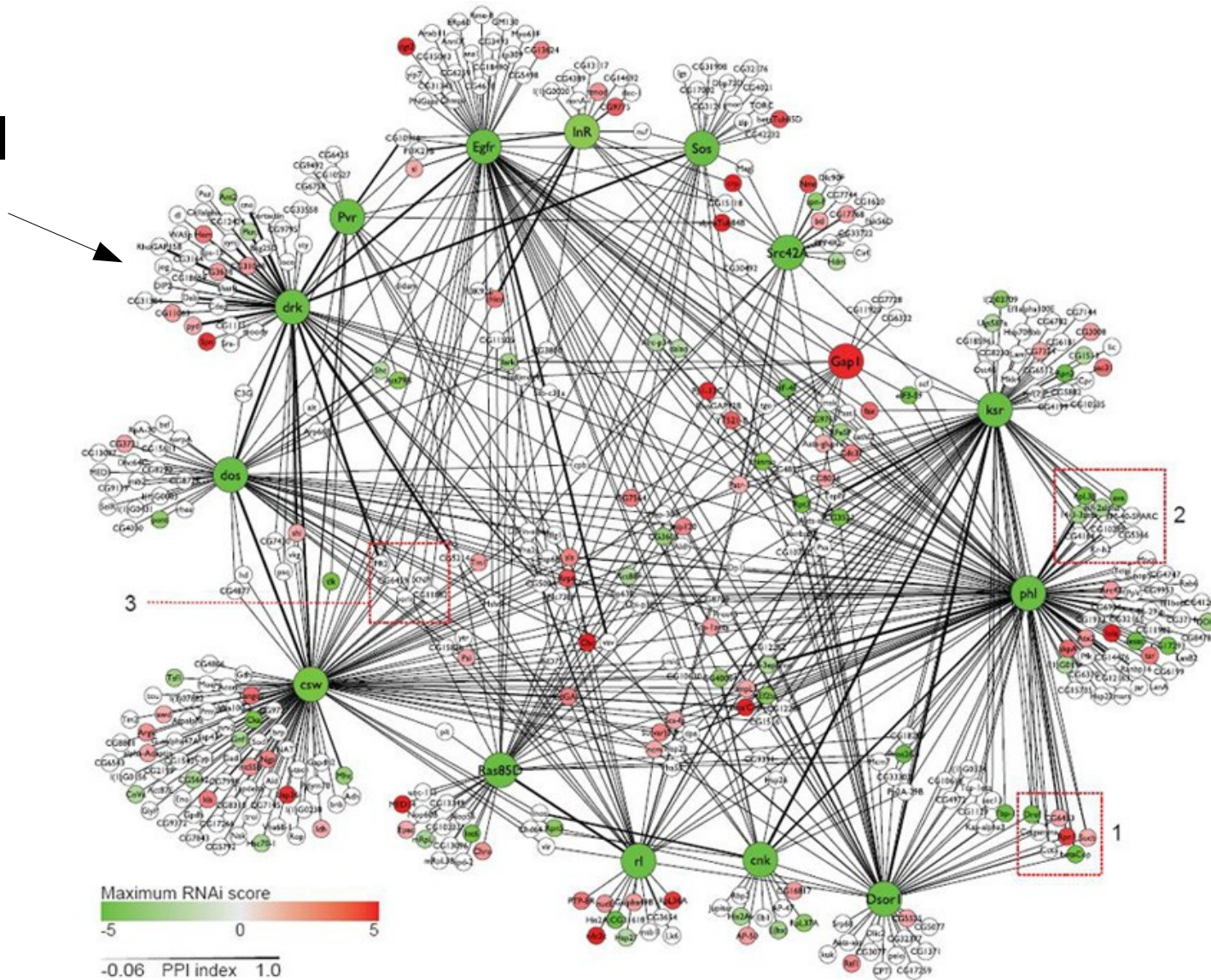
Signaling network determines output



Friedman et al (2011) *Sci Signal*

Signaling network determines output

External signal

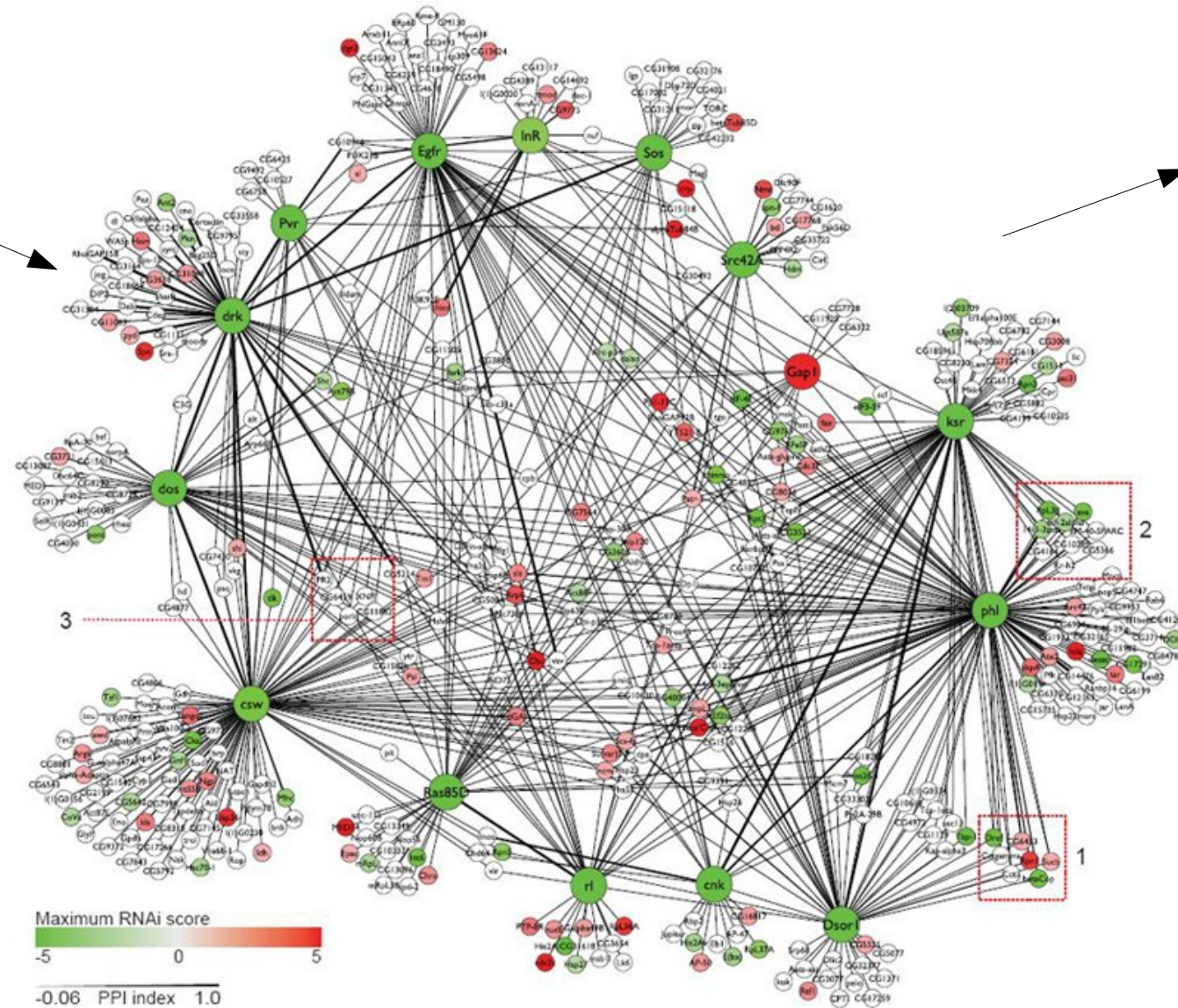


Friedman et al (2011) *Sci Signal*

Signaling network determines output

External
signal

Division

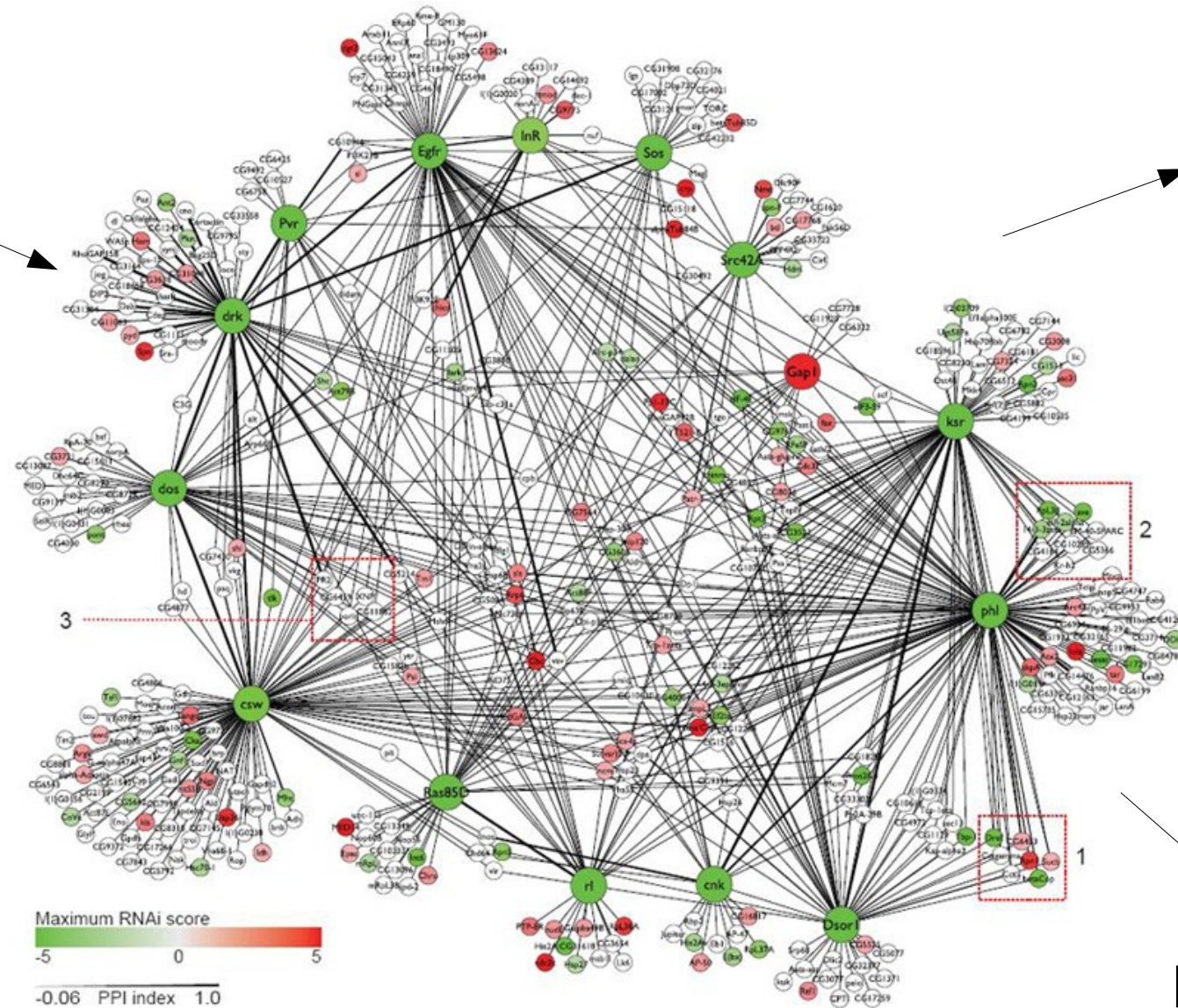


Friedman et al (2011) *Sci Signal*

Signaling network determines output

External
signal

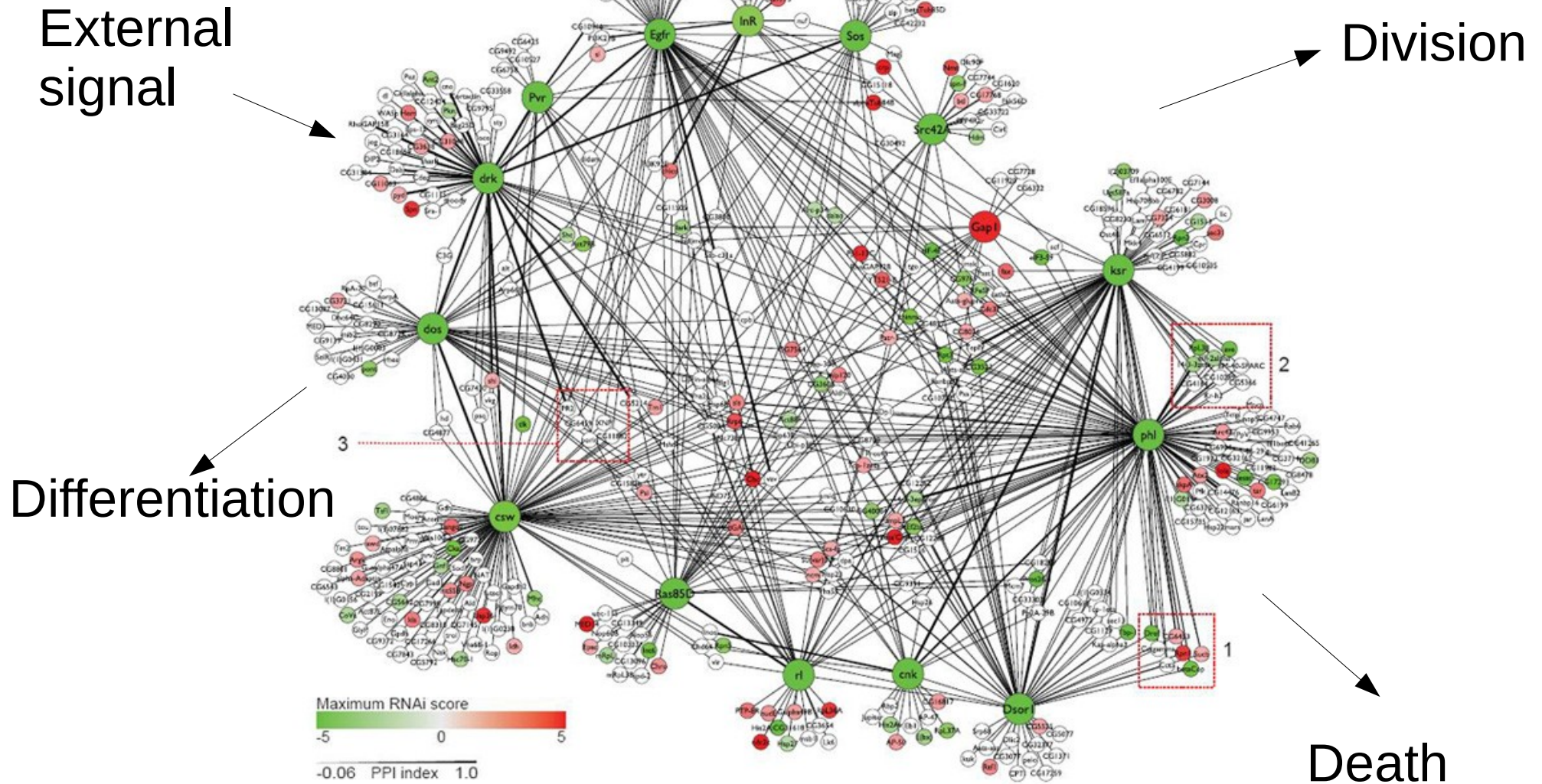
