

Mechanism and function of stochastic pulse regulation

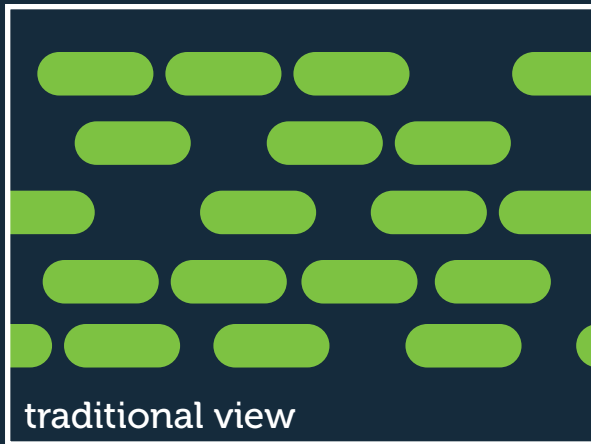
Stephanie Unna, Locke Lab.



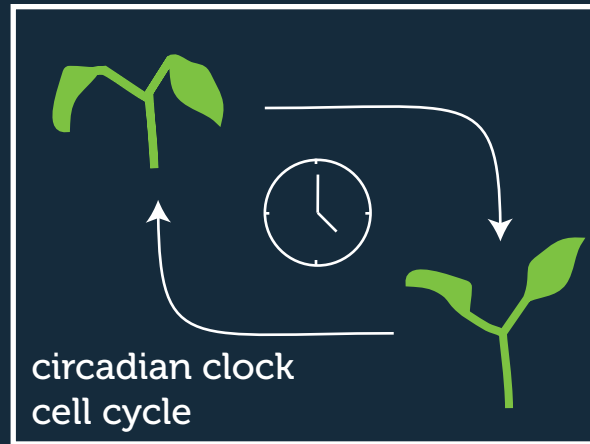
Gene expression is noisy

even under constant environmental conditions

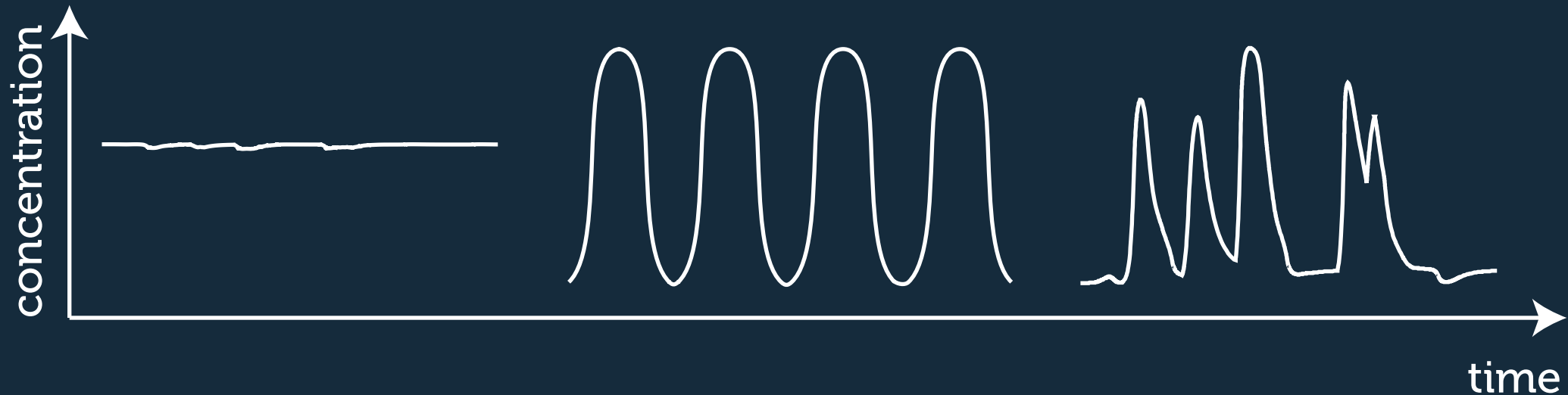
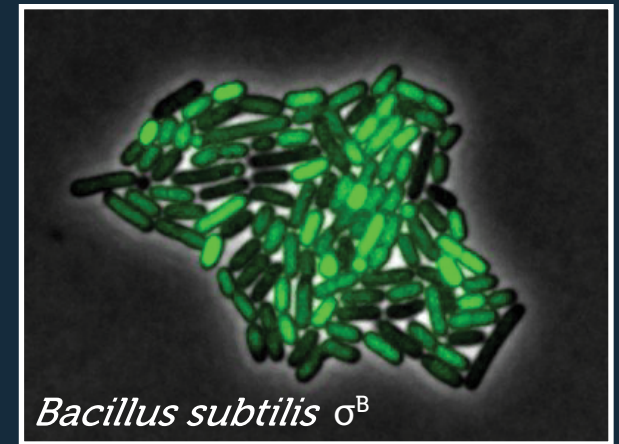
equilibration



oscillation



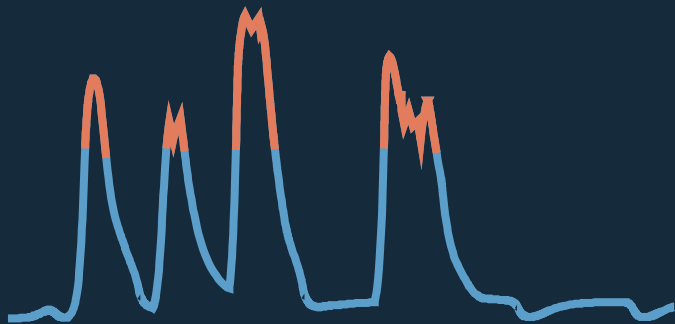
stochastic



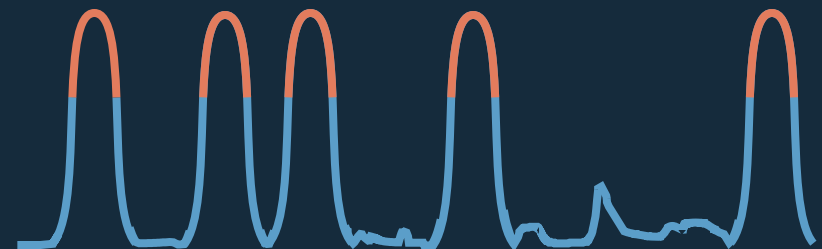
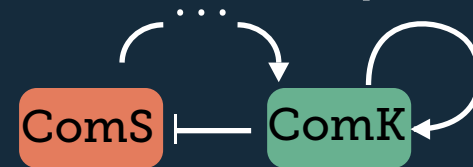
Noise is functionally important



- Circuit
 - stochastic
 - threshold
 - positive feedback
- Persistence *E. coli*
 - HipBA toxin/antitoxin
 - antibiotic resistance

