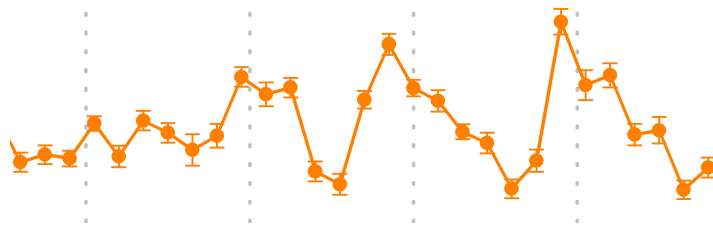
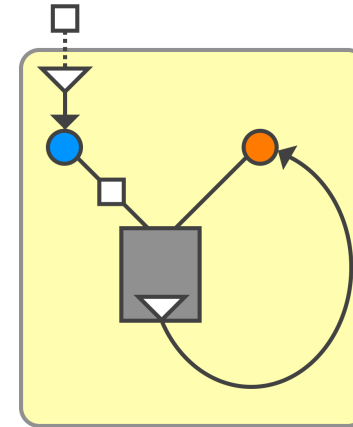


The watchmaker's apprentice:

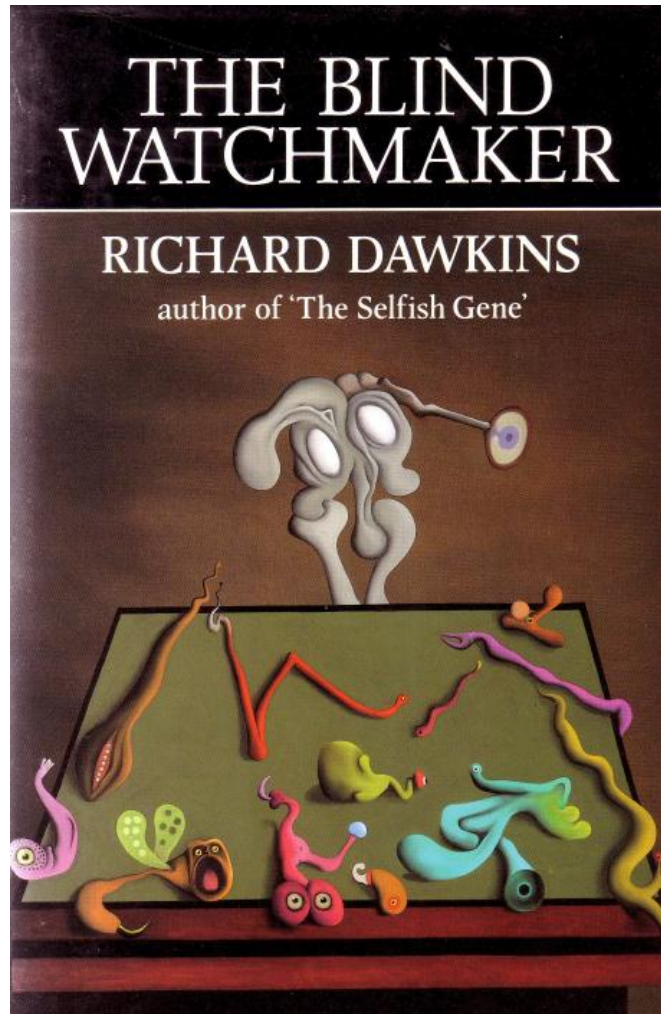
Building a synthetic genetic oscillator with parts borrowed from nature



Mukund Thattai
Simons Centre for the Study of Living Machines
NCBS/TIFR

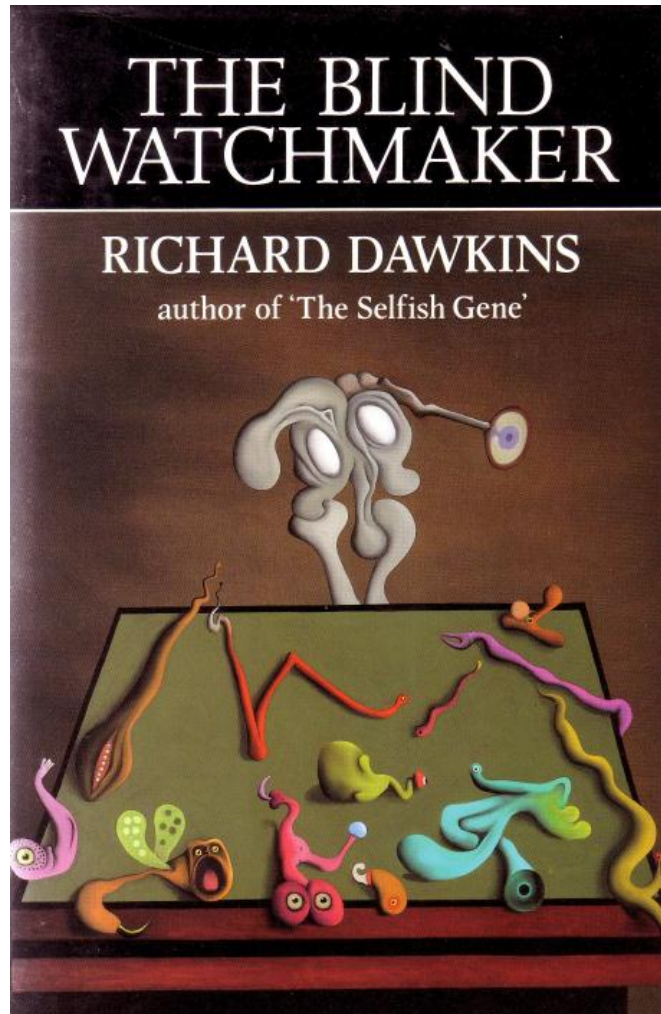
September 2013

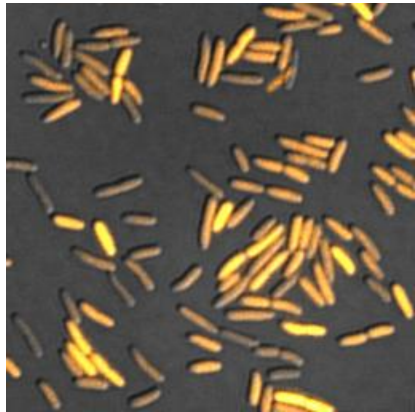
The blind watchmaker
generates complexity through
the undirected processes of
mutation and selection...



The blind watchmaker generates complexity through the undirected processes of mutation and selection...

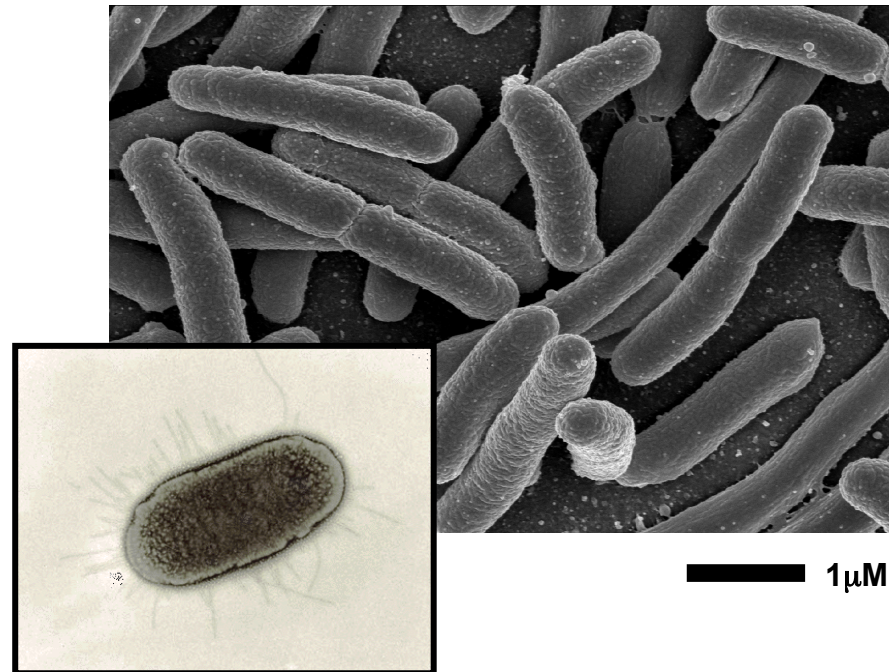
... while the watchmaker's apprentice has a design in mind, and uses the complex parts invented by nature to realize it