Division



Death

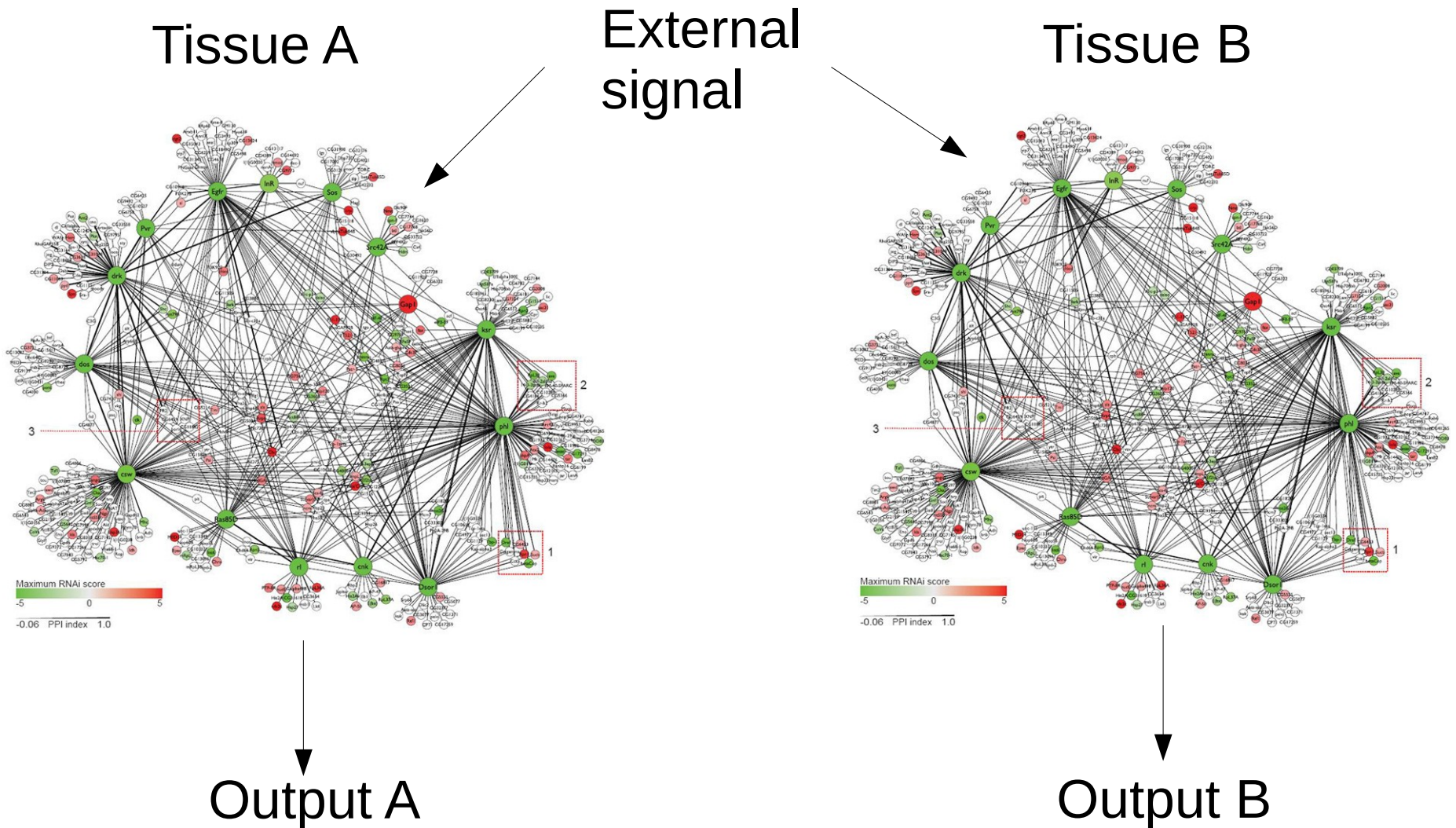
Friedman et al (2011) *Sci Signal*

Signaling network determines output

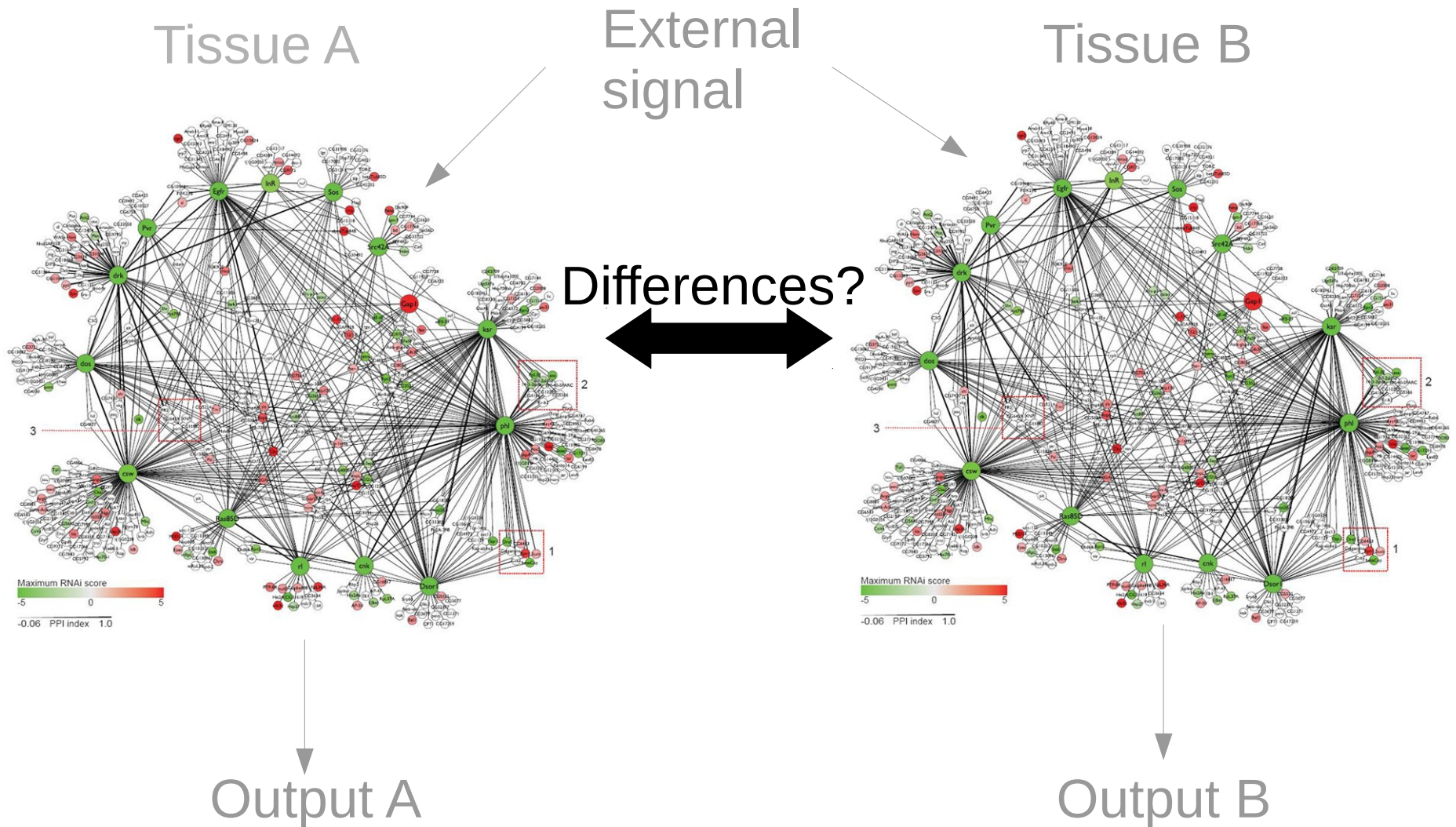


Friedman et al (2011) *Sci Signal*

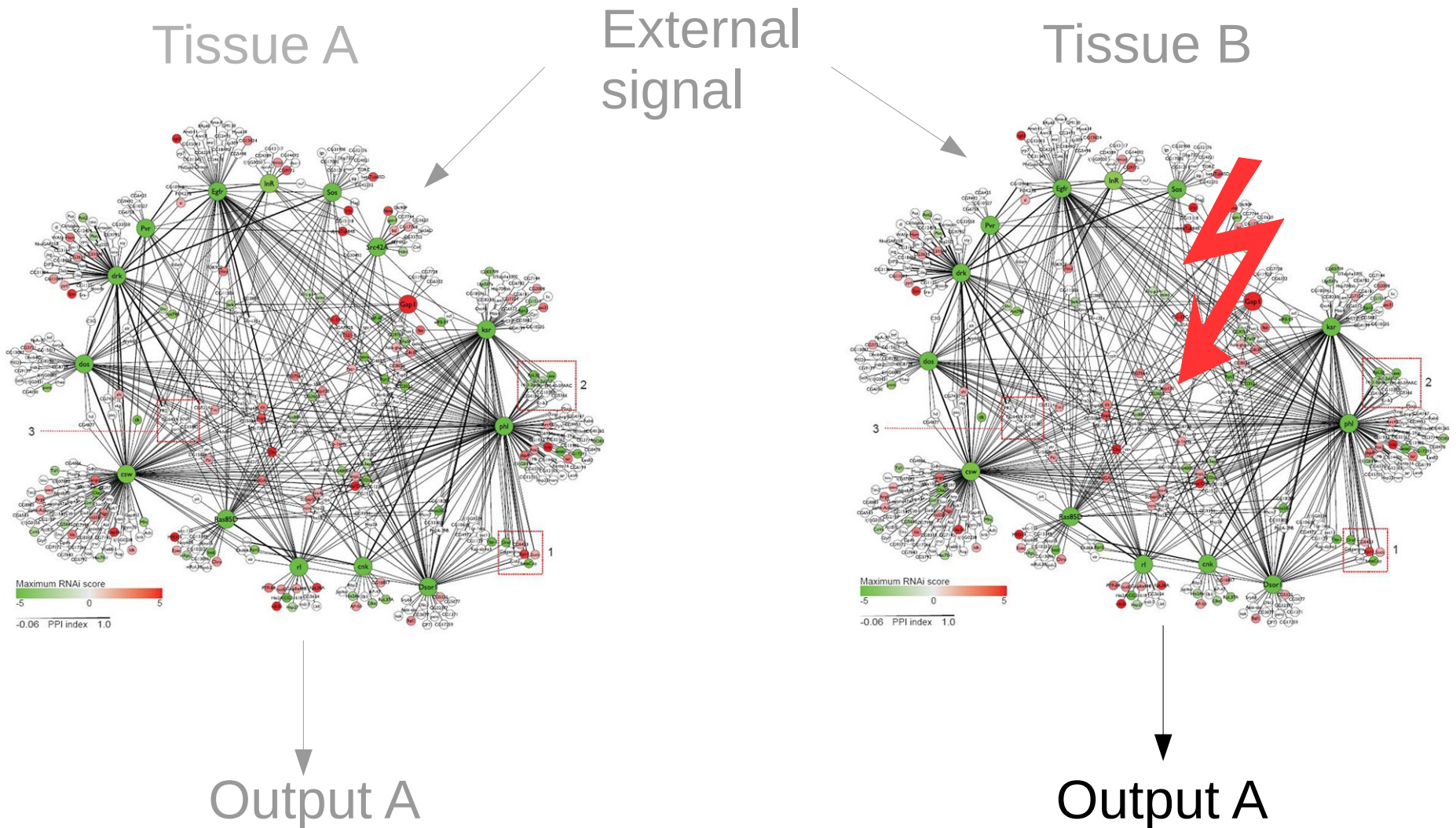
Tissue-specific response



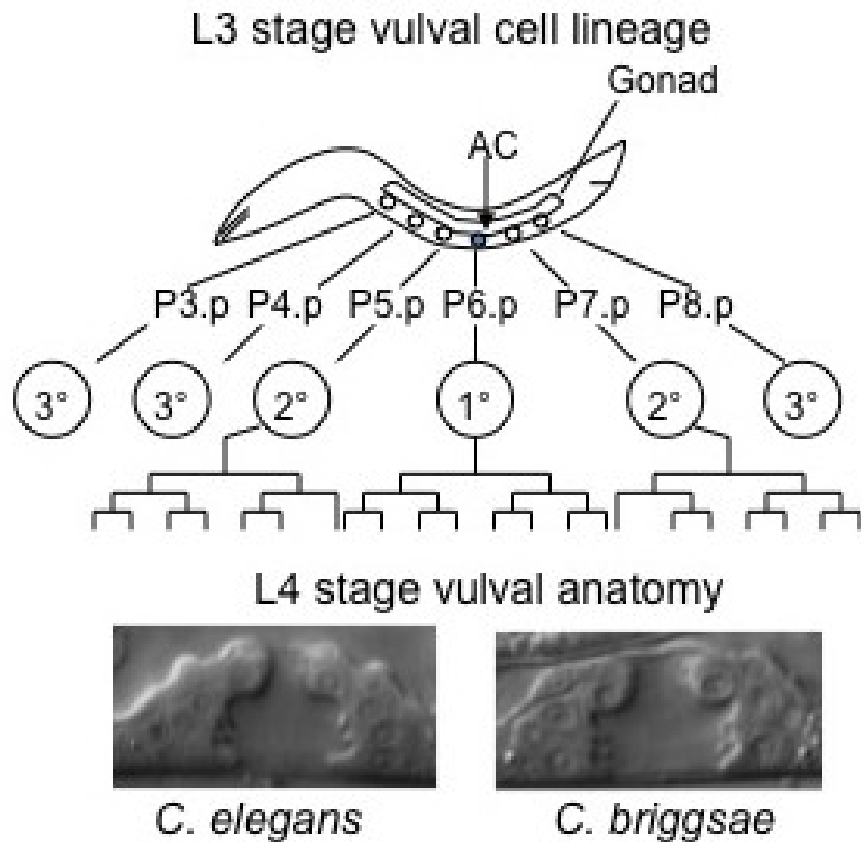
What accounts for observed differences?



Tune output by targeted manipulations?



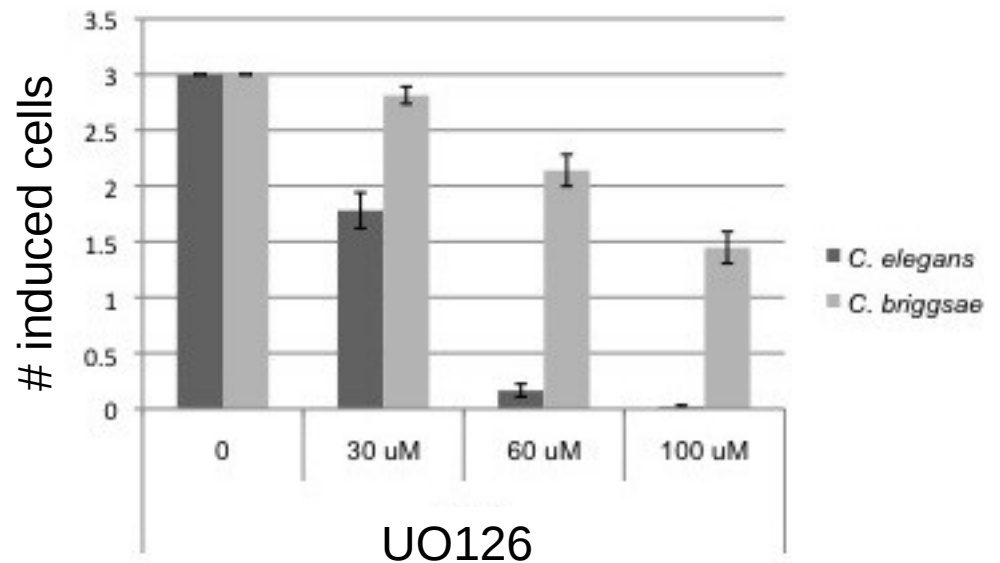
Vulval development in *Caenorhabditis*



- Egg laying structure
- VPC = Vulval Precursor Cell
- Anchor Cell (AC): signal source
- 1°, 2°: Induced
- 3°: Not induced
- Identical WT development in *C. elegans* and *C. Briggsae*
- Conservation of process and factors: EGF (inductive signal), Notch (lateral signal) and Wnt

VPC response to perturbation

- Drug treatment: gradual reduction in target activity
- Species-specific response
- Control: UO126 affects other processes