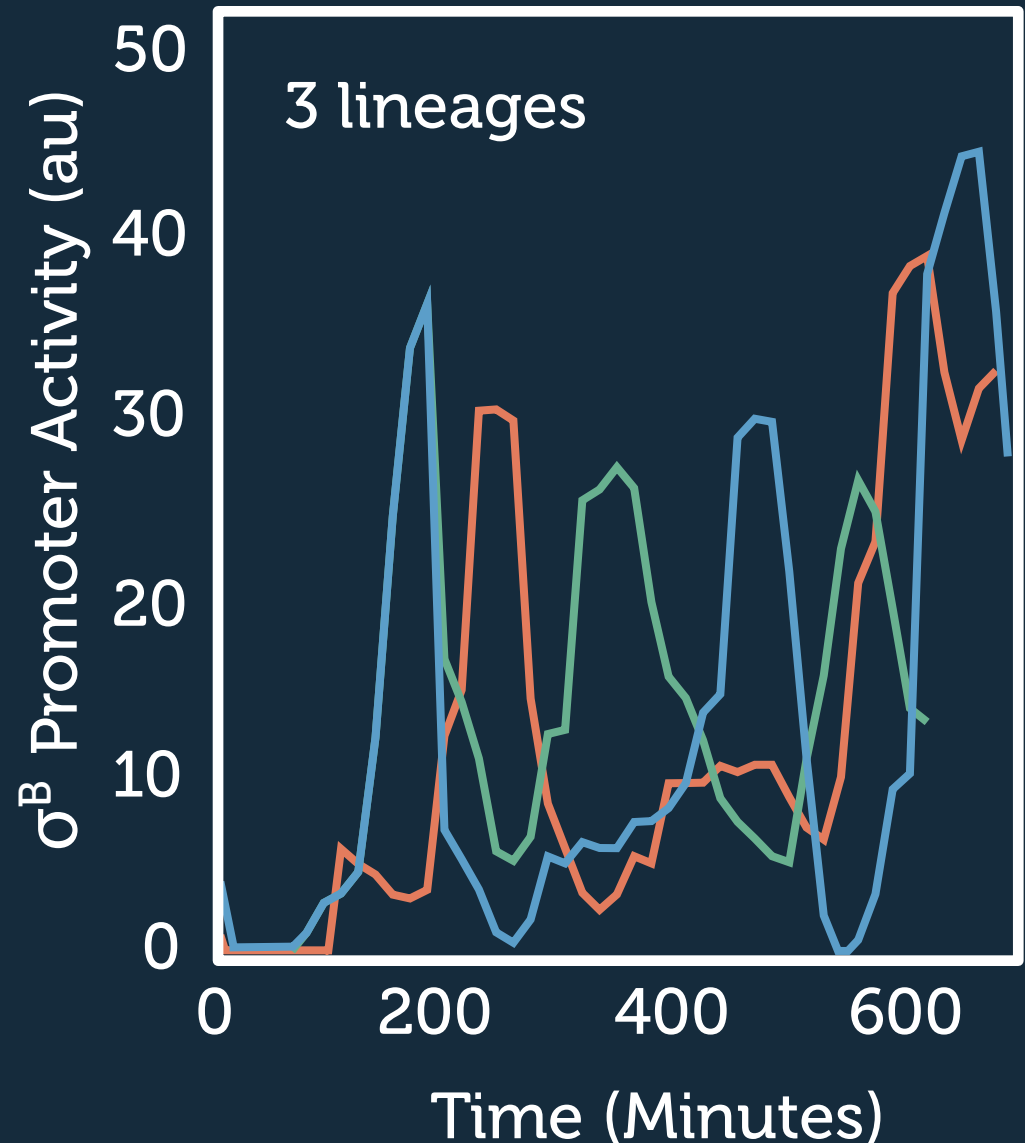


- Characteristic response
- Circuit:
 - stochastic process
 - threshold
 - fast/slow feedback
- Bacterial competence



A different kind of noise

stochastic pulsing



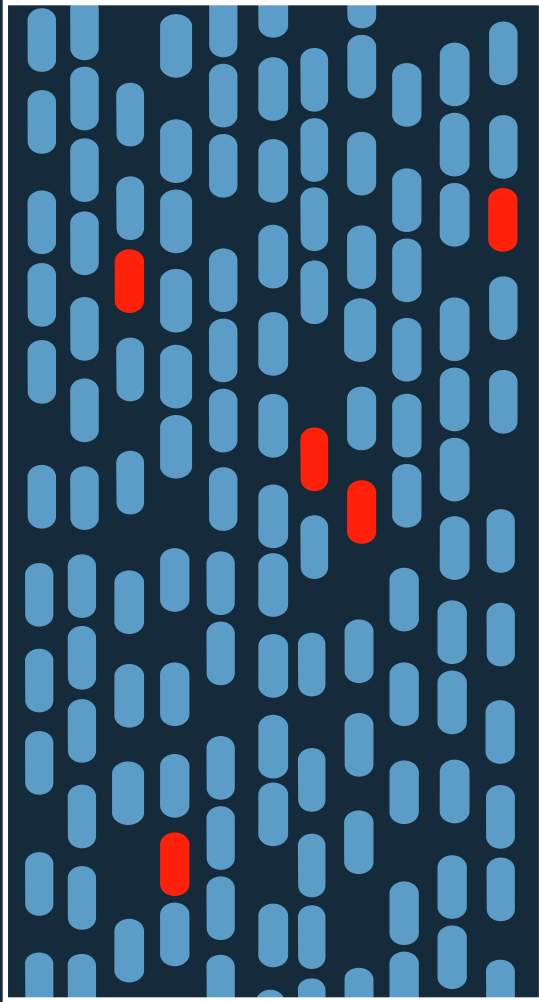
Bacillus subtilis P_{σ^B}-yfp

energy stress

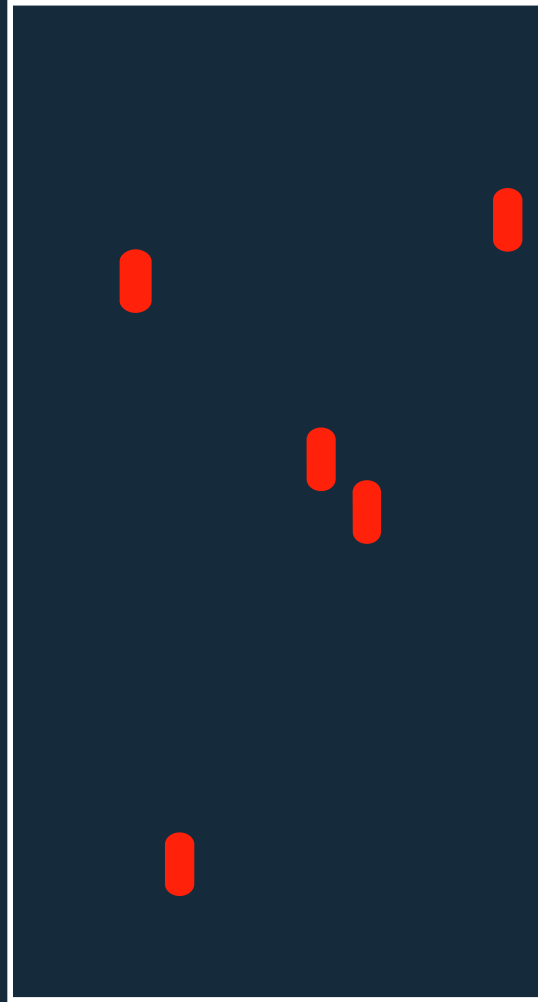
40μg/L MPA

What is the role of stochastic pulsing?