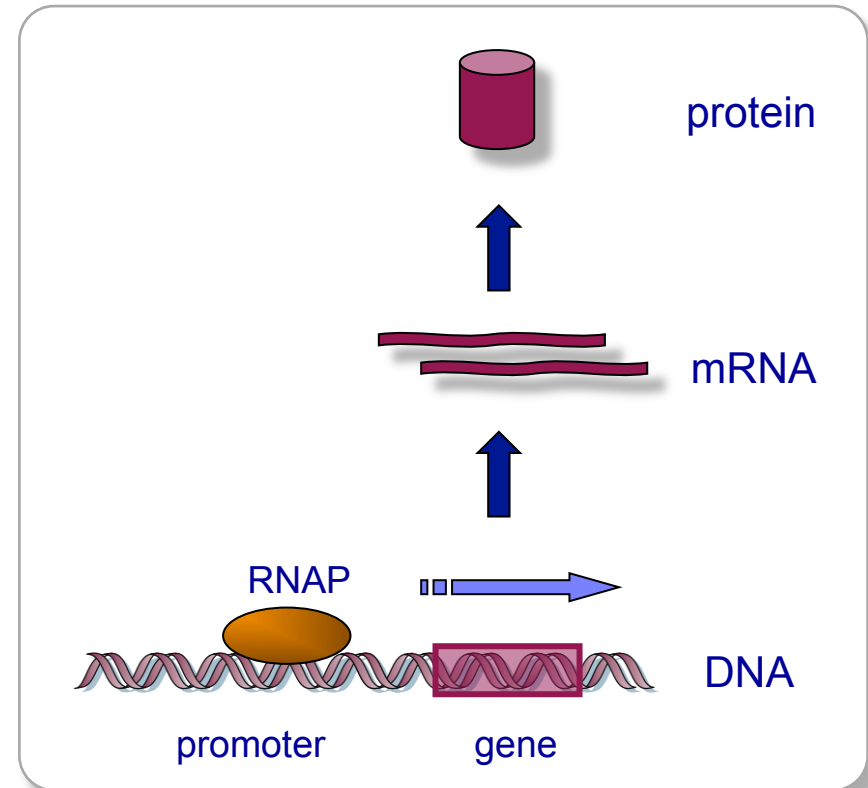
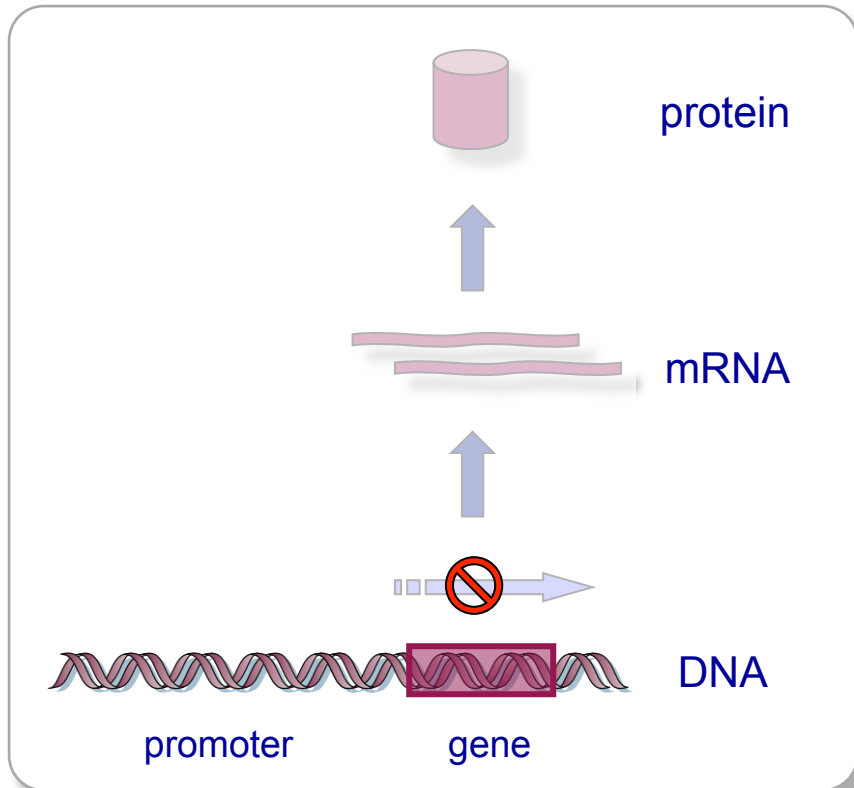
A brief introduction to genetic networks

The *E. coli* genome contains about 4000 genes...

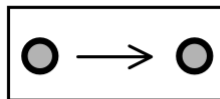
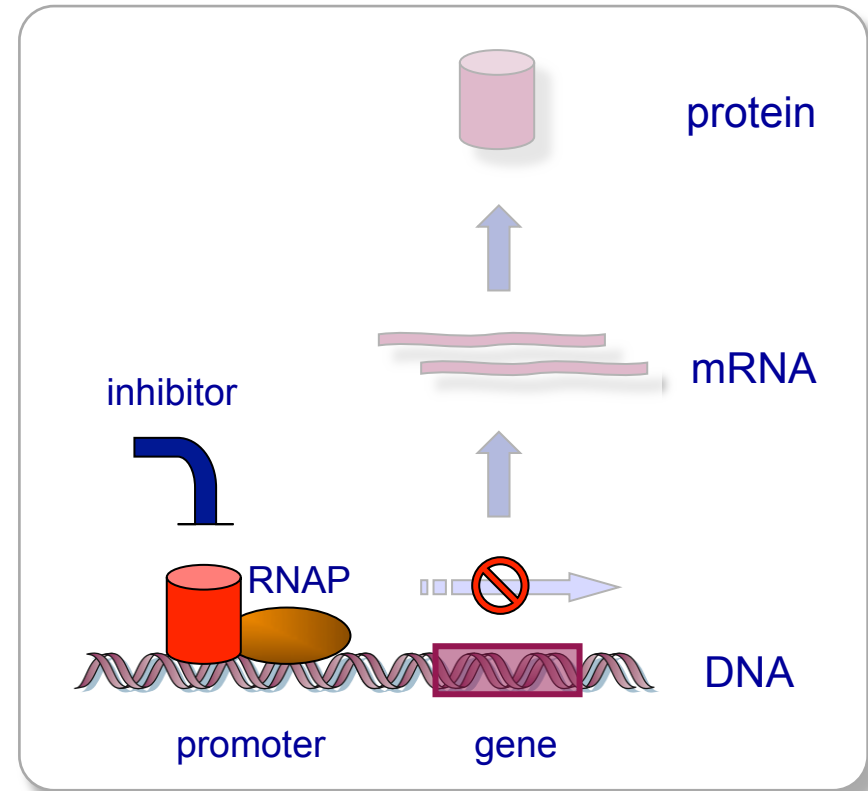
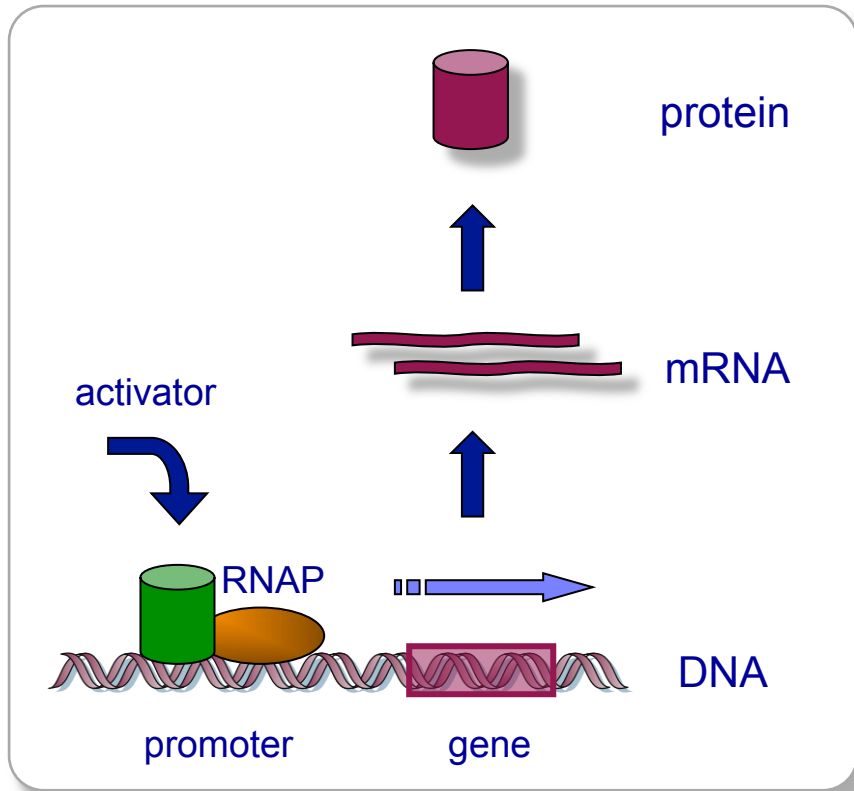