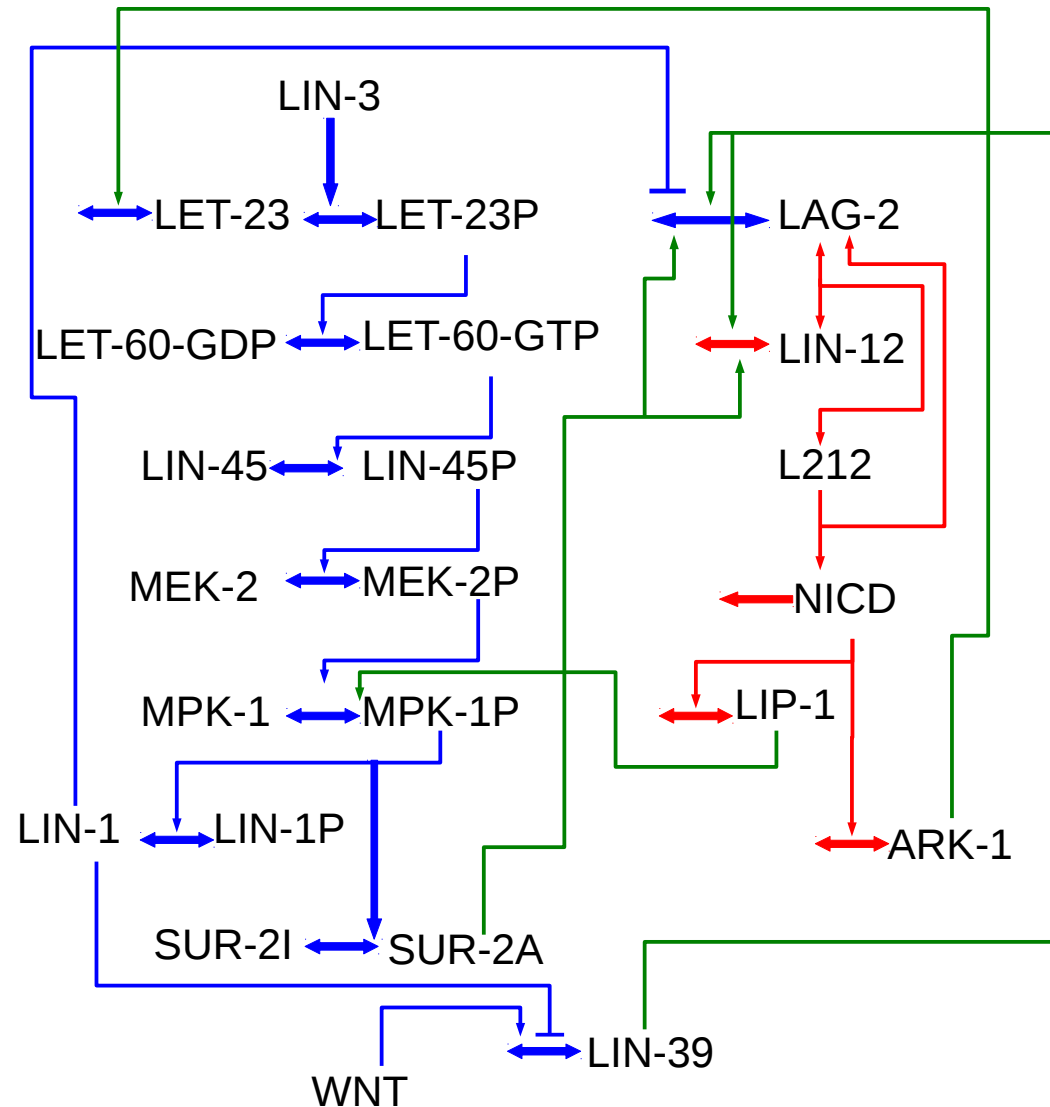


Big Question #1

What factors are responsible for the species-specific response?

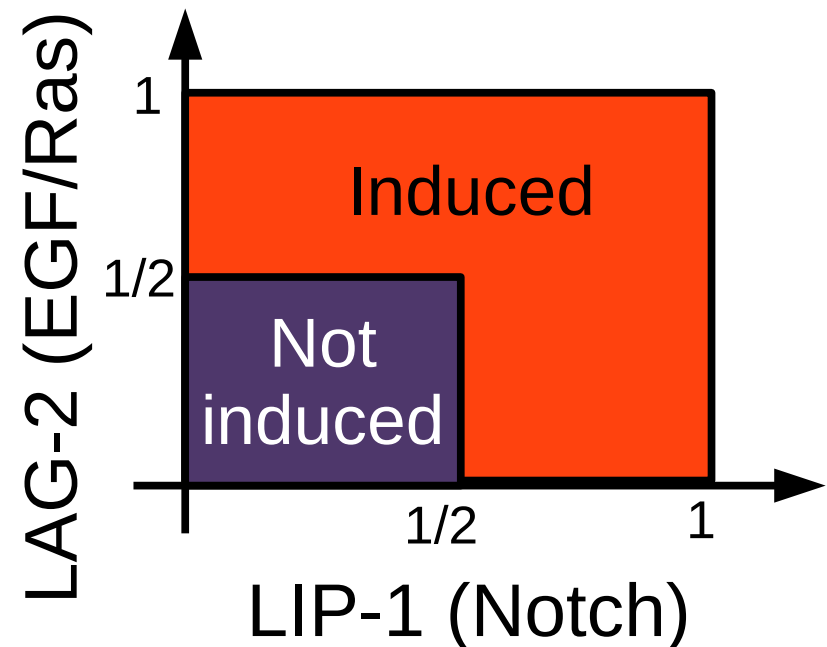
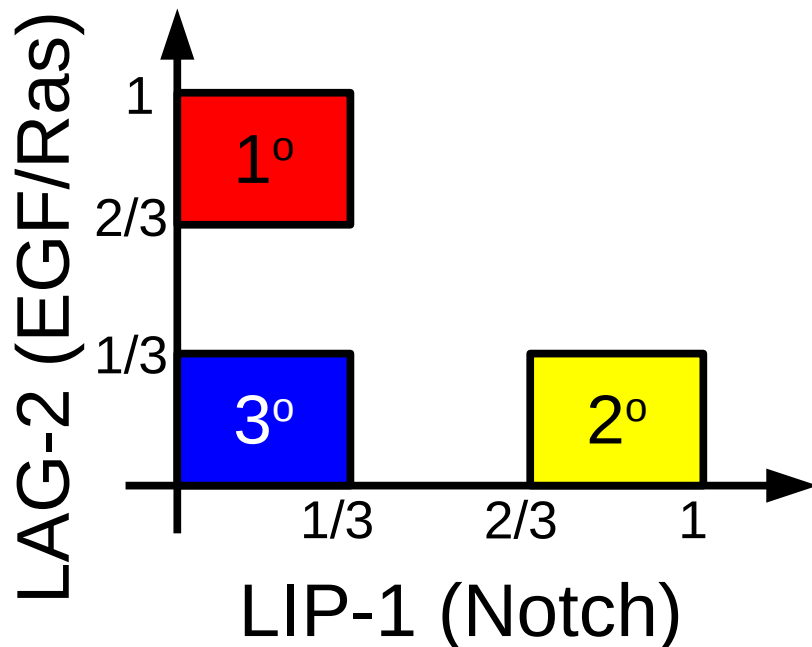
Biologically based model

- Most data from *C. elegans*: conservation
- **Blue = EGF/Ras**
- **Red = Notch**
- **Green = Crosstalk**
- Identical system of equations implemented in each VPC (6 x 15 eqns, 71 params)
- Spatially varying EGF, Wnt signal



Model Assumptions

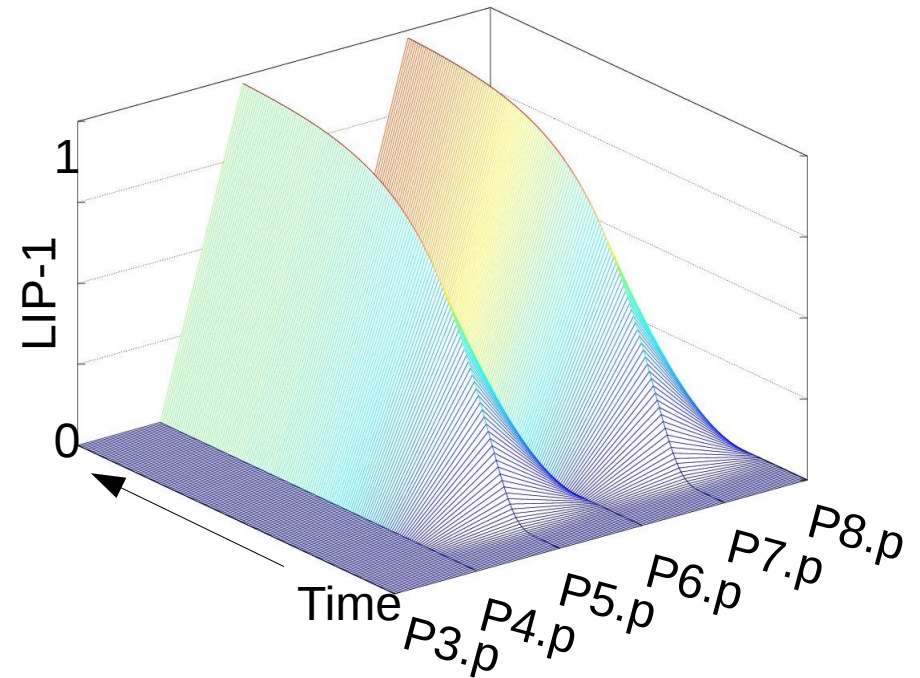
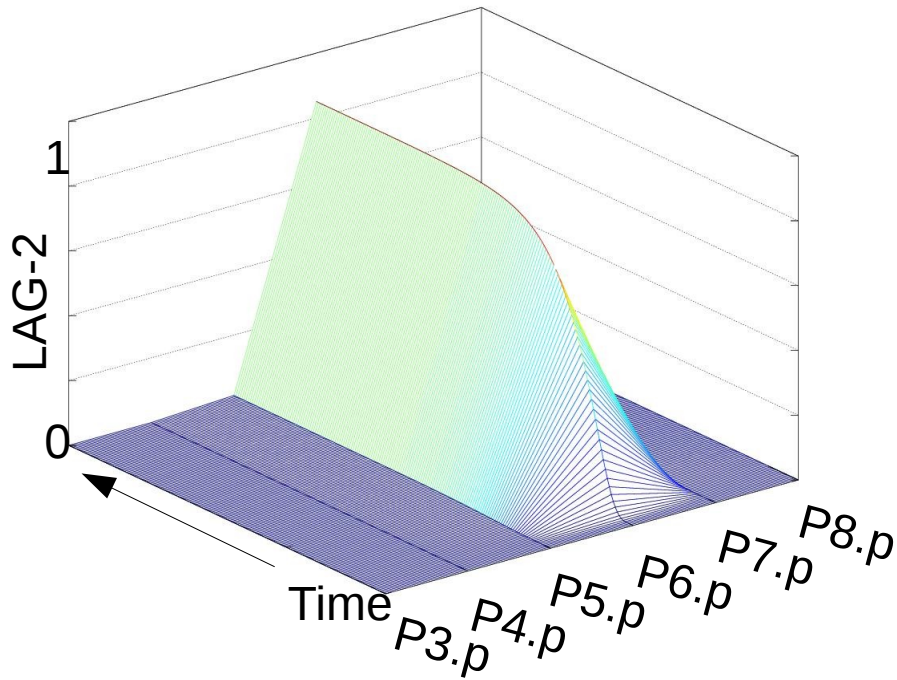
- Kinase/phosphatase activity follows Hill-type dynamics (saturating)
- All other dynamics linear/mass-action
- Cell fate determination:



Model recapitulates wild-type time course

EGF/Ras output

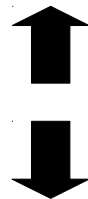
Notch output



LAG-2

LIP-1

Primary



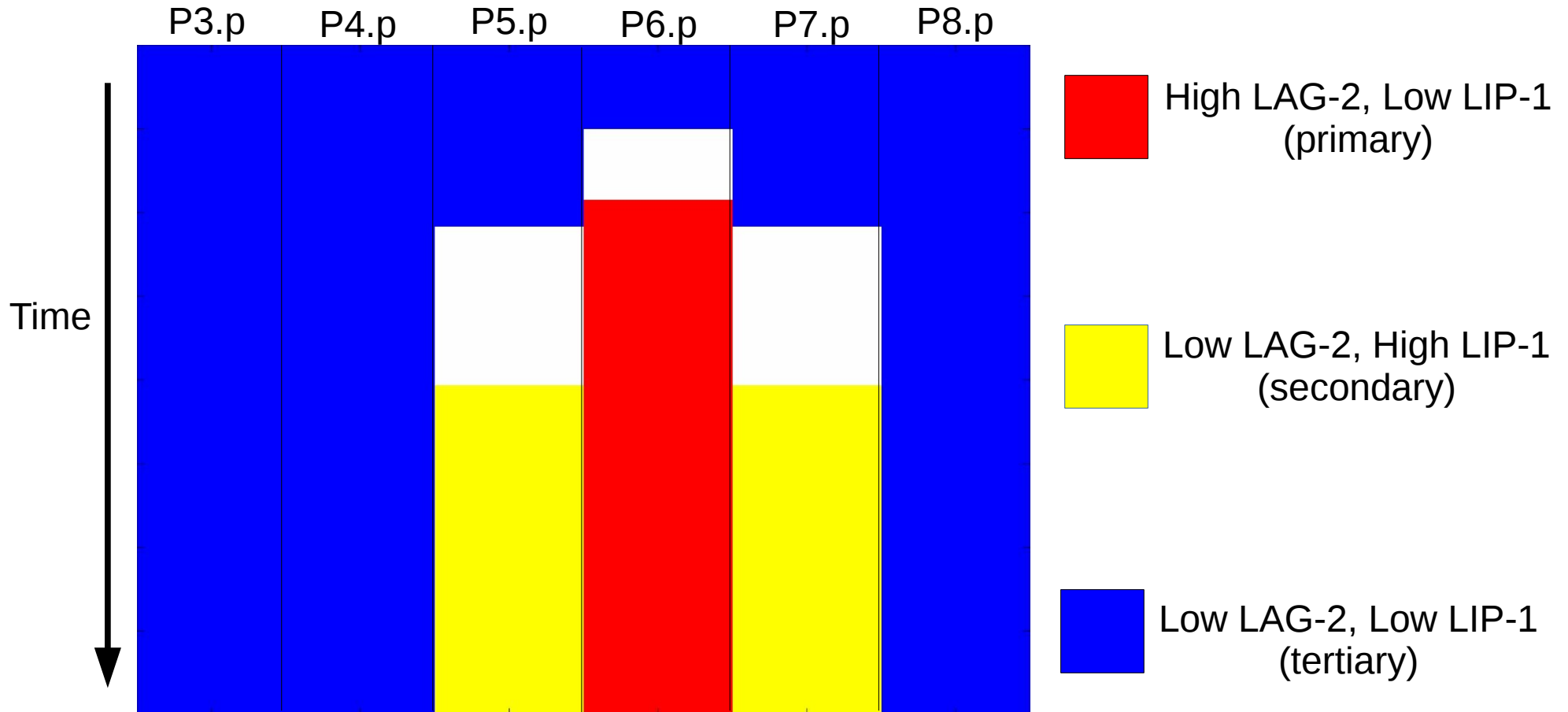
Secondary



Tertiary

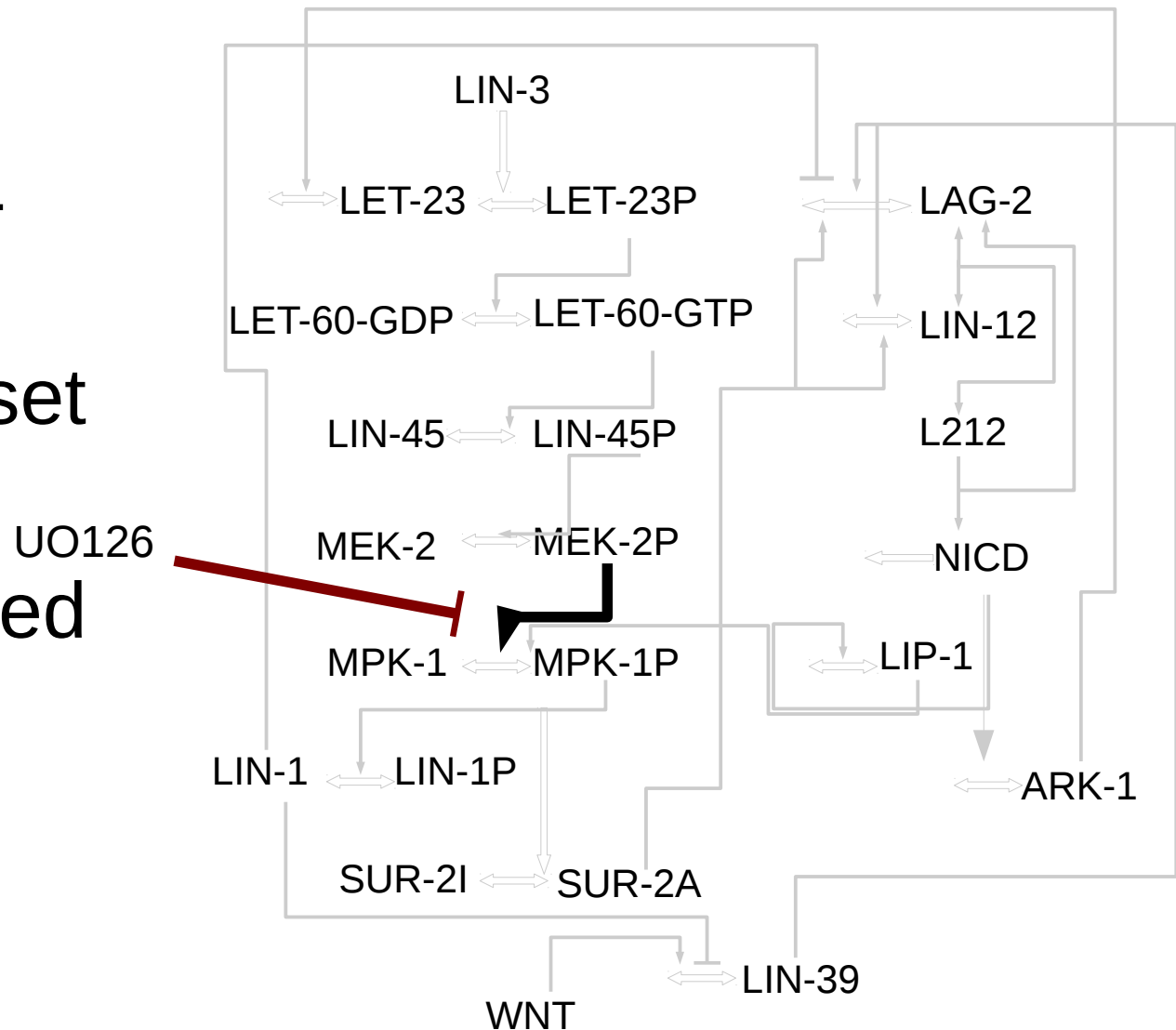


WT time course



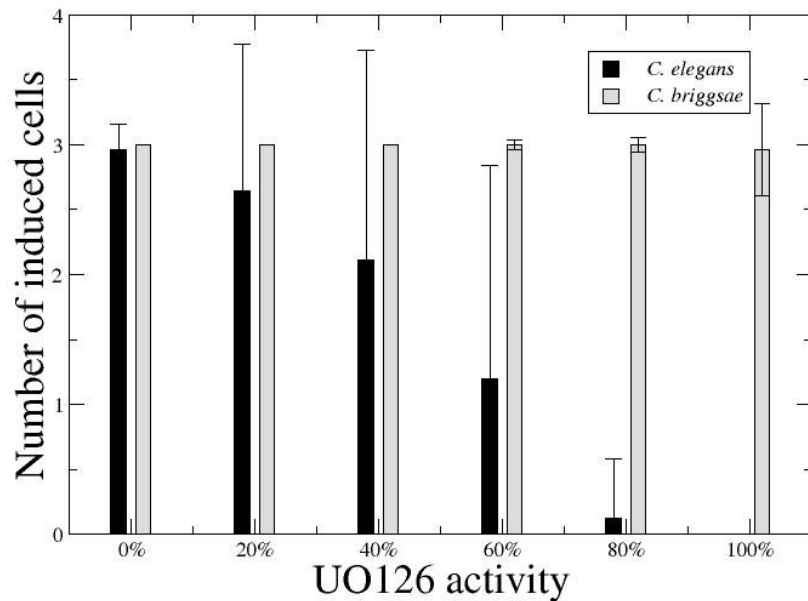
Role of UO126

- MEK-2 inhibitor
- Prevents MPK-1 phosphorylation
- Test parameter set responses:
number of induced cells