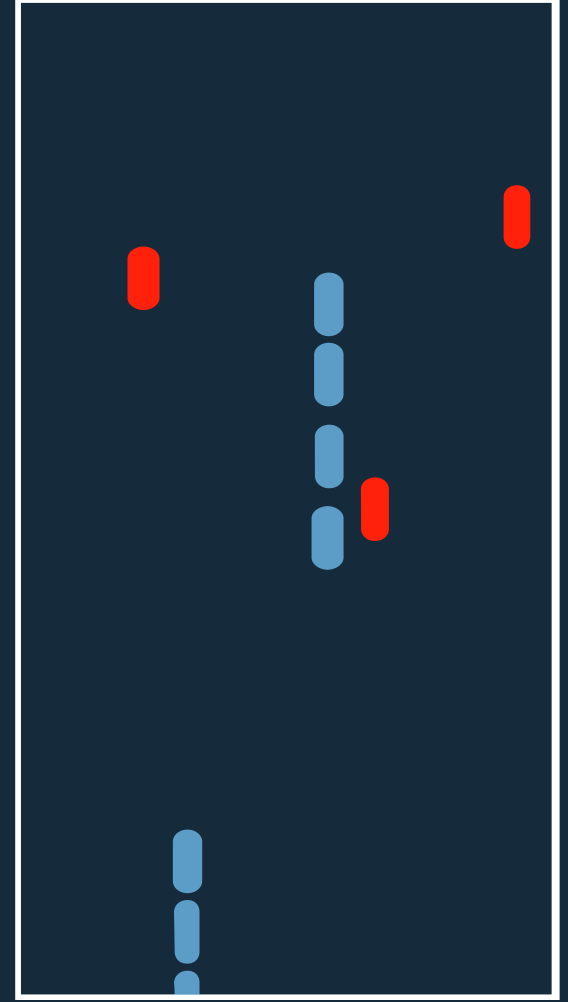
Bet-hedging in an uncertain environment



Optimal growth
conditions

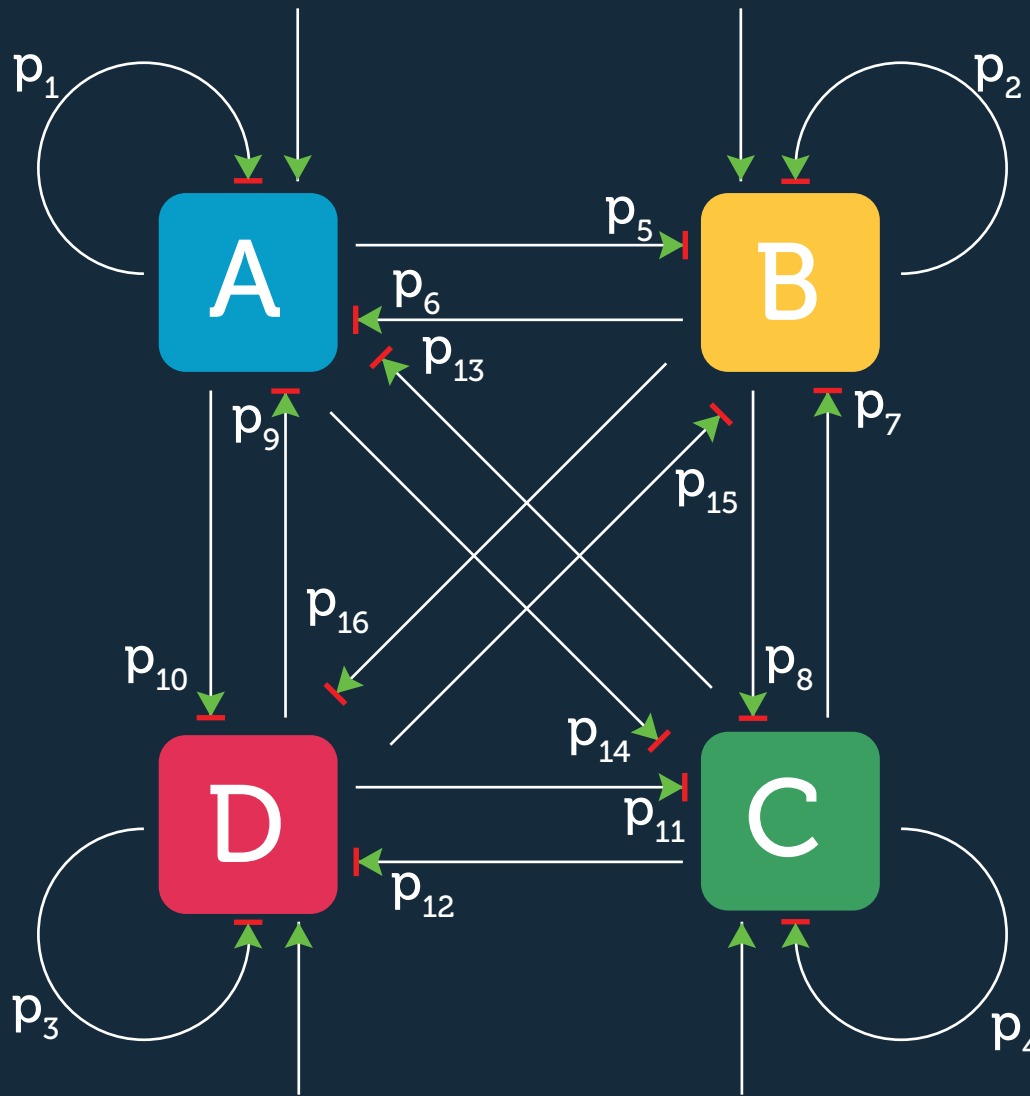


Catastrophic
event



Optimal growth
conditions

Modelling a gene regulatory network



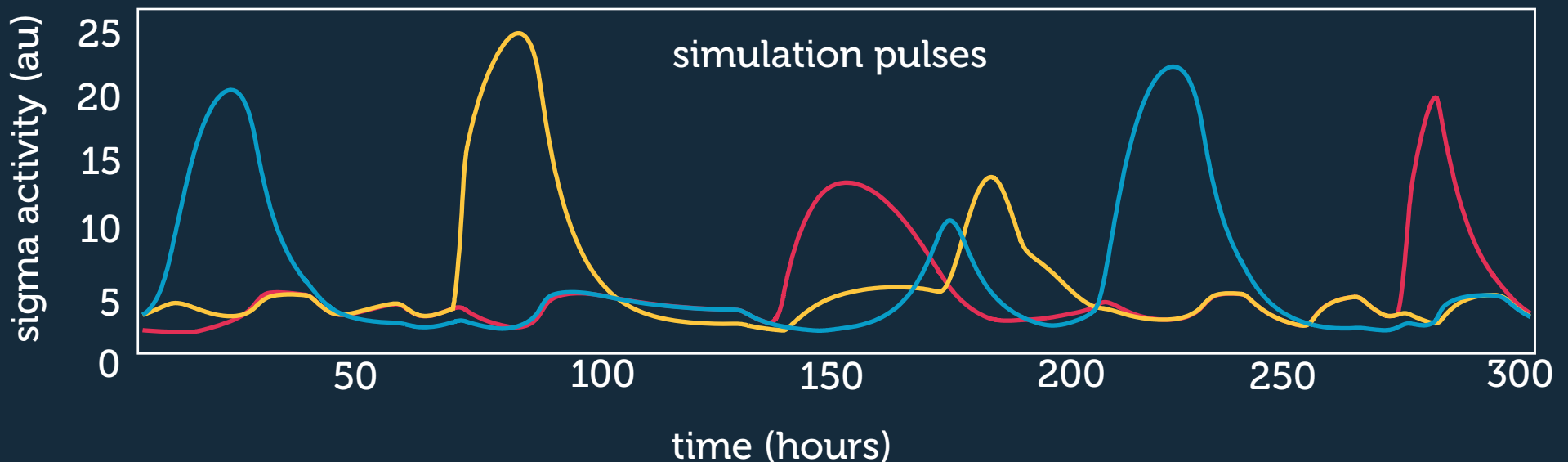
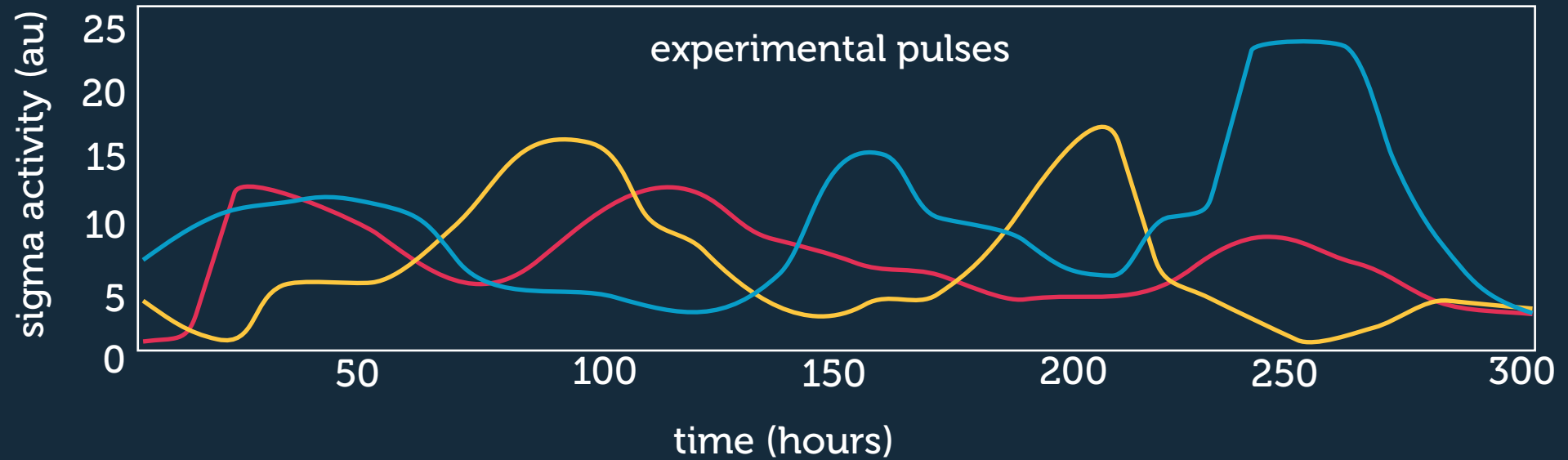
Interactions between genes define network

$$N = \{p_1 p_2 p_3 p_4 \dots p_{16}\}$$

For each path, p :

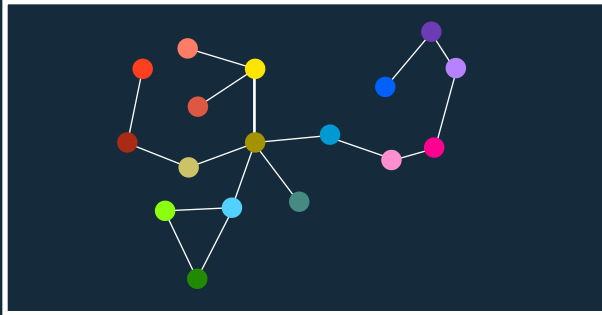
- present / absent {bool.}
- activation / repression {bool.}
- response curve {real}
- lag {real}

Simulating network interactions



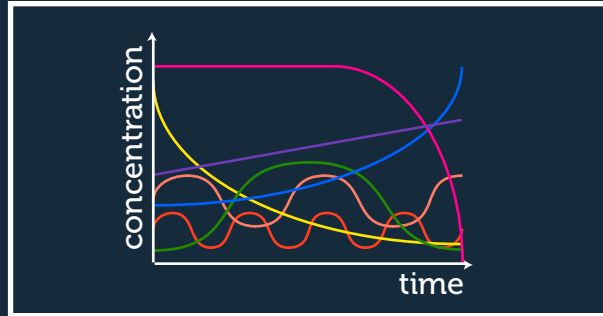
Emulating biological evolution

1. Initiation



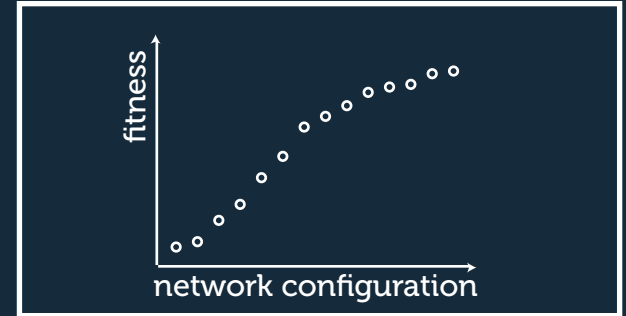
generate random networks

2. Simulation



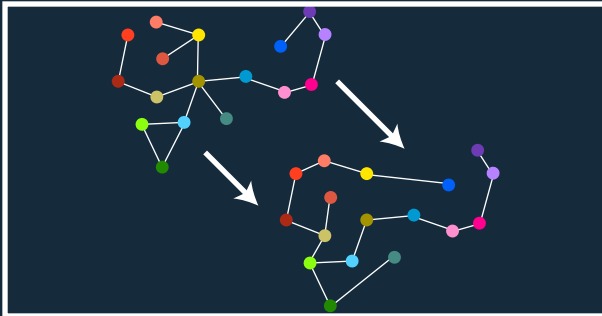
run simulations to get
timeseries of gene
concentrations