... and the human genome contains about 24000 genes

Genes can be turned on and off, allowing combinatorial complexity

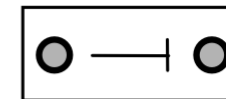


Genes can be turned on and off, allowing combinatorial complexity



0 → 0

1 → 1



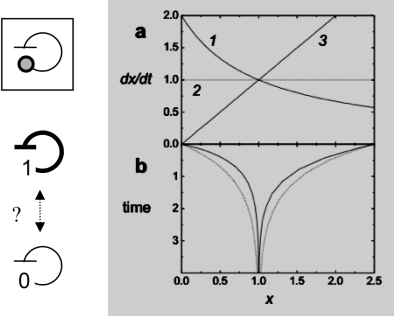
0 → 1

1 → 0

Genes can be connected into networks with useful dynamical properties

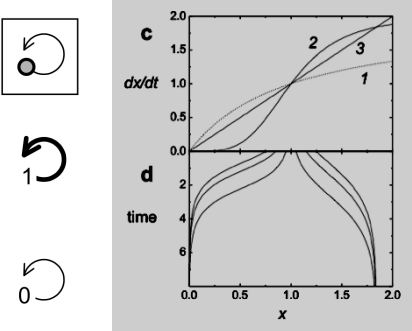
Noise reduction

Negative feedback

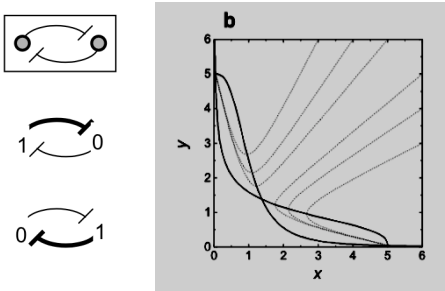


Memory

Positive feedback

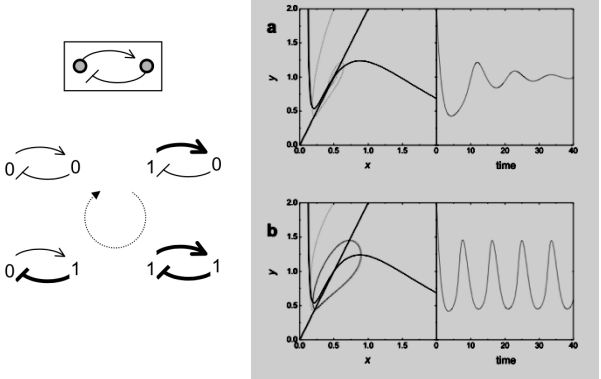


Flip-flop

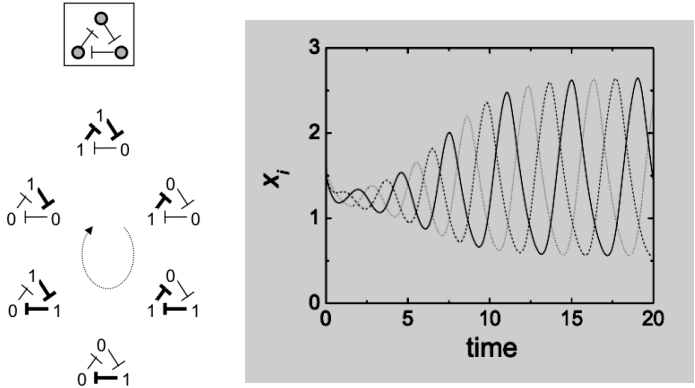


Oscillations

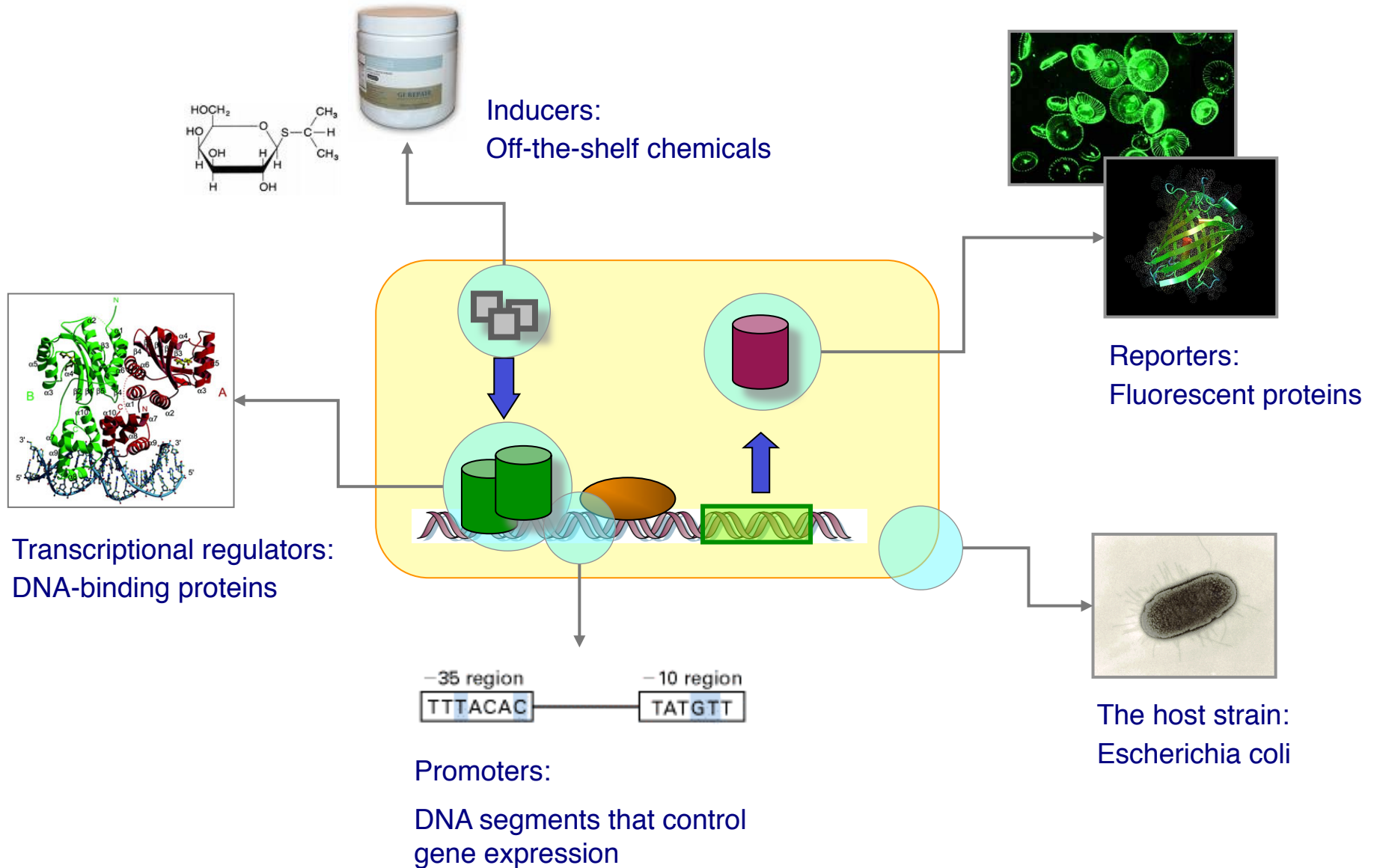
Hysteretic oscillator



Ring oscillator

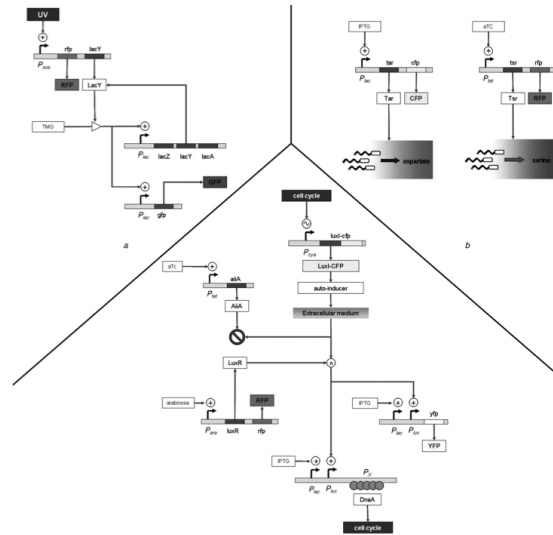


We can build networks from basic parts...



... but getting things to work is hard!

In principle, UV exposure would trigger transient expression of lacY, and the resulting TMG uptake would turn on lac expression. In practice, the design did not function as expected. We soon realised that the presence of extraneous lacI binding sites on the plasmid backbone had perturbed the system.

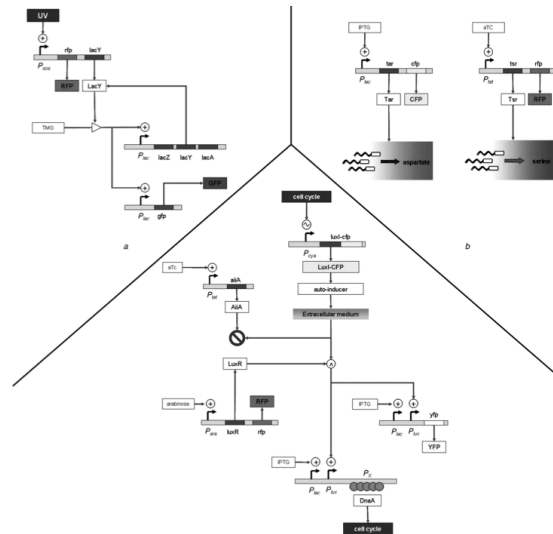


However, implementing such a gradient turned out to be far from trivial. Moreover, we were not able to rescue chemotaxis in the knockout strain.

As expected, the RFP control signal did not oscillate. However, synchronisation could not be achieved because the presence of DnaA boxes did not seem to affect cell-cycle progression.

... but getting things to work is hard!