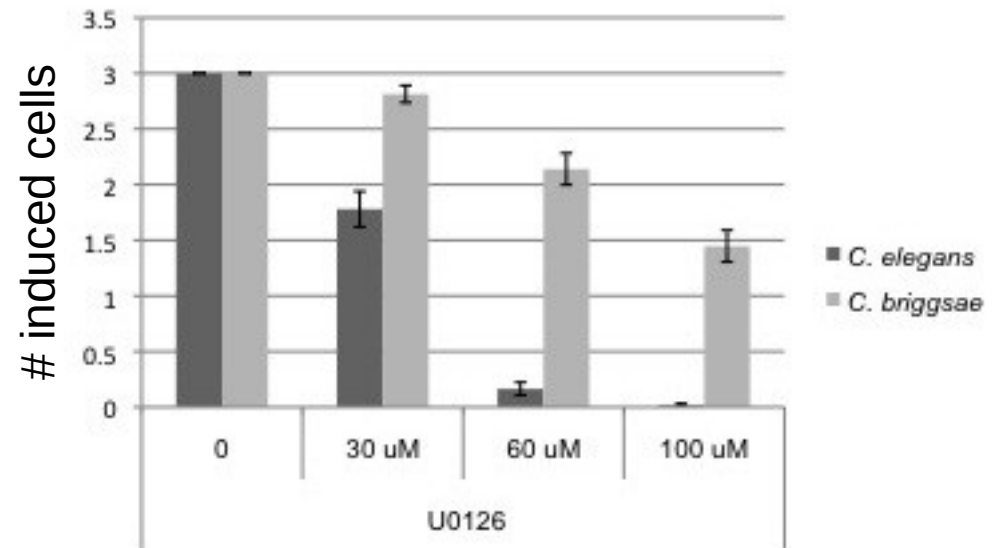


Simulations replicate response to UO126

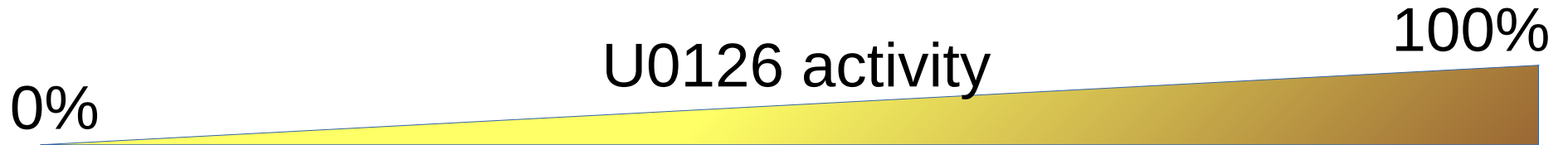
Numerical data



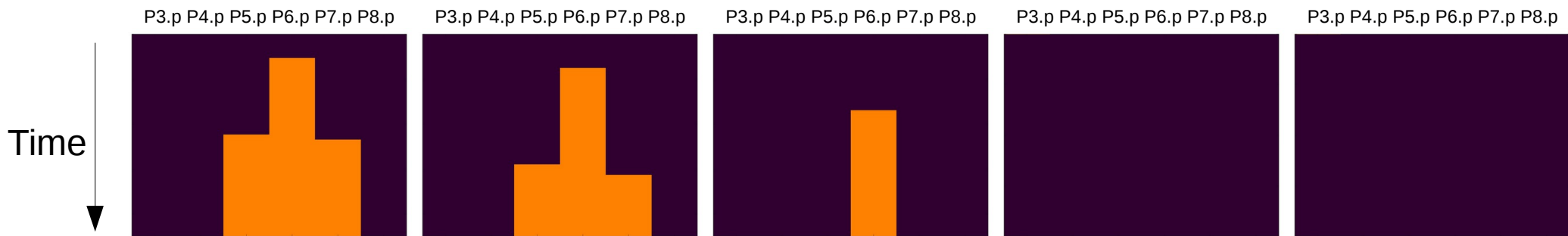
Experimental data



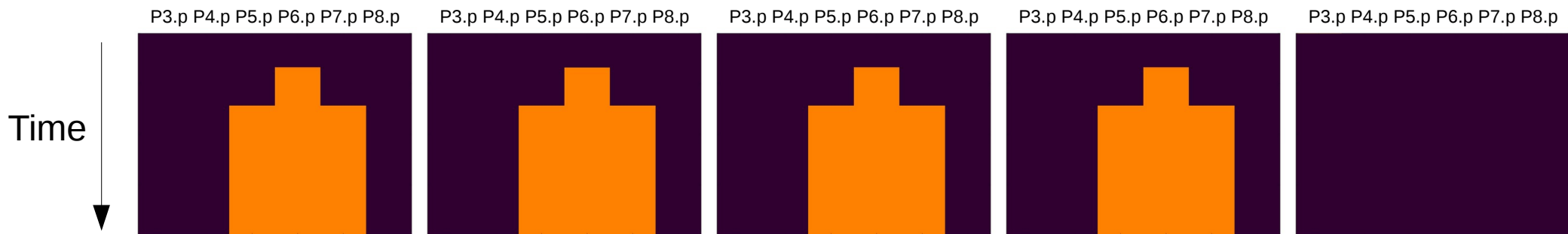
Simulated response to UO126



C. elegans



C. briggsae

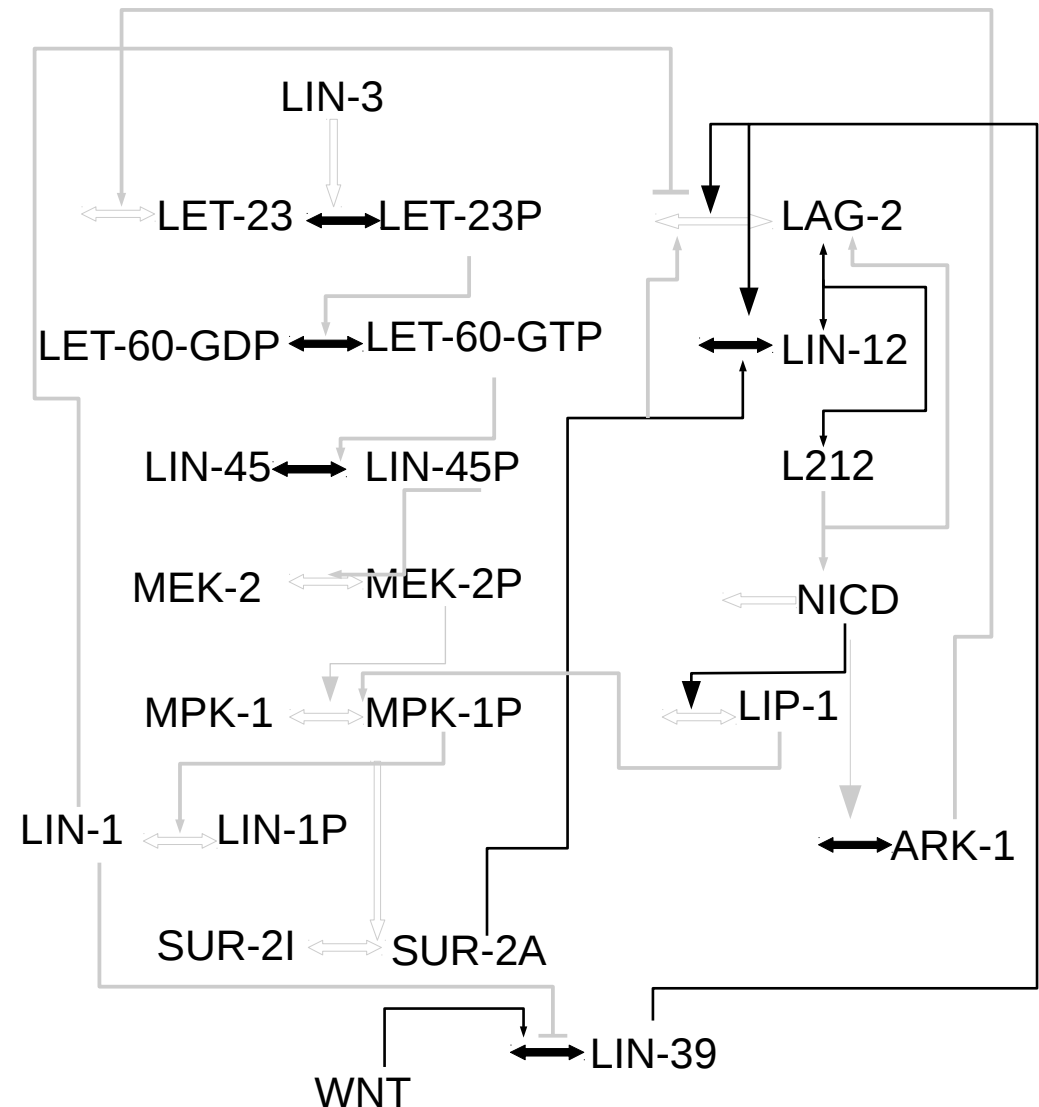


Induced

Not induced

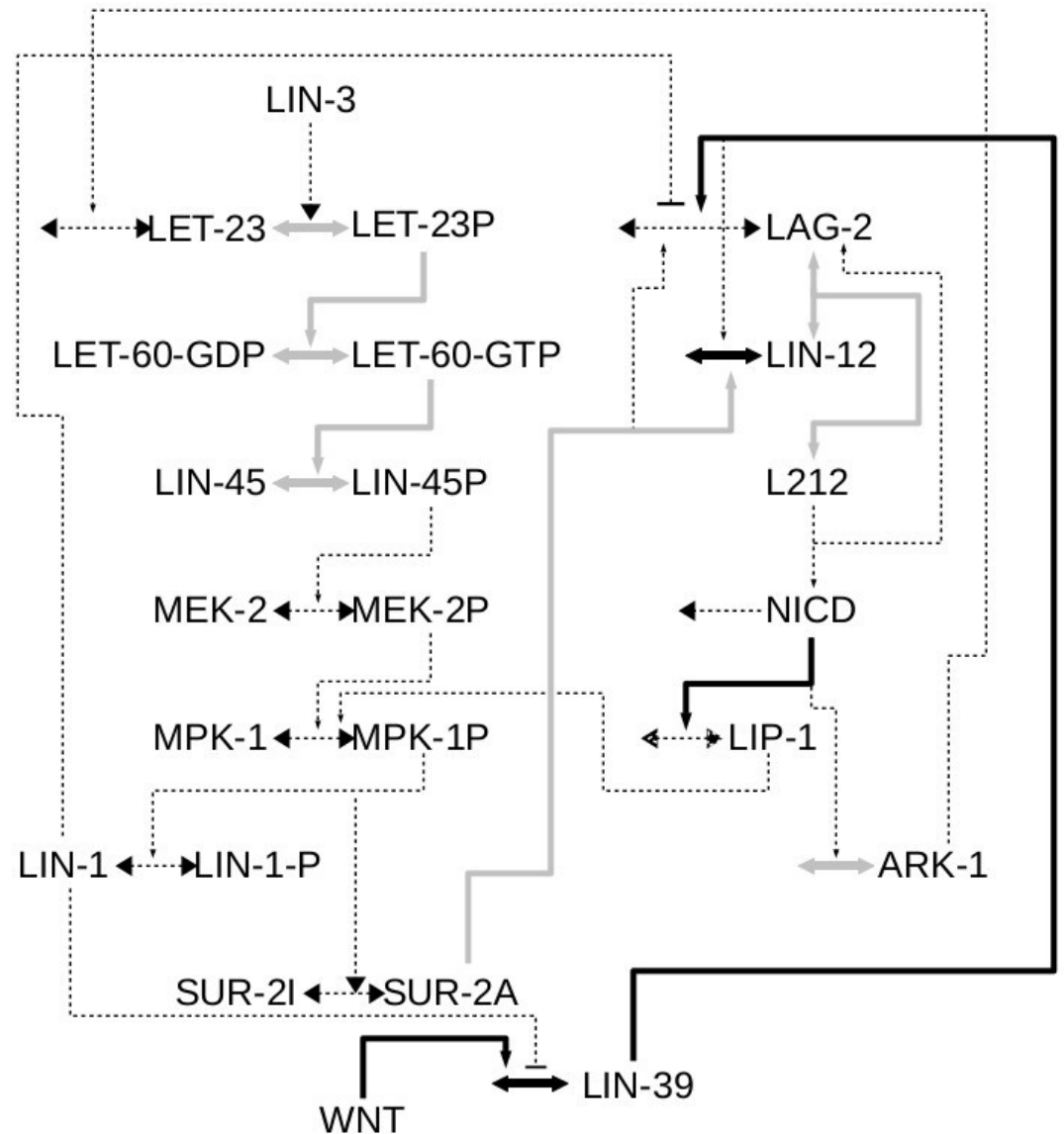
Parameter differences in EGF synthesis, Notch activation