3. Fitness



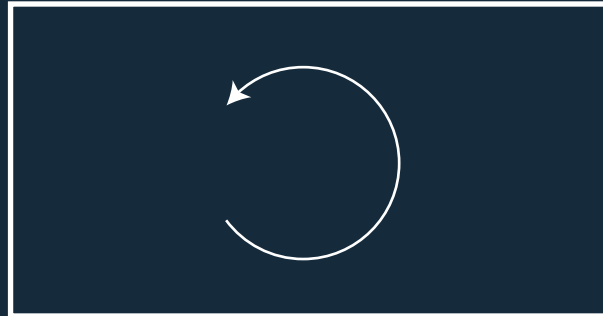
rank all networks according
to fitness of output (choose
function)

4. Reproduction



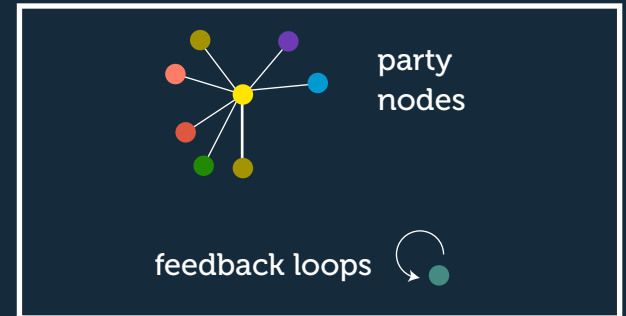
mutations, recombination
and clonal reproduction

5. Evolve



repeat steps 2 - 4 over
many generations

6. Analysis







look for common themes
in evolved networks

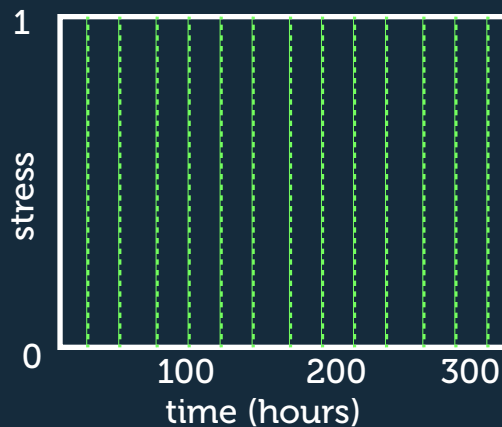
Testing for bet-hedging



population
size of
K cells
in colony
(not to scale)

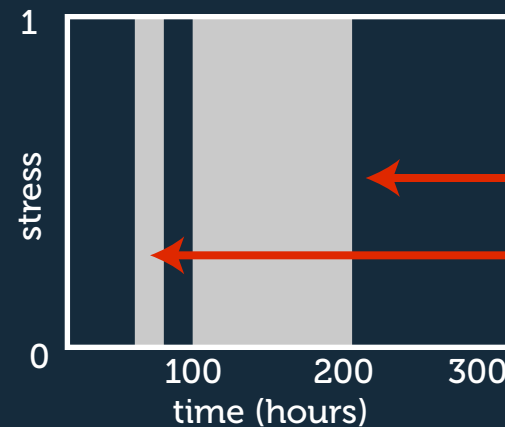
		resistance gene		
		on	off	
antibiotic	on			four possible scenarios for phenotype and environment
	off			

Cell divisions



all cells
expressing gene
1 during
cell division will
not reproduce
(but will survive)

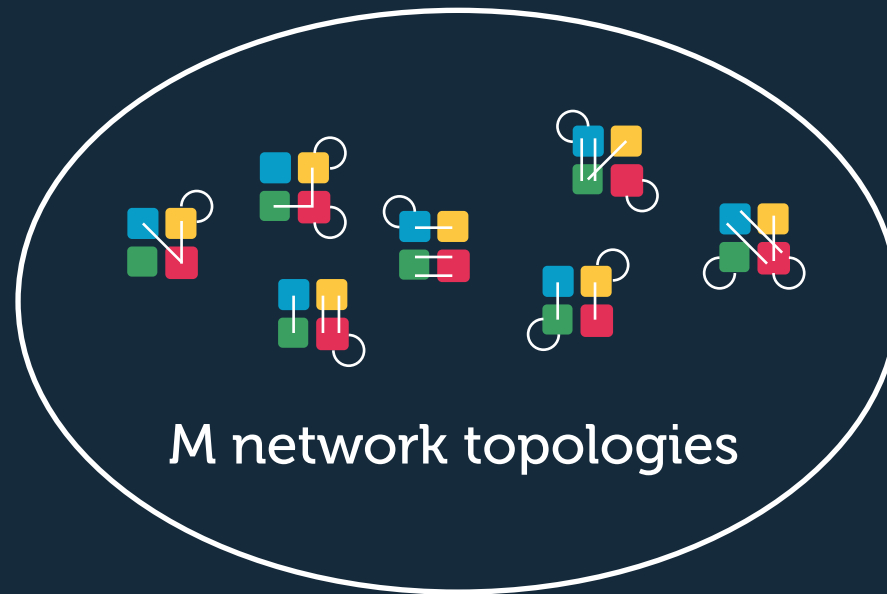
Antibiotic windows



all cells not
expressing
gene 1 in
these
windows
will die

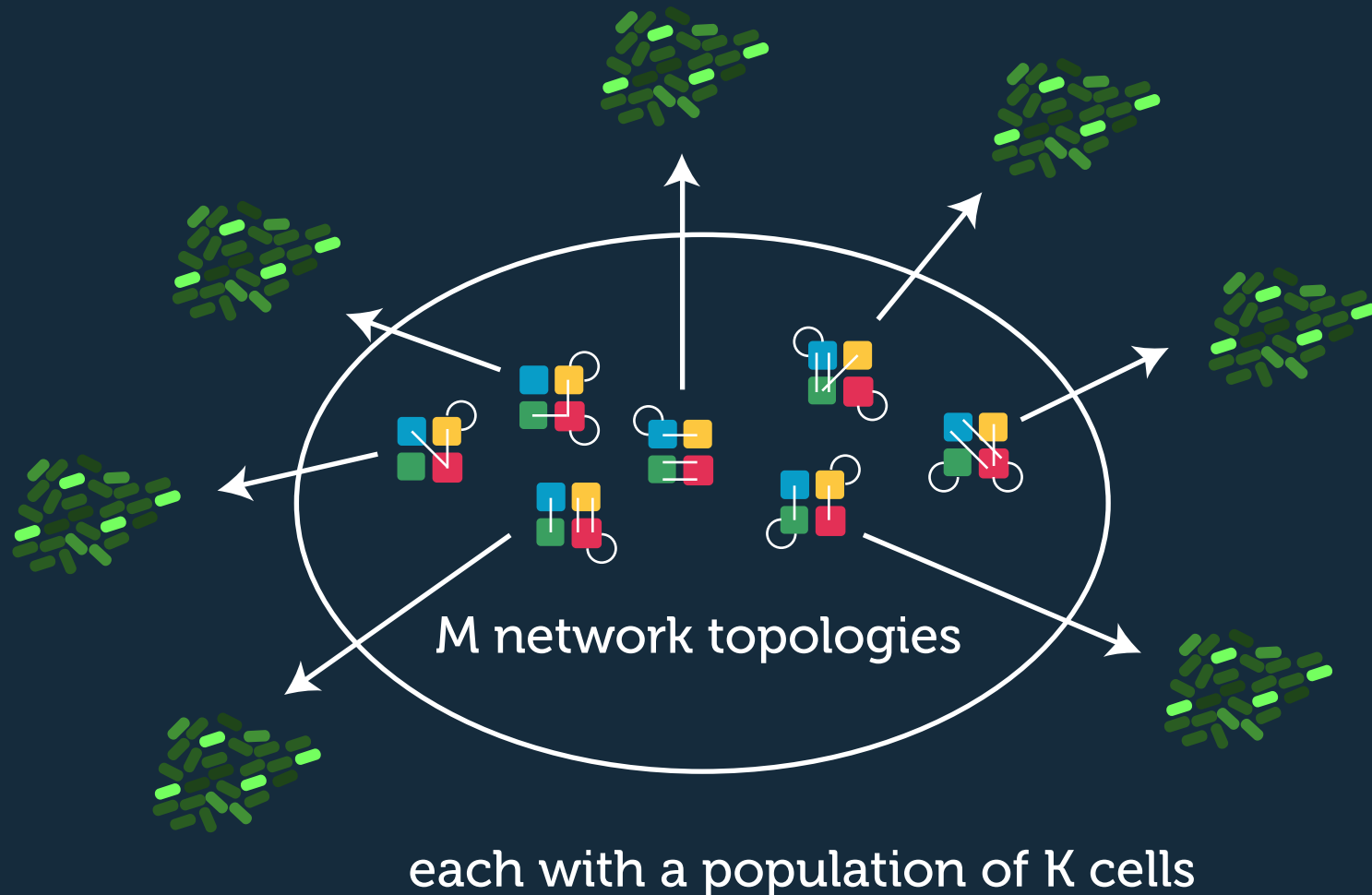
This is all very slow

For each generation:



This is all very slow

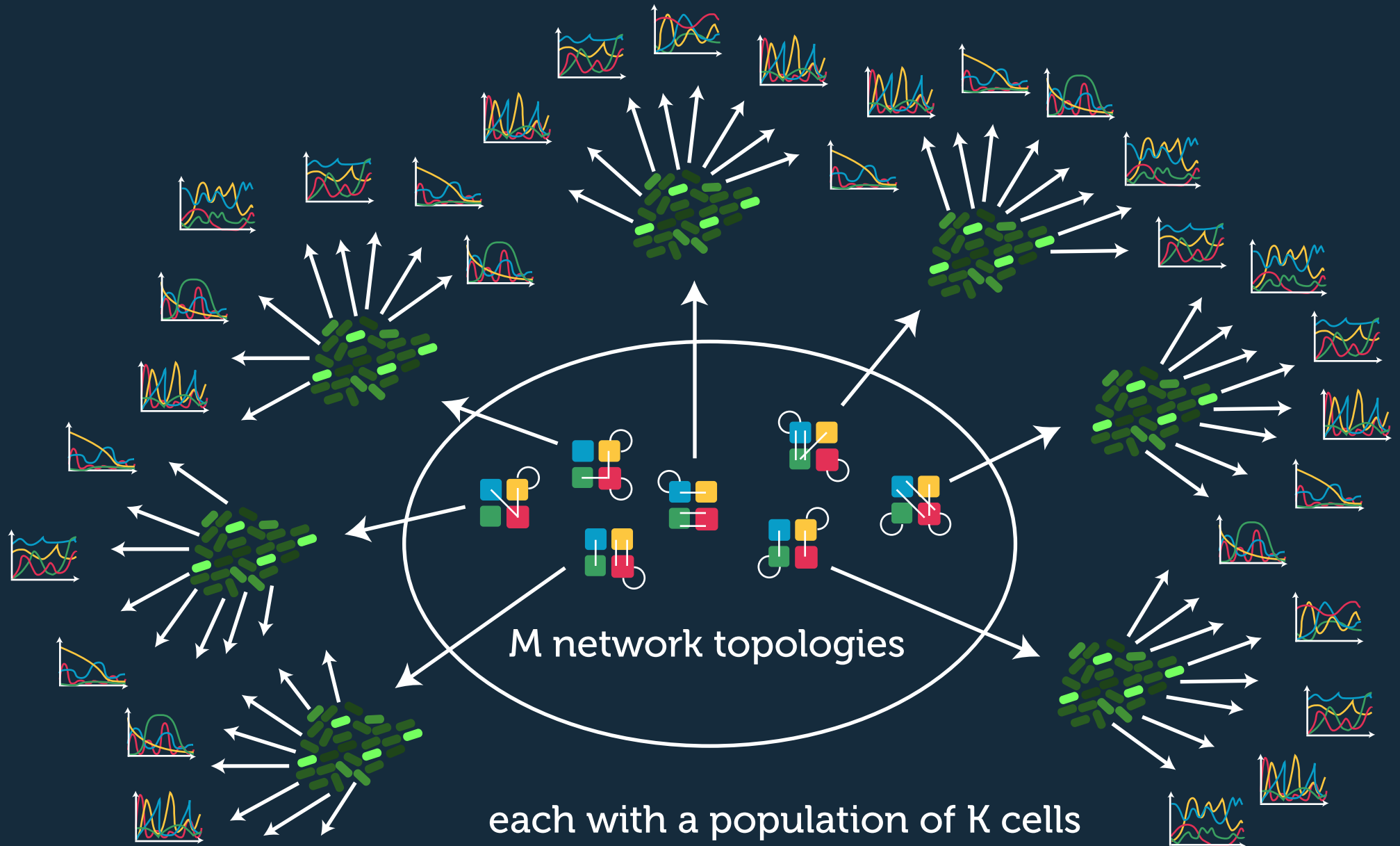
For each generation:



This is all very slow

For each generation:

each with multiple
dynamic simulations
over a few days

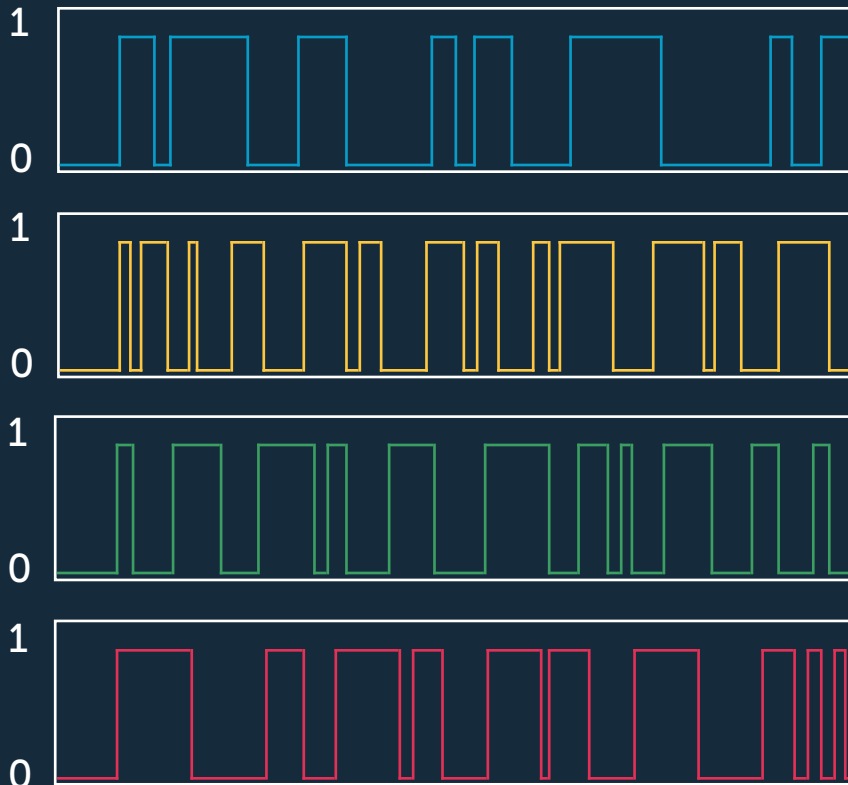


Boolean Approach



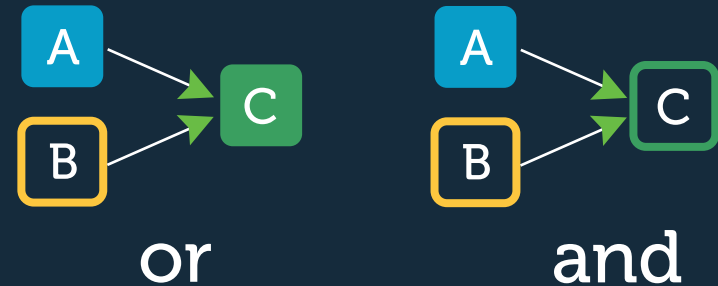
Gene concentrations

Dynamic simulation:



paths 0 or -1 or 1

logic gates



$0 < \text{lag} < 60 \text{ minutes}$

environmental signal
and response

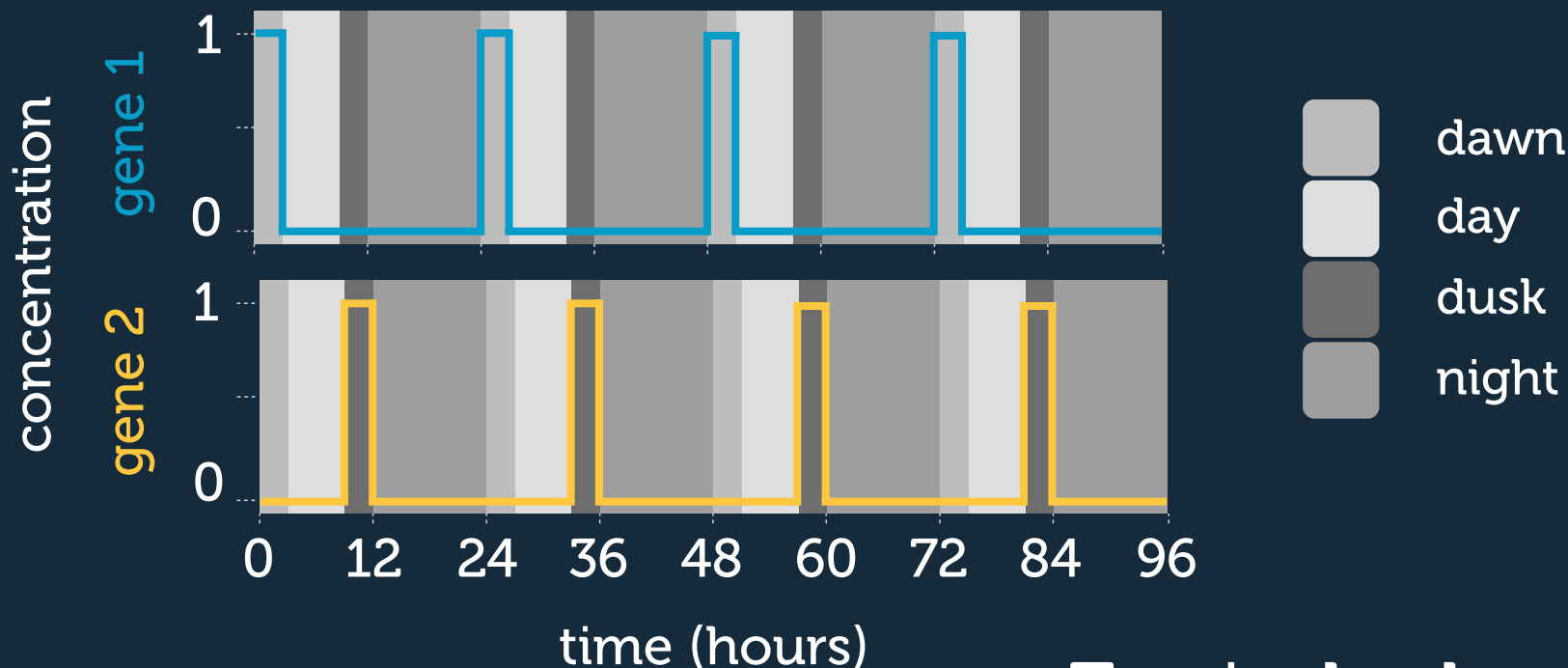


Validating the boolean approach

can we evolve a clock?

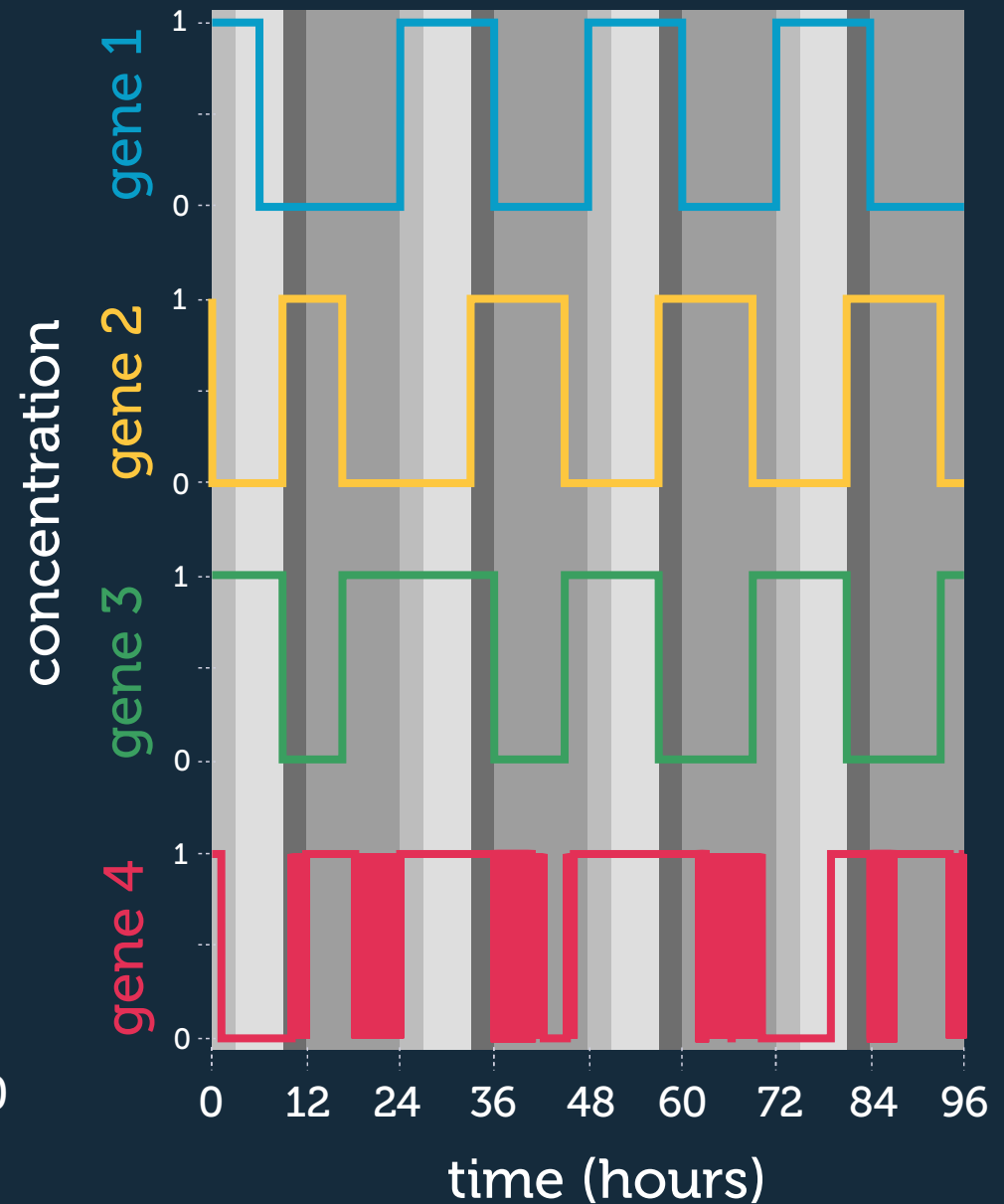
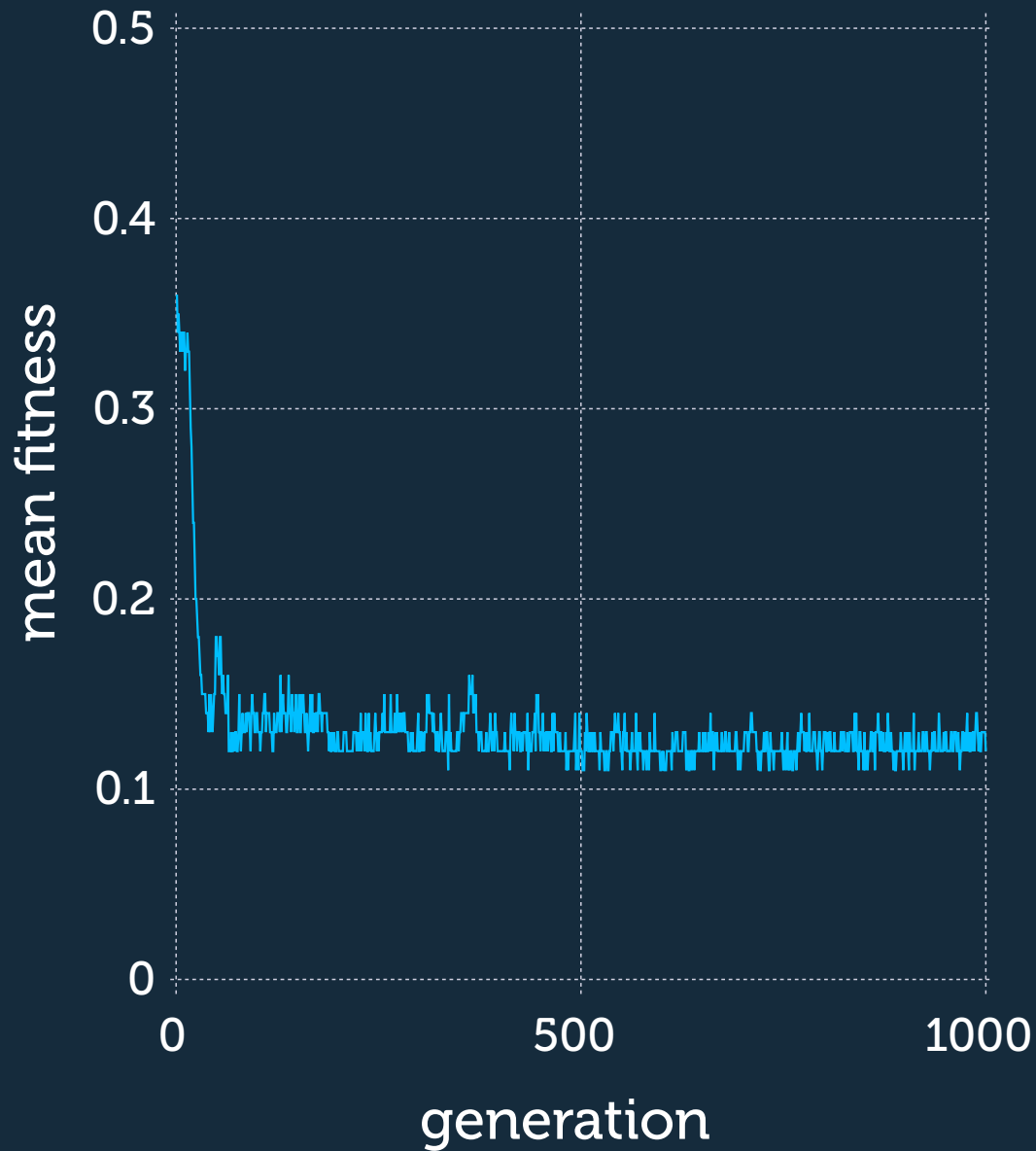