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METHODOLOGY

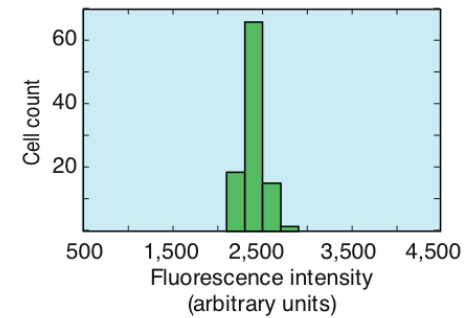
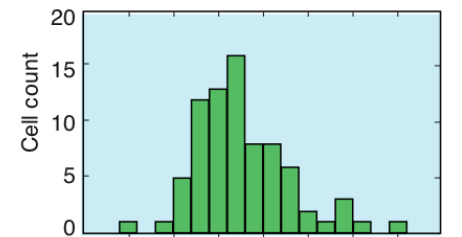
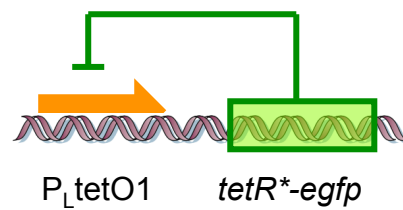
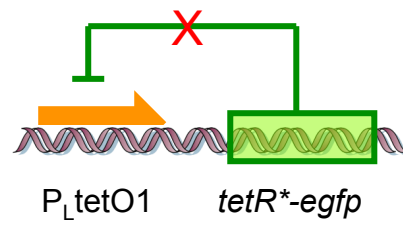
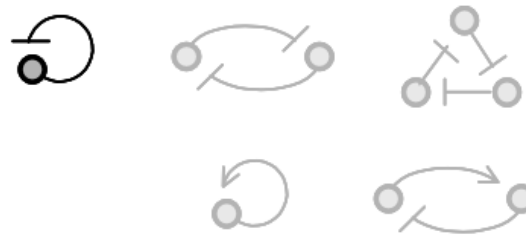
Introduction of customized inserts for streamlined assembly and optimization of BioBrick synthetic genetic circuits

Julie E Norville^{1,2,3*}, Ratmir Dersa^{4,5,6}, Saurabh Gupta², Kelly A Drinkwater^{1,2}, Angela M Belcher^{2,3}, Andres E Leschziner⁷, Thomas F Knight Jr^{1,8}

BioBrick standard biological parts are freely available to researchers through the Registry of Standard Biological Parts [42]. Although BioBricks have been used to construct a large variety of genetic circuits [12,43-54], these circuits often require optimization [44,55-59] and currently, there is no standard methodology for optimizing BioBrick circuits.

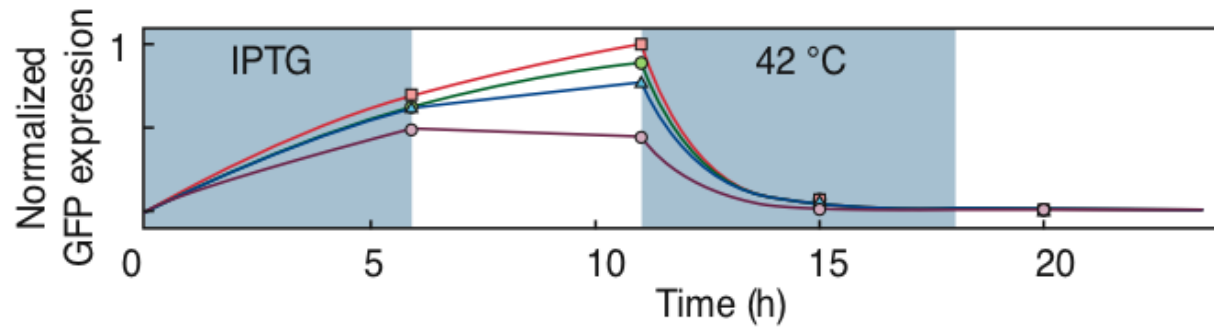
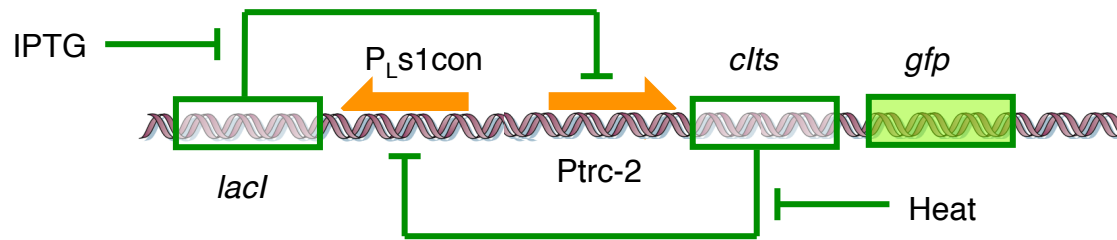
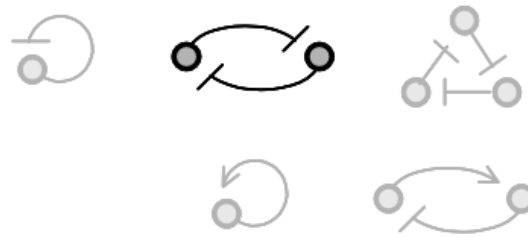
Negative feedback

Noise reduction



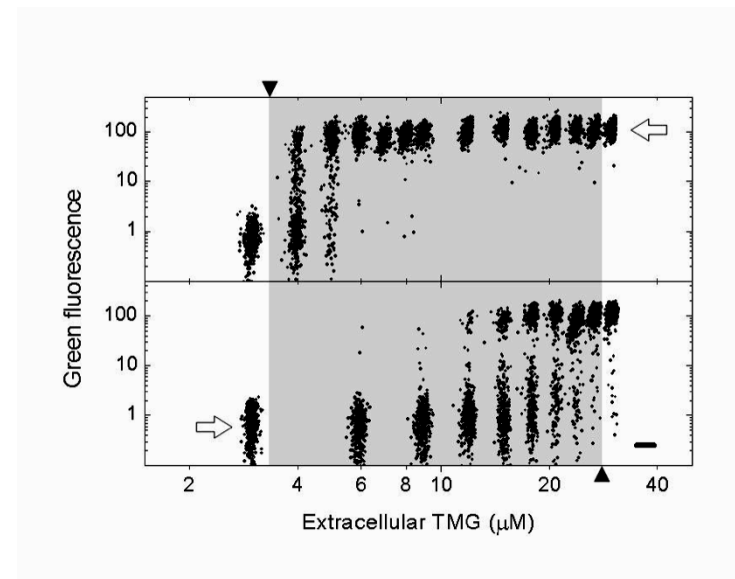
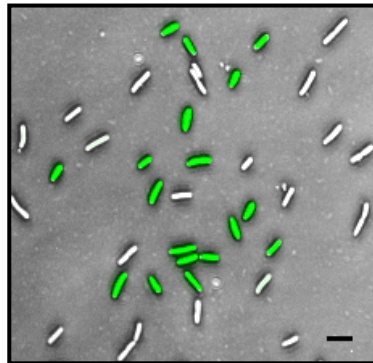
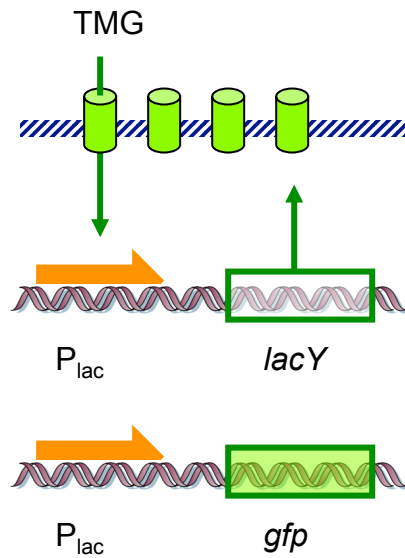
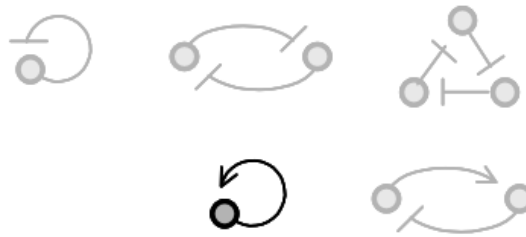
Flip flop

Memory



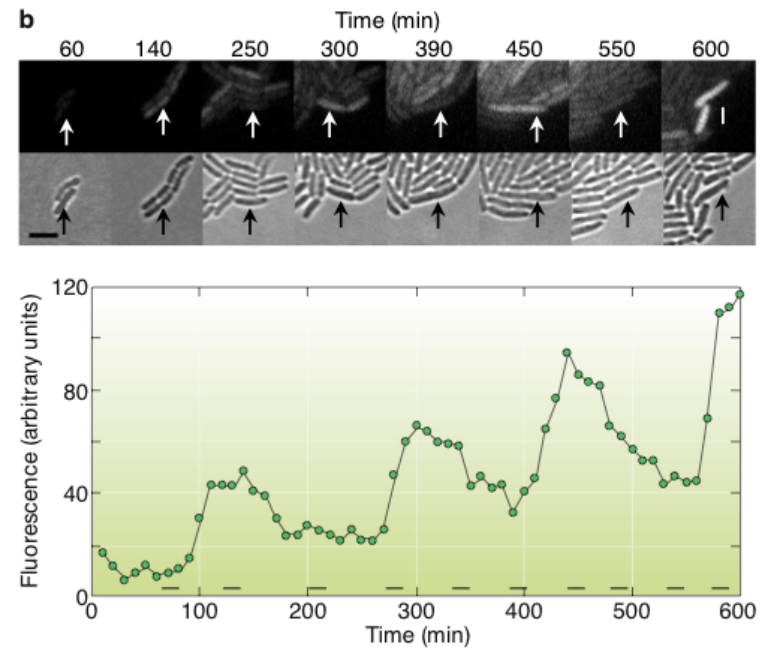
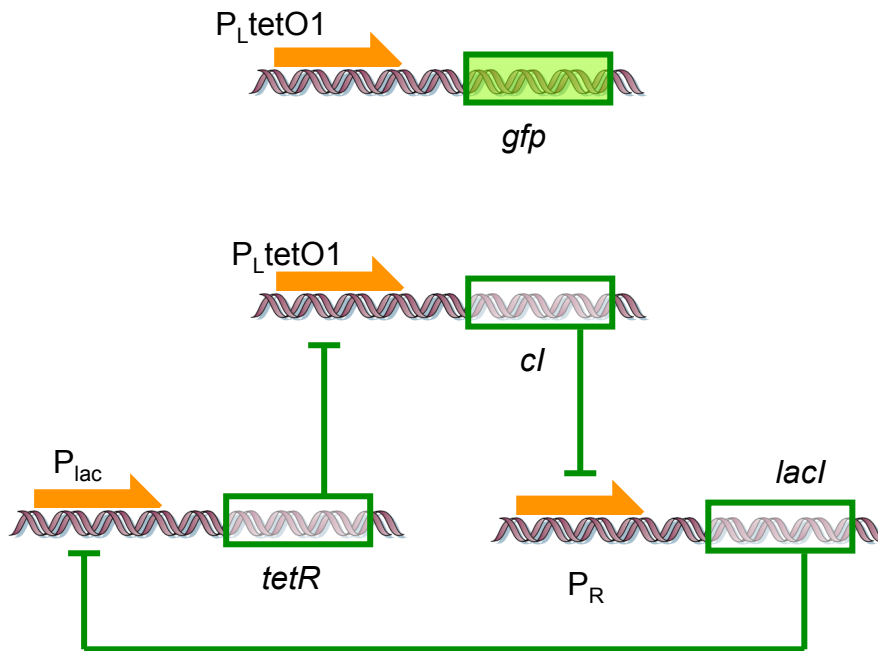
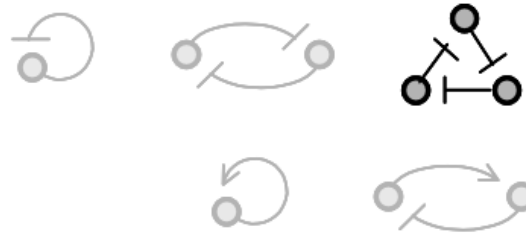
Positive feedback