- Two sample t-test between *C. elegans* and *C. briggsae*-like sets



Phenotype “switching” by key parameters

- Parameters modulated one at a time and response tested



Results #1

- Constructed biologically based model that recapitulates wild-type behaviour
- External Wnt signal identified as key parameter for species-specific response

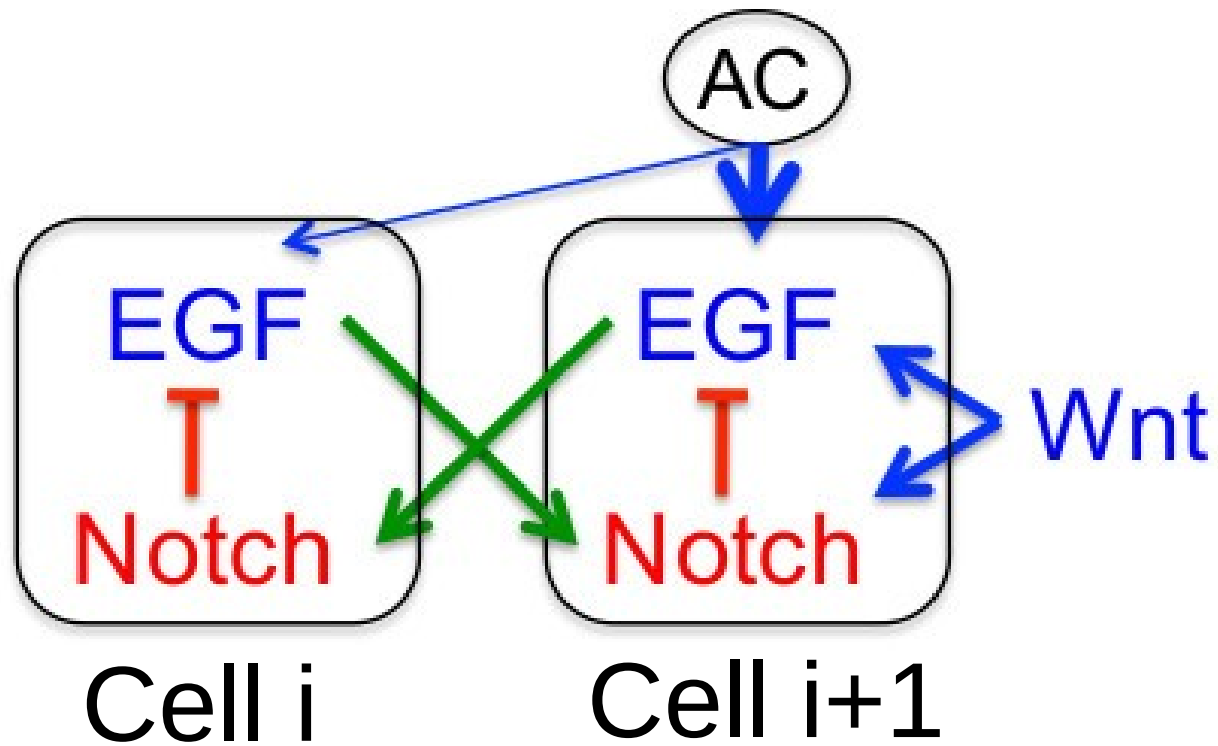
Big Question #2

How is the Wnt signal affecting cell fates?



Carly Williamson

Simplified network



Model equations

$$\text{Notch} \quad \frac{dN_i}{dt} = W_i f(\bar{E}) - N_i$$

$$\text{EGF} \quad \frac{dE_i}{dt} = \gamma(\lambda_i g(N_i) - E_i)$$

$$\text{where} \quad \bar{E}_i = \frac{1}{m} \sum_j E_j$$

$$f(\sigma) = \frac{\sigma^k}{\alpha + \sigma^k}$$

$$g(\sigma) = \frac{1}{1 + \beta \sigma^h}$$

Model equations

$$\begin{array}{l} \text{Notch} \quad \frac{dN_i}{dt} = W_i f(\bar{E}) - N_i \\ \text{EGF} \quad \frac{dE_i}{dt} = \gamma (\lambda_i g(N_i) - E_i) \end{array} \quad \gamma = \frac{k_E}{k_N}$$

$$\text{where} \quad \bar{E}_i = \frac{1}{m} \sum_j E_j$$

$$f(\sigma) = \frac{\sigma^k}{\alpha + \sigma^k}$$

$$g(\sigma) = \frac{1}{1 + \beta \sigma^h}$$

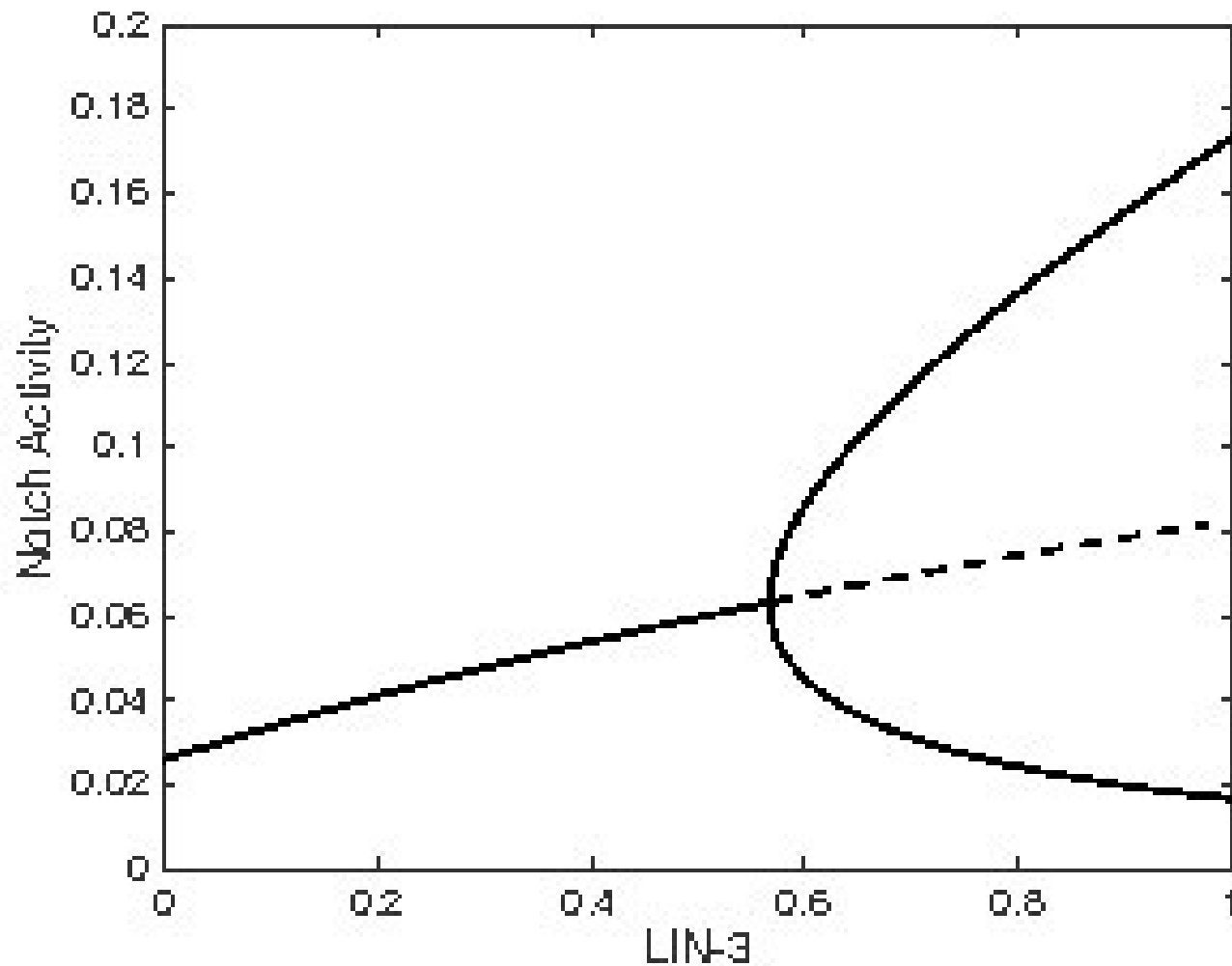
Linear stability analysis

- Solve for steady state E^* , N^*
- Linearize system of equations and find eigenvalues
- Bifurcation point:

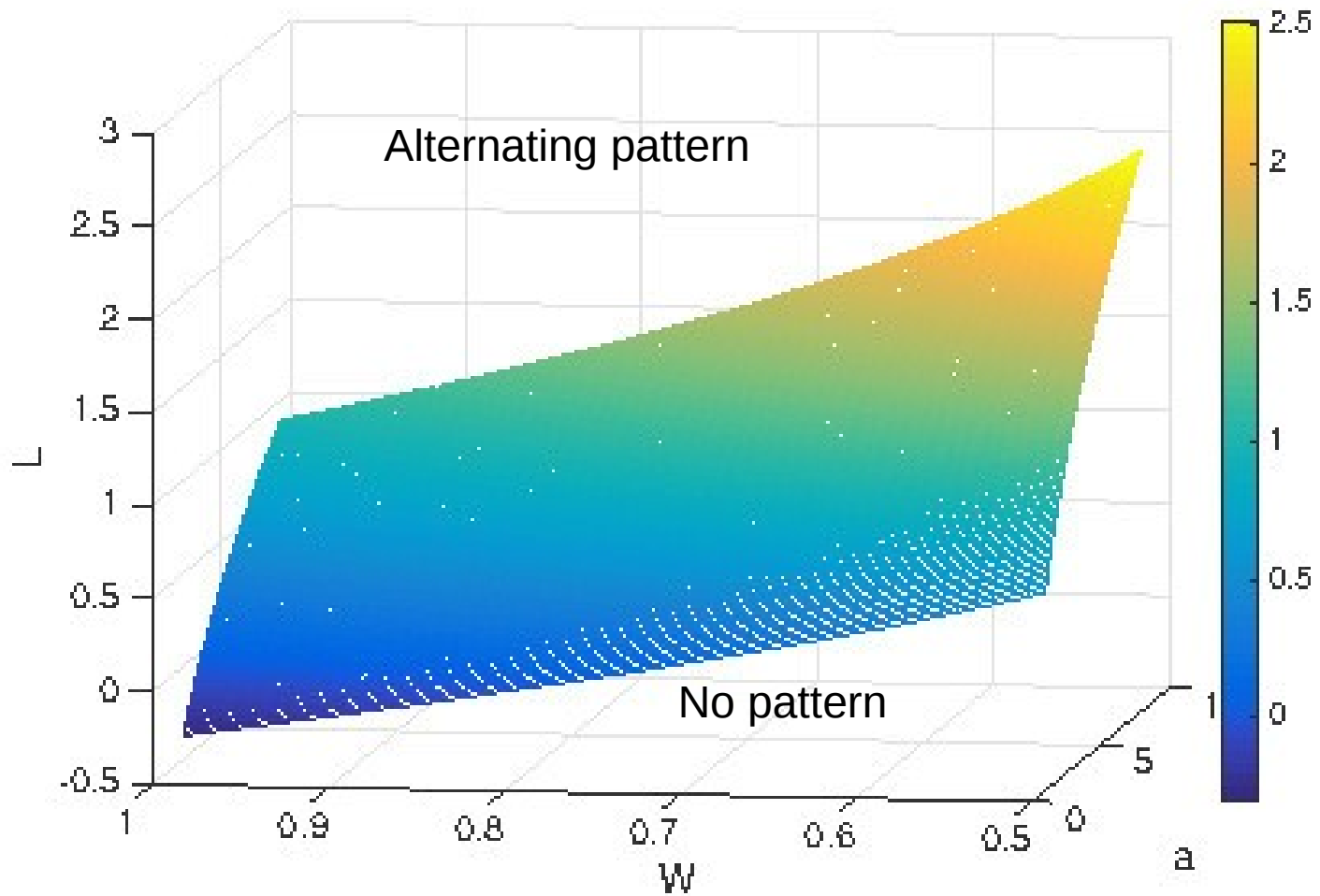
$$L = \frac{-1}{\mu} \left(\frac{\mu + 1}{W f'(E^*) g'(N^*)} + W \right)$$

Bifurcation diagram

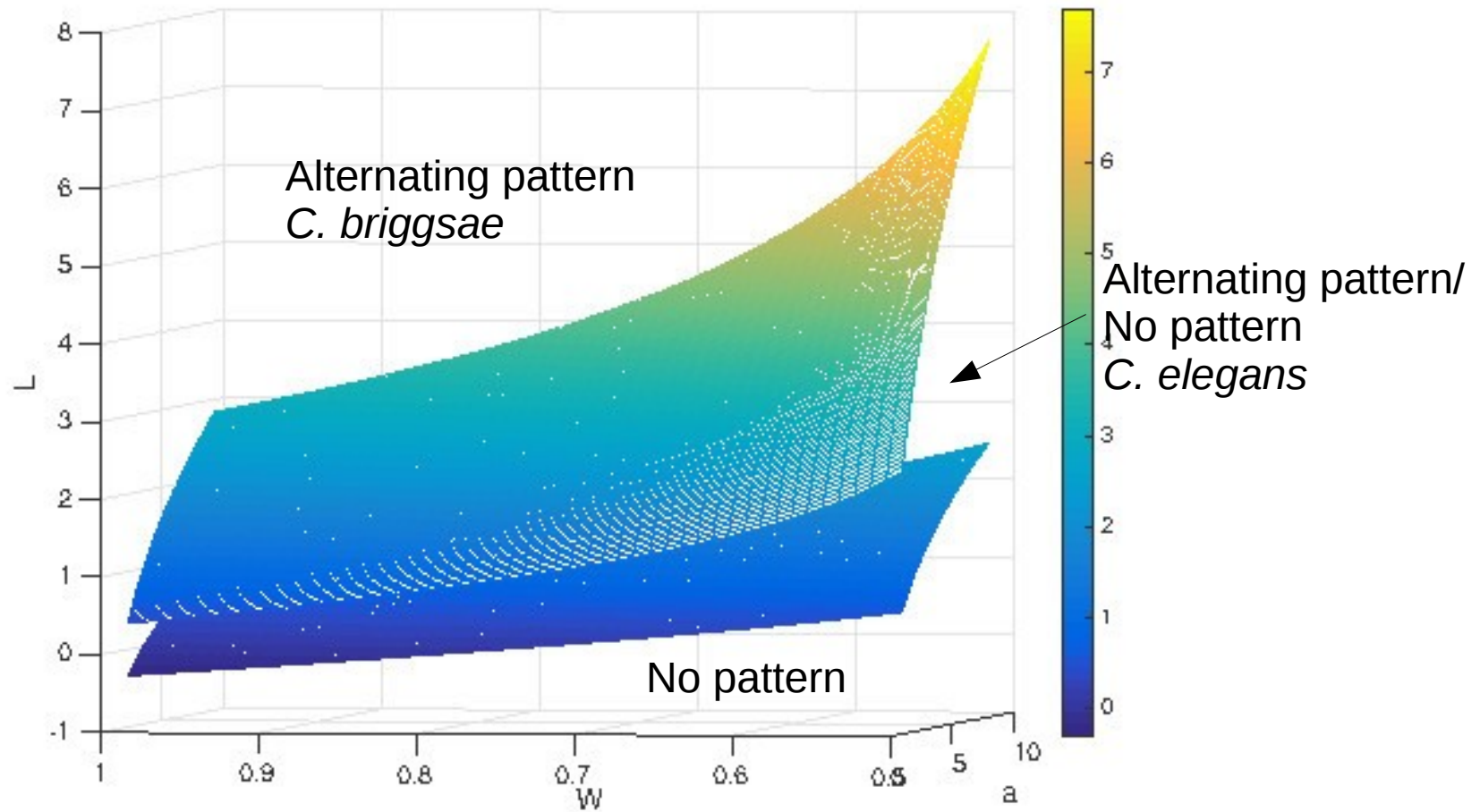
- Supercritical pitchfork bifurcation



Bifurcation plane



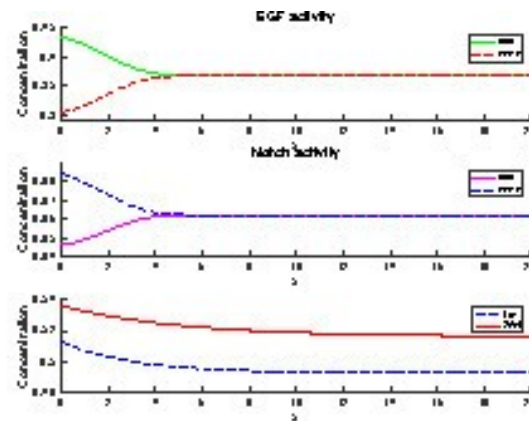
Bifurcation plane with EGF knock down



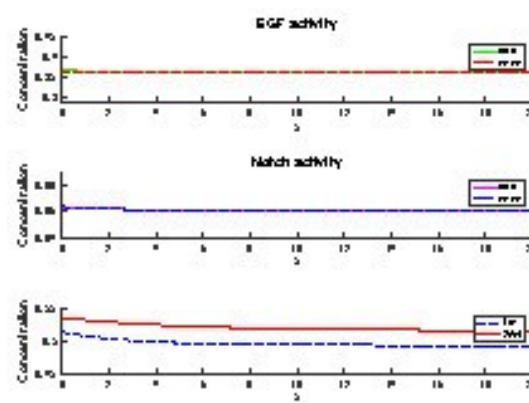
Asymptotics about bifurcation point

$$L = l + \delta^2 L(x)$$
$$W = w + \delta^2 W(x)$$

C. briggsae



C. elegans



Results #2

- Simplified model recapitulates loss of pattern
- Interrelationship between LIN-3, Wnt and sensitivity to external signals dictates loss of pattern
- Tuning Wnt signal can rescue alternating pattern

Open questions

- What if γ is very large?
- Consider $\varepsilon \rightarrow 0$

$$\begin{aligned} \text{Notch} \quad \frac{dN_i}{dt} &= W_i f(\bar{E}) - N_i \\ \text{EGF} \quad \frac{dE_i}{dt} &= \gamma (\lambda_i g(N_i) - E_i) \\ \Rightarrow \quad \frac{dN_i}{dt} &= W_i f(\bar{E}) - N_i \\ \varepsilon \frac{dE_i}{dt} &= \lambda_i g(N_i) - E_i \end{aligned}$$

Open questions

- Does this system satisfy conditions of Tikhonov's theorem?
- How does small parameter assumption affect bifurcation dynamics? Asymptotics?
- Do we gain new insight into the system dynamics by taking the limit?

Acknowledgments

Lab members:

Valerie Coffman

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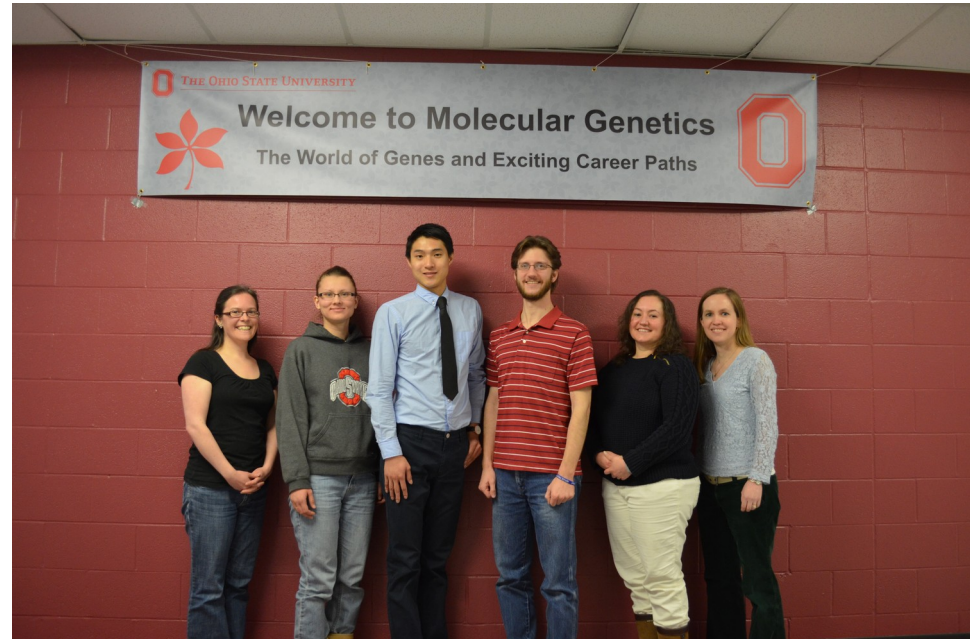
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David Ignacio

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Lawrence Small

Carly Williamson



Chamberlin Lab