$$\sum_{\text{dawn}} G_1 + \sum_{\text{dusk}} G_2 - \sum_{\text{not dawn}} G_1 - \sum_{\text{not dusk}} G_2$$



Troein, Locke et al., 2009

Genetic algorithm



Summary and next steps

Done

- Constructed a generic framework for network evolution
- Established novel boolean network evolution system
- Written GA and fitness functions
- Evolved a clock (sort of)

Doing

- Try starting with networks that are already clocks
- Try starting with networks that evolved in the clock evolution paper

To do

- Change the environmental input pattern
- Look at the interplay between sensory and anticipatory pathways

Acknowledgements



James Locke

Carl Troein

Locke Lab

Thanks for listening

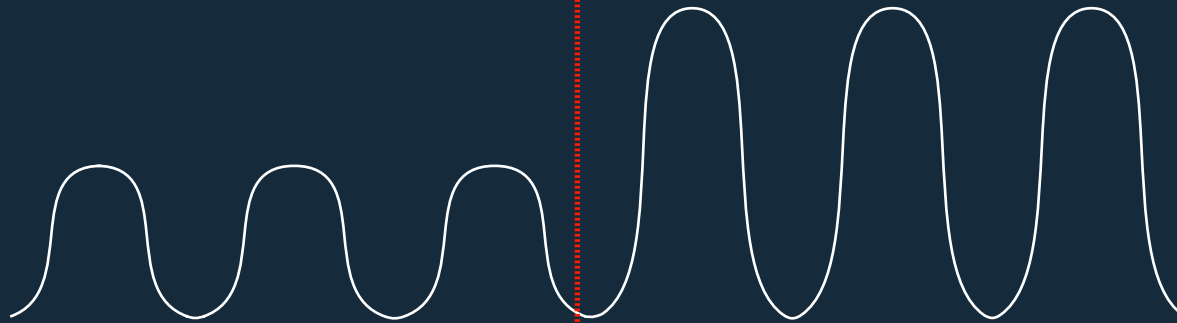
Bet-hedging requires whole colony



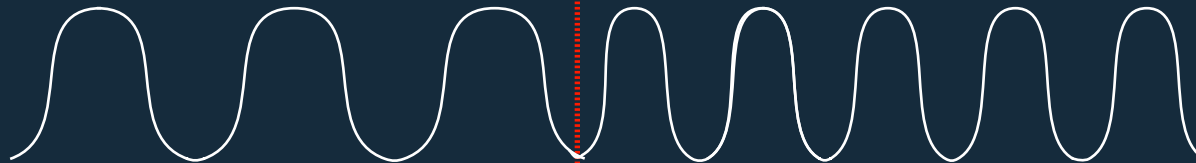
Multiple stochastic simulations for every generation (slow)

Pulse modulation

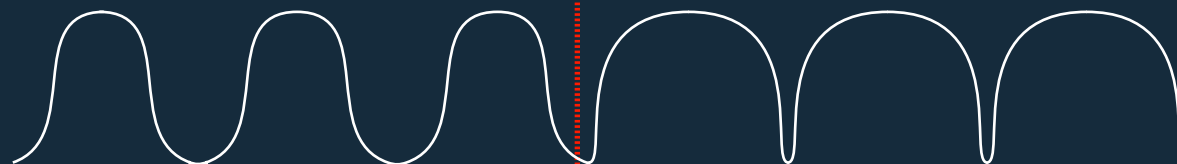
Amplitude modulated (AM)



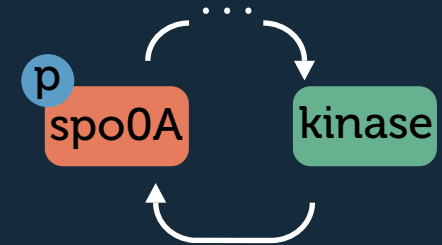
Frequency modulated (FM)



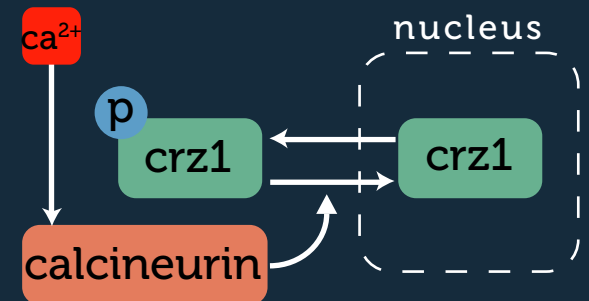
Duration modulated (DM)



Bacillus subtilis spo0A



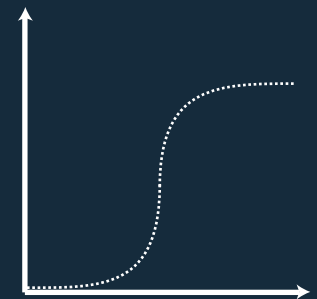
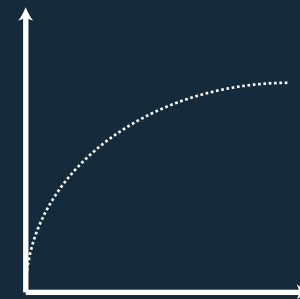
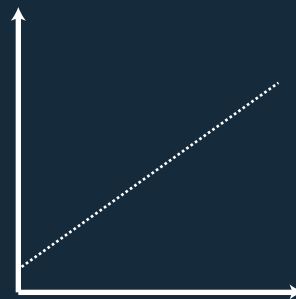
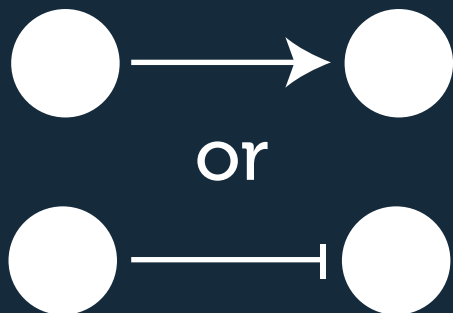
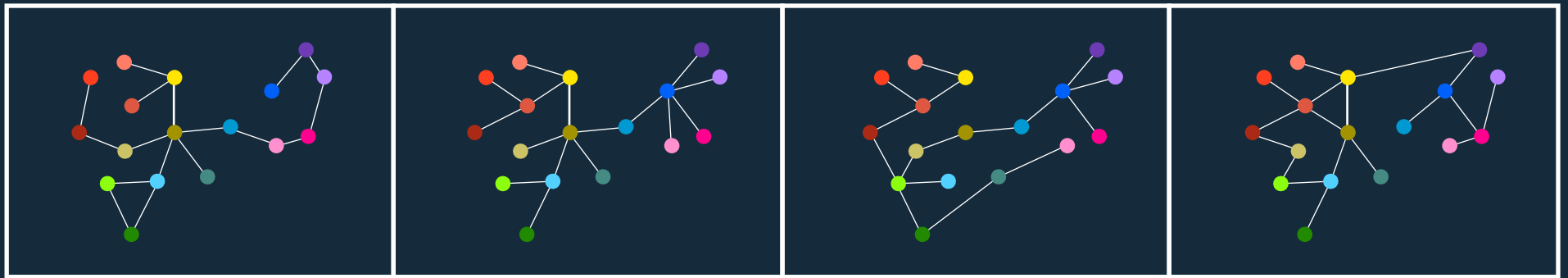
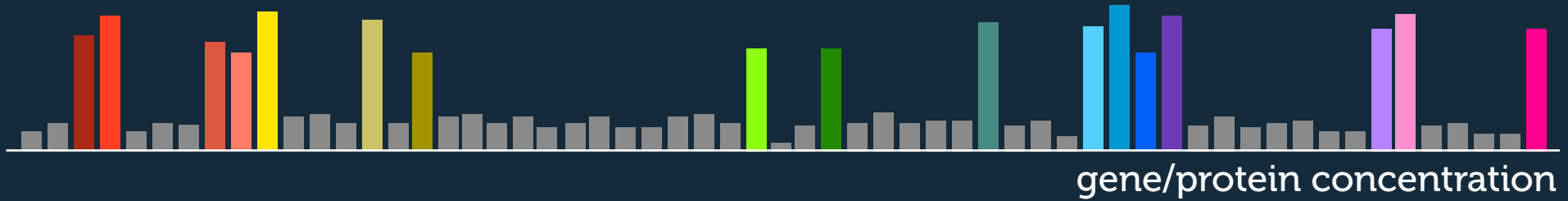
Yeast Crz1



?