Memory



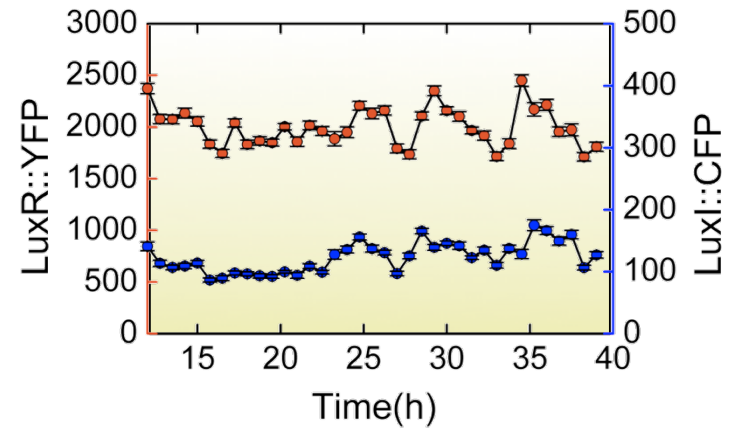
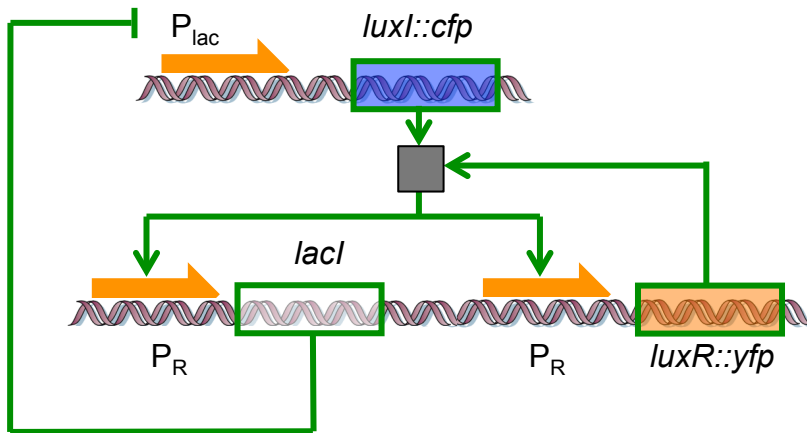
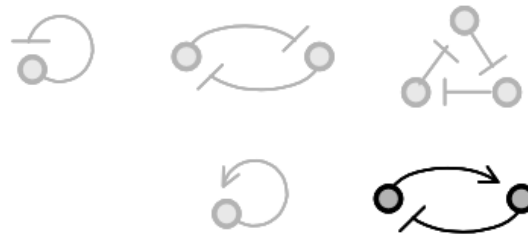
Ring oscillator

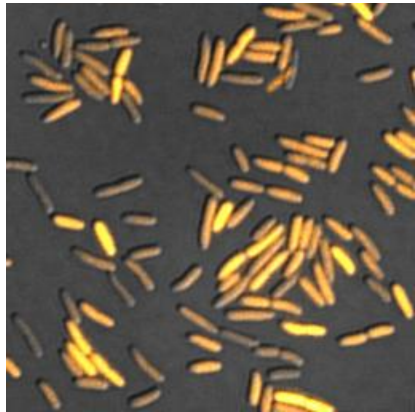
Oscillations



Hysteretic oscillator

Synchronized oscillations





The basic parts:
Borrowing from a cell
communication system

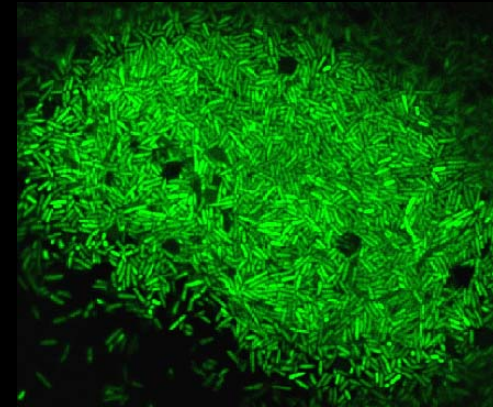
Bacteria can change their behaviour depending on local cell density



Density-dependent bioluminescence in *Vibrio harveyi*



Symbiosis between bioluminescent *Vibrio fischeri* and the hawaiian bobtail squid



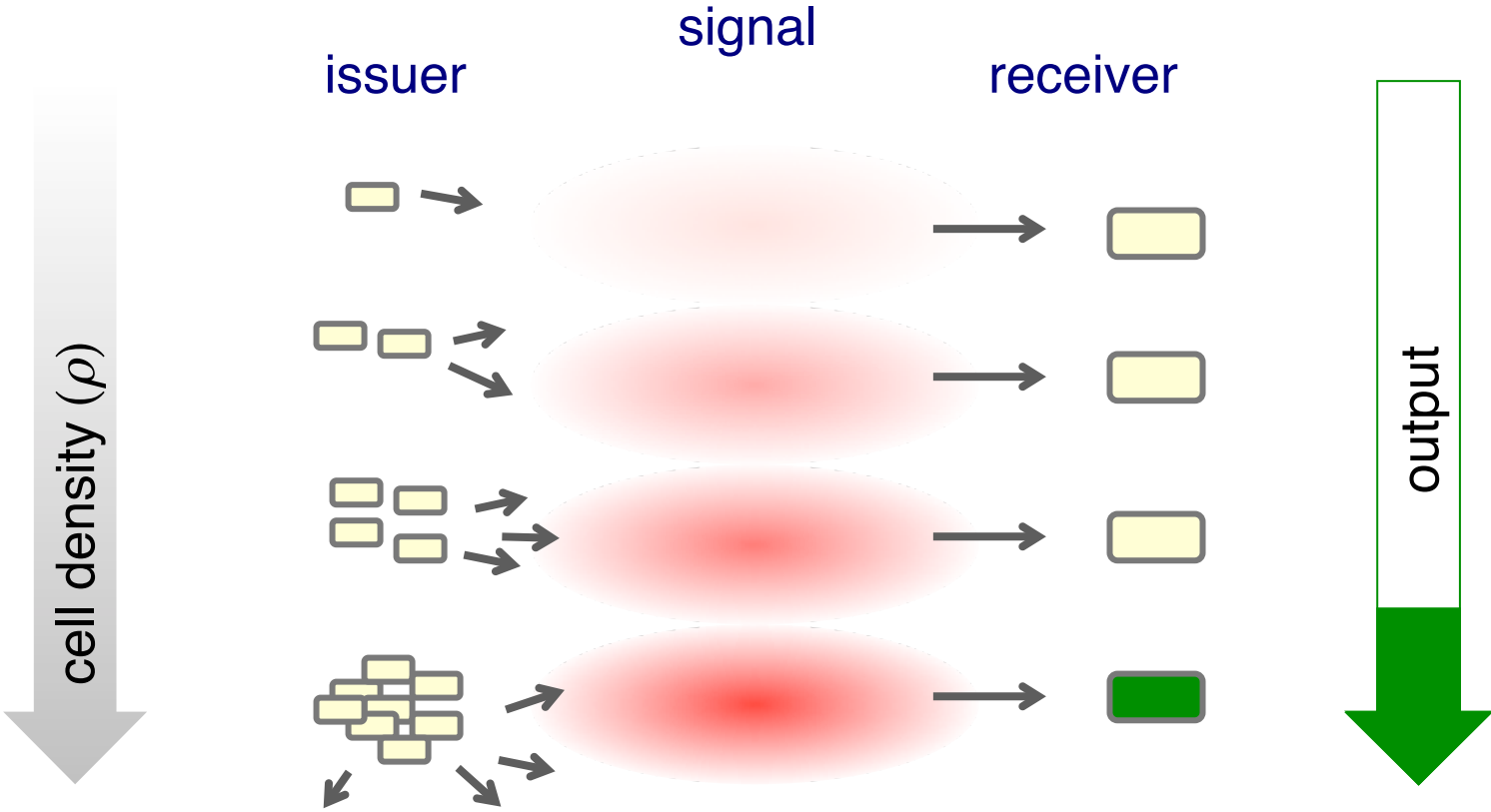
Quorum sensing during biofilm formation in *Pseudomonas aeruginosa*

📖 MJ McFall-Ngai & EG Ruby, University of Hawaii

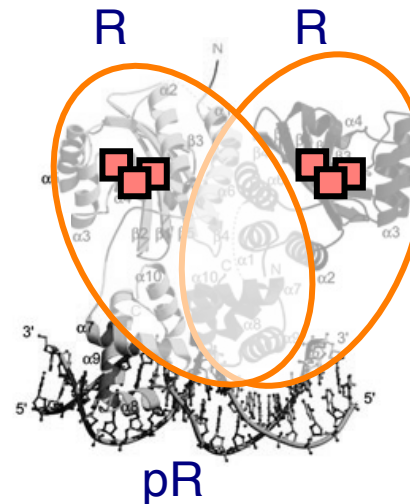
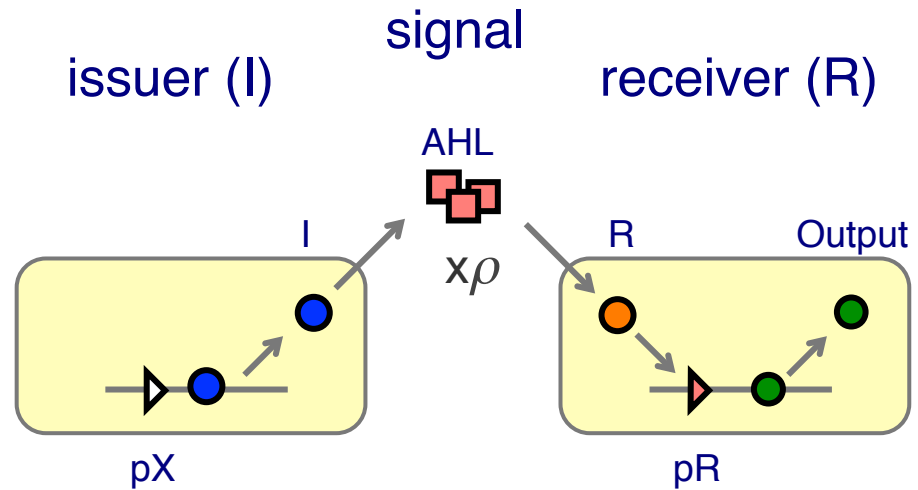
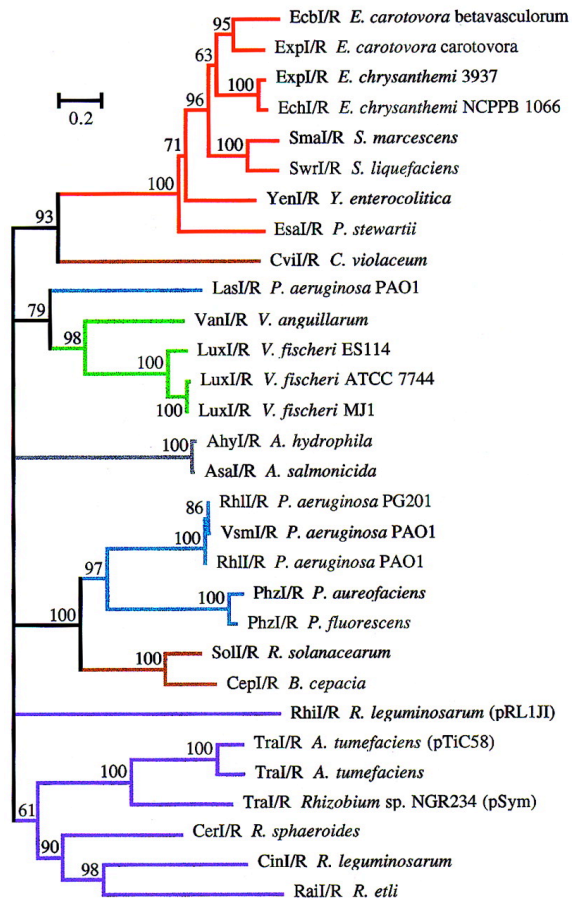
📖 F Ausubel, MGH

📖 MB Miller & BL Bassler, 2001

“Quorum sensing” is a form of chemical cell-to-cell communication

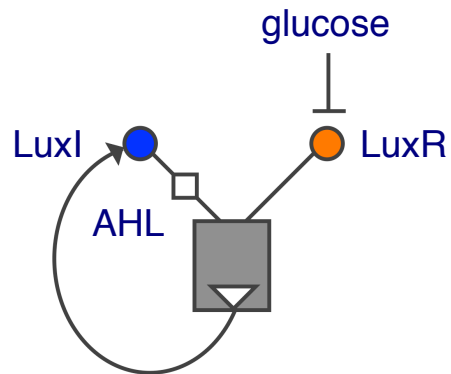


Quorum-sensing systems are built from a handful of conserved molecular components

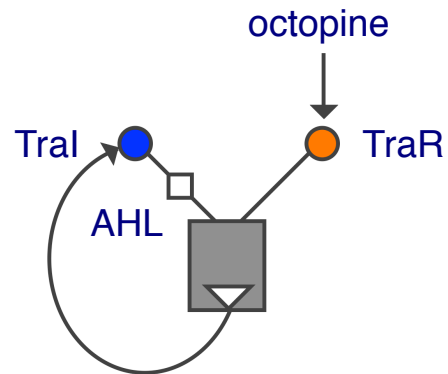


A topological puzzle: All characterized natural QS systems use I-feedback

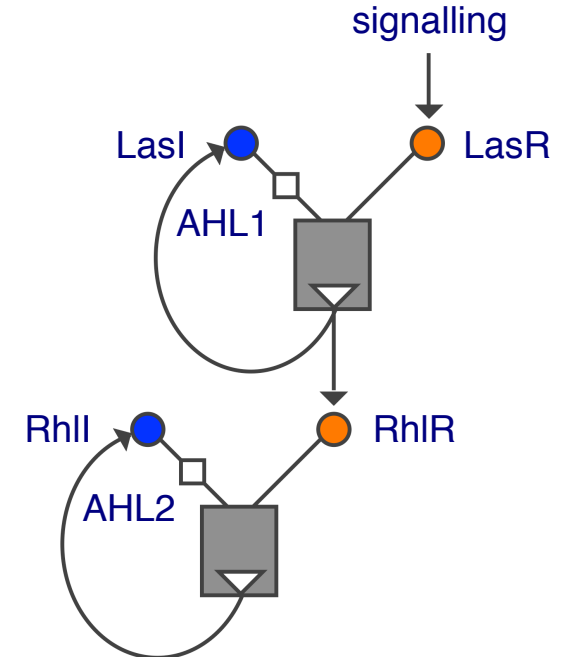
Species: *Vibrio fischeri*
Niche: marine endosymbiont
Process: bioluminescence

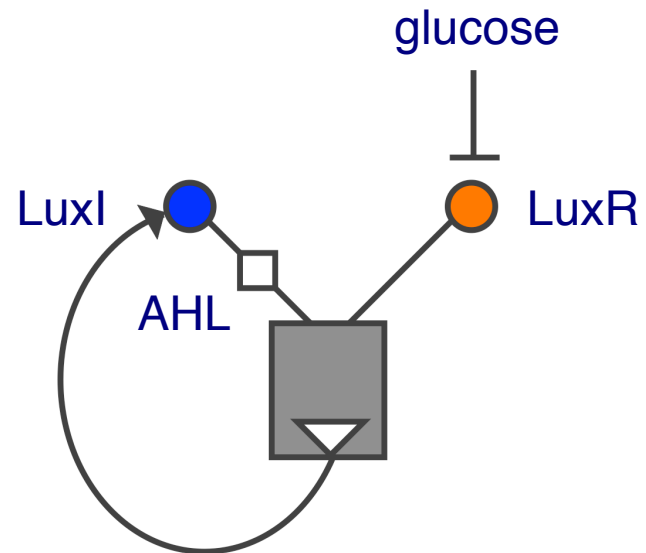


Species: *Agrobacterium tumefaciens*
Niche: plant pathogen
Process: plasmid conjugation



Species: *Pseudomonas aeruginosa*
Niche: human pathogen
Process: virulence; biofilm formation



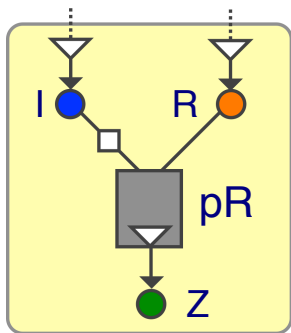


What is happening inside the biochemical black box?

What is the role of transcriptional feedback?

Why is one feedback topology preferred over another?