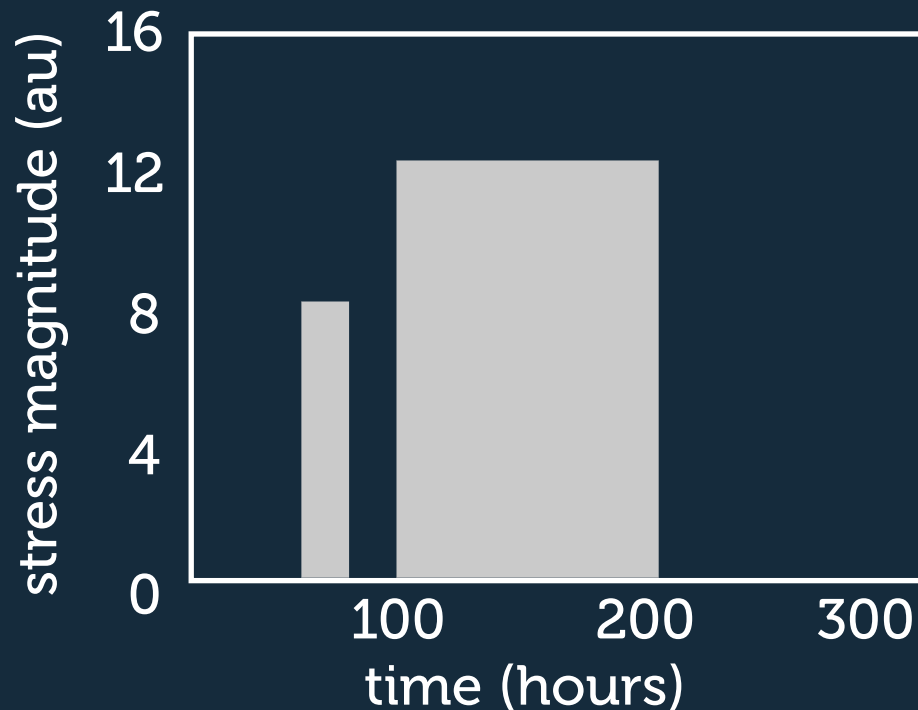
Approach

inferring networks from data



response curves

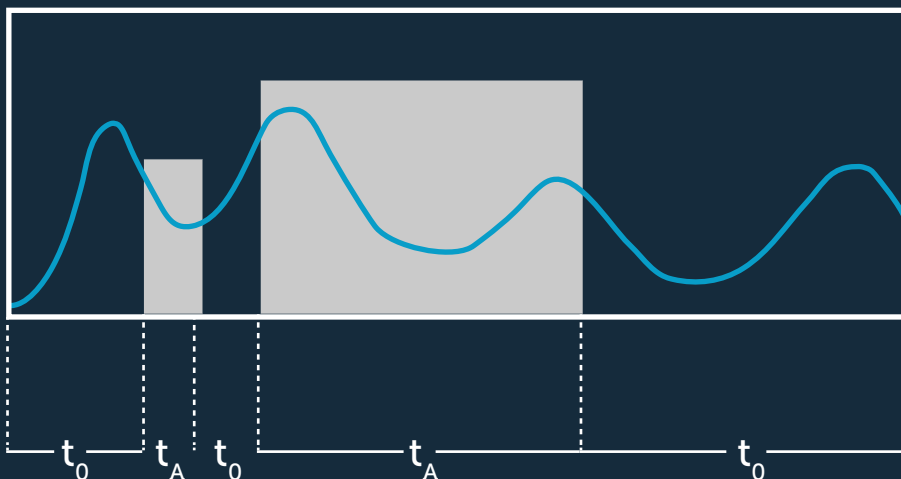
Testing for bet-hedging



A antibiotic resistance gene

(a) severe cost for no/low expression during stress

(b) slight cost for expression otherwise (slowed growth)



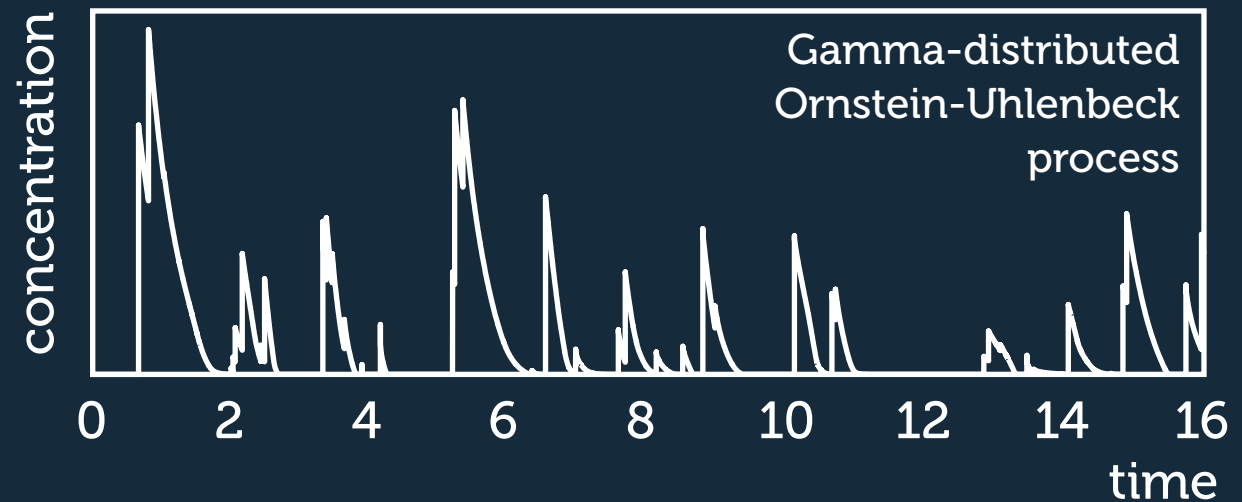
Fitness will be proportional to:

(a) $\int_{t_A} G_1$

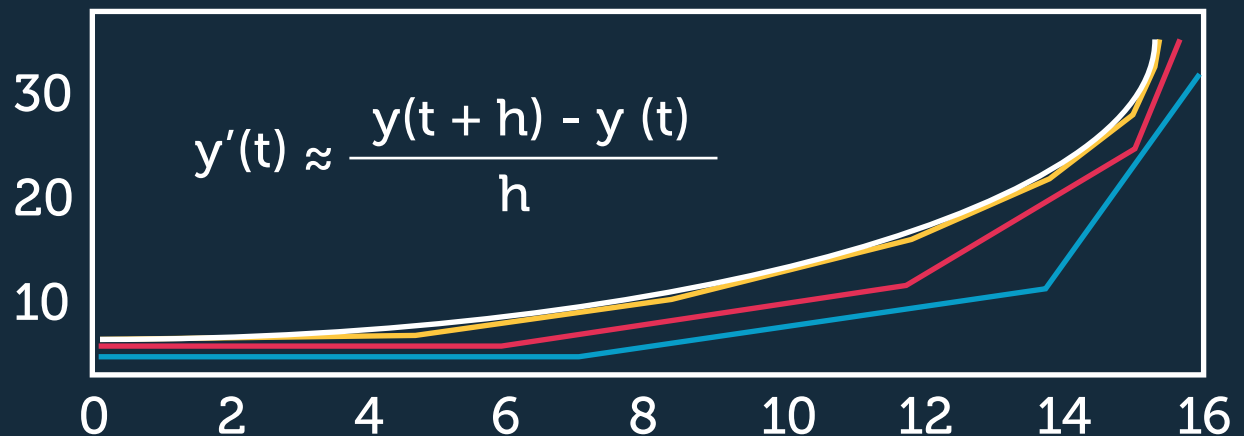
(b) $-\int_{t_0} G_1$

Bet-hedging requires stochasticity

Option 1:
Discretise and
solve piecewise



Option 2:
Create an
SDDE solver



Option 3:
Simulate transcription and translation using SDE solver with no lag.