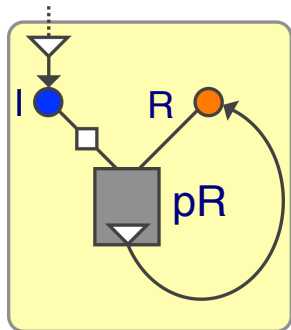
Inferring feedback behavior from feedforward data



Feedforward

$$\frac{1}{\gamma_Z} \frac{dY_Z}{dt} = Q_Z f(\mu \rho \bar{Y}_I, \bar{Y}_R) - Y_Z$$

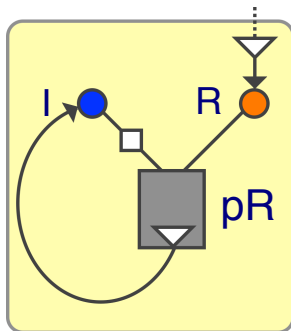
extract biochemistry
(two regulated inputs)



R-feedback

$$\frac{1}{\gamma_R} \frac{dY_R}{dt} = Q_R f(\mu \rho \bar{Y}_I, Y_R) - Y_R$$

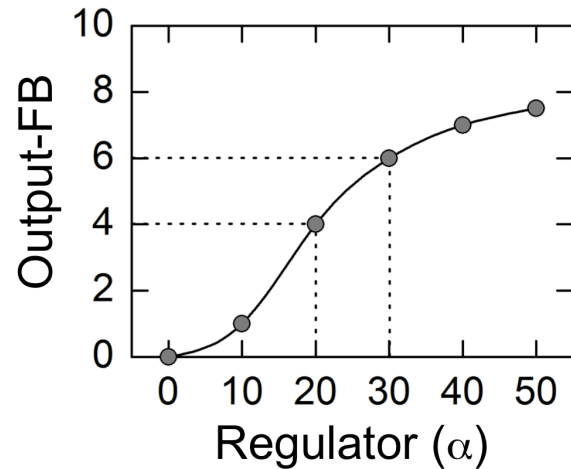
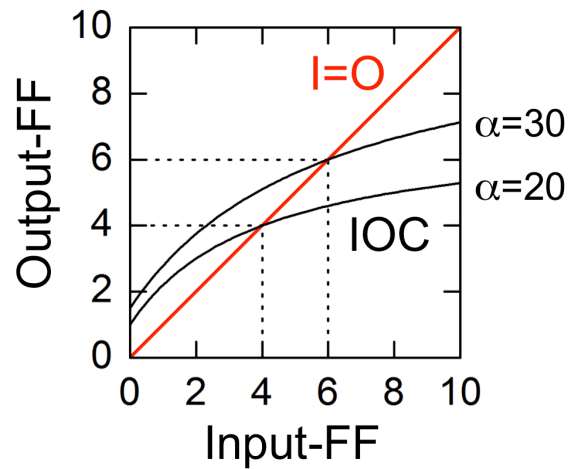
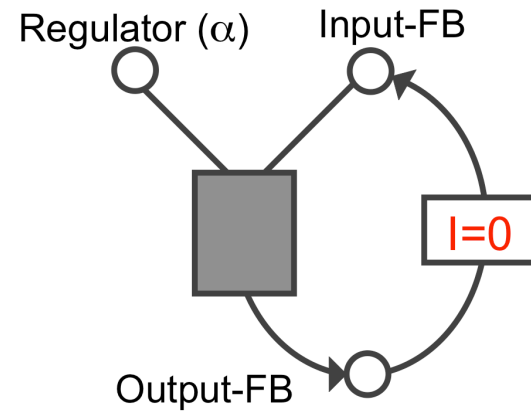
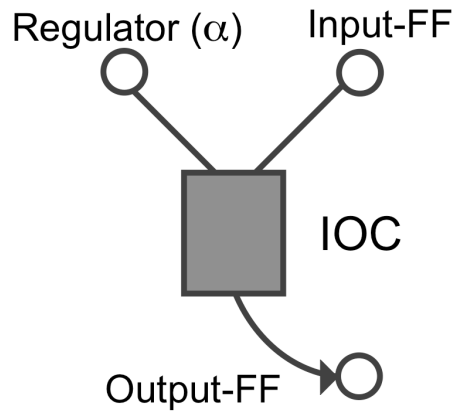
predict
feedback
responses
(single
regulated
input)



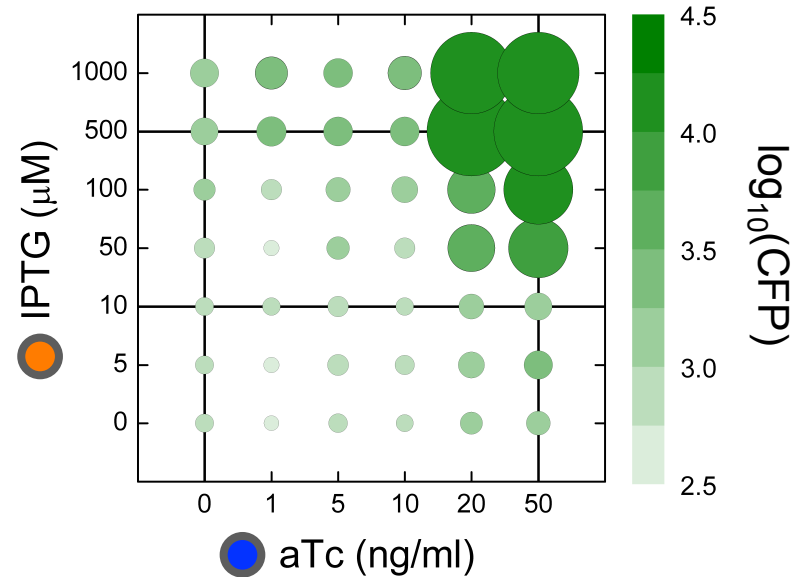
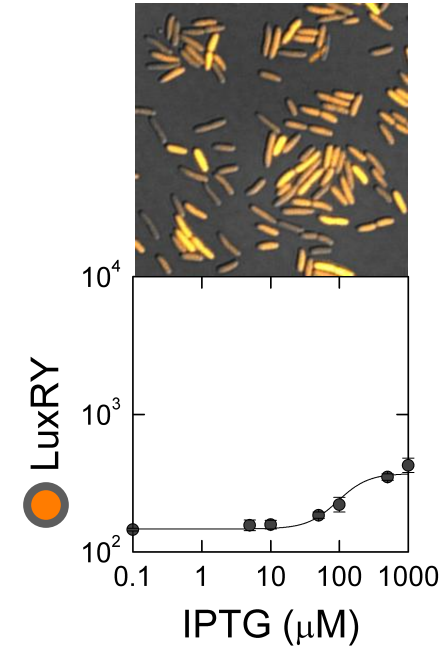
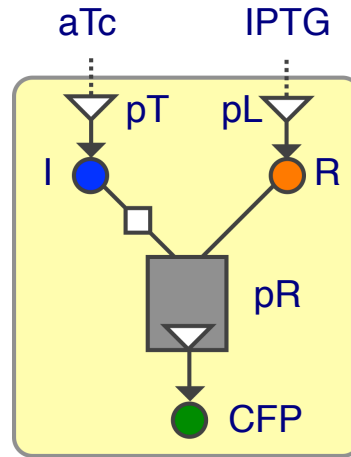
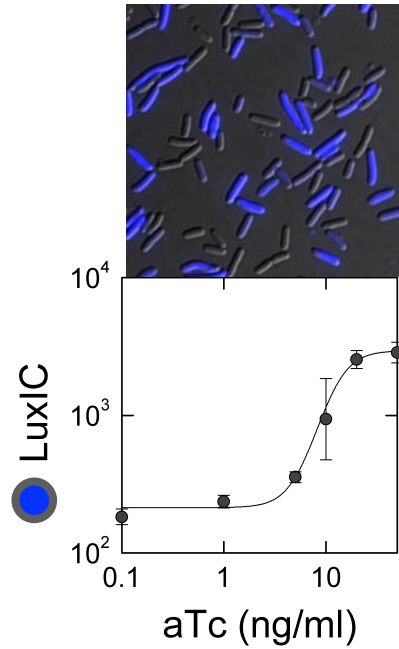
I-feedback

$$\frac{1}{\gamma_I} \frac{dY_I}{dt} = Q_I f(\mu \rho Y_I, \bar{Y}_R) - Y_I$$

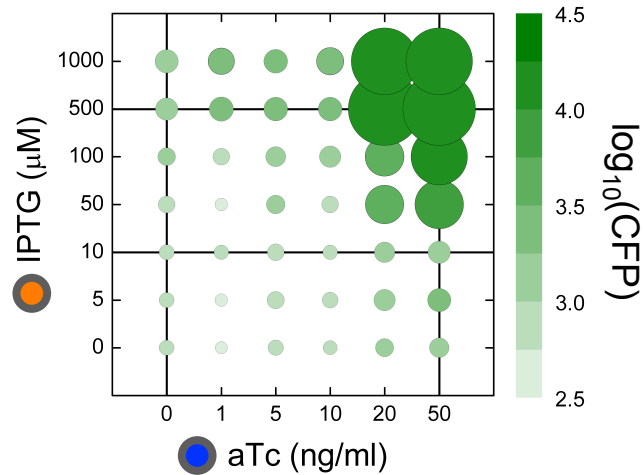
Inferring feedback behavior from feedforward data



Extracting biochemistry: the feedforward response

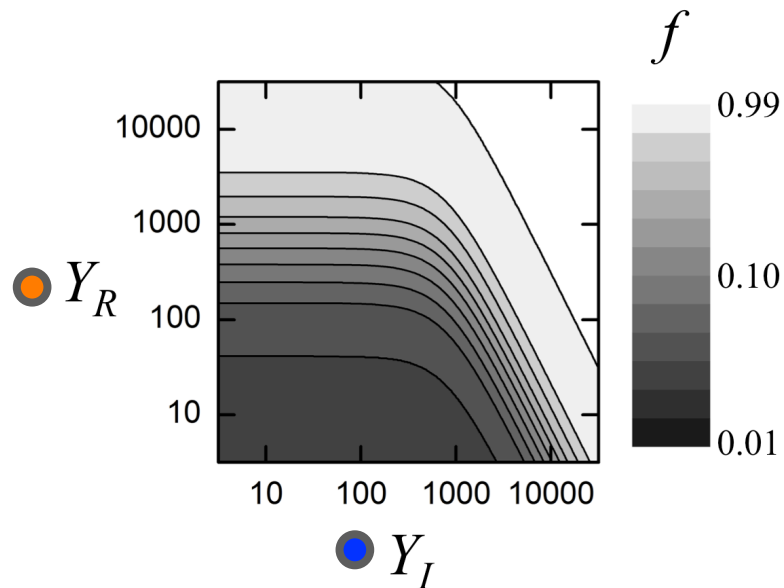


The 2-D promoter logic function of pR

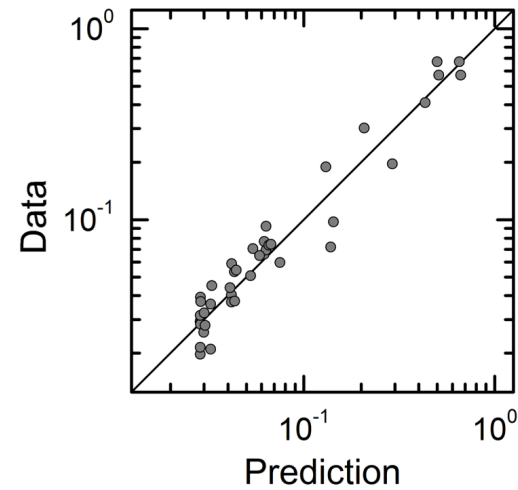


$$f(\mu\rho Y_I, Y_R) = \beta + (1 - \beta) \frac{Y_R^n (\tilde{\delta} + (\tilde{\mu}\rho Y_I)^m)}{1 + Y_R^n (\tilde{\delta} + (\tilde{\mu}\rho Y_I)^m)}$$

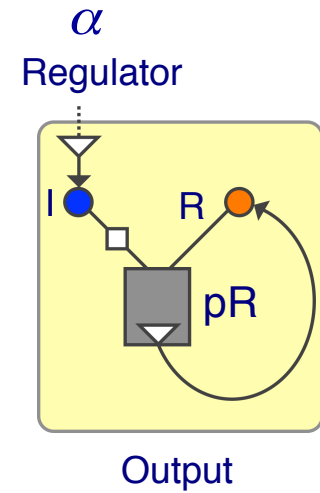
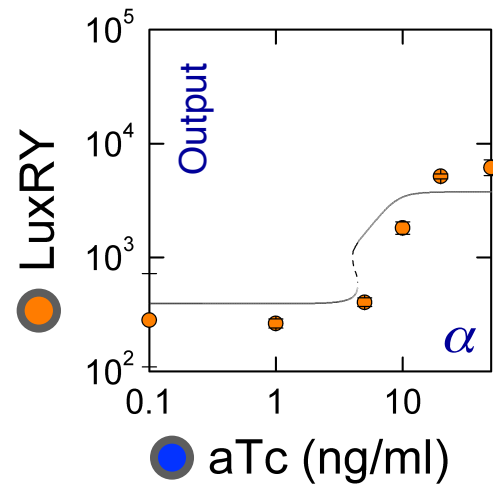
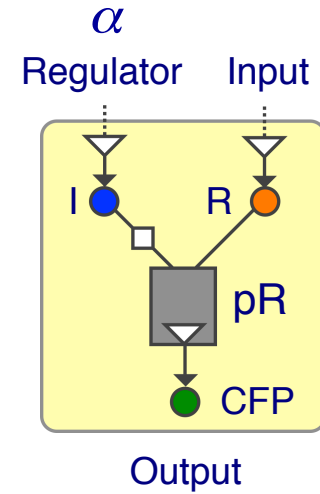
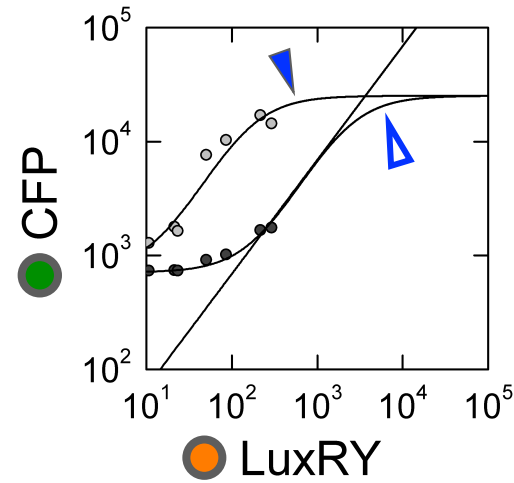
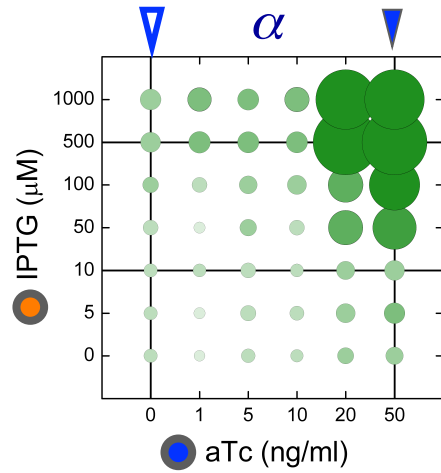
n : Hill coefficient of LuxR-DNA binding



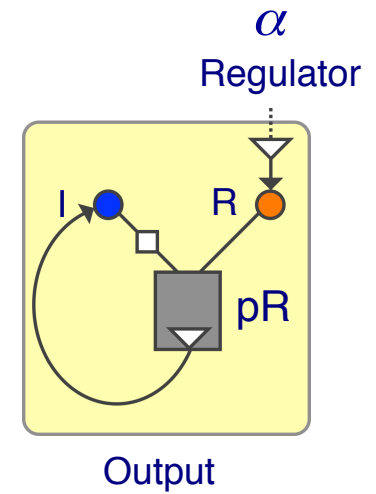
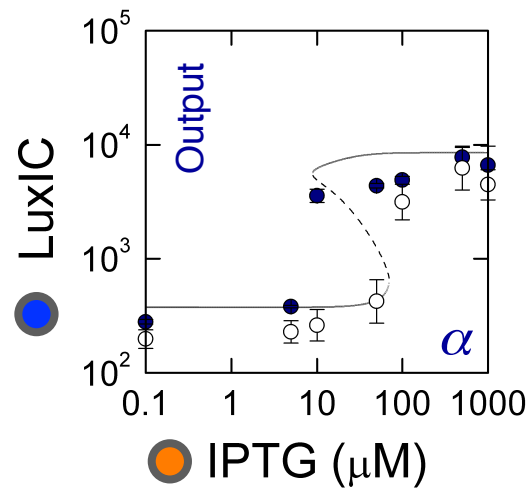
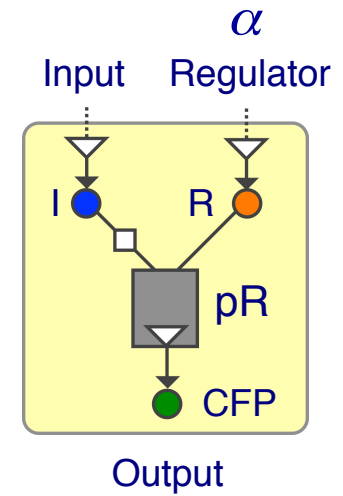
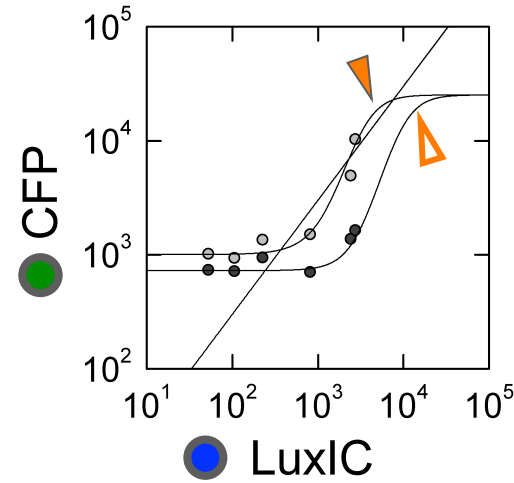
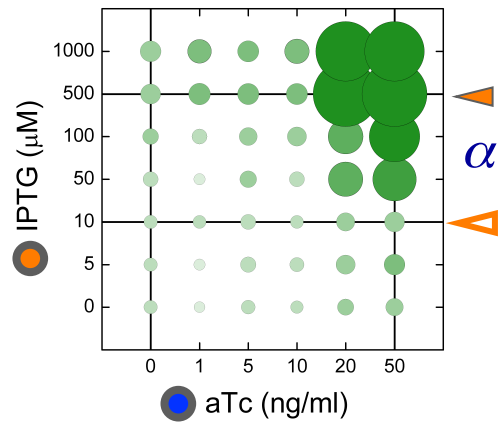
5 parameter fit to data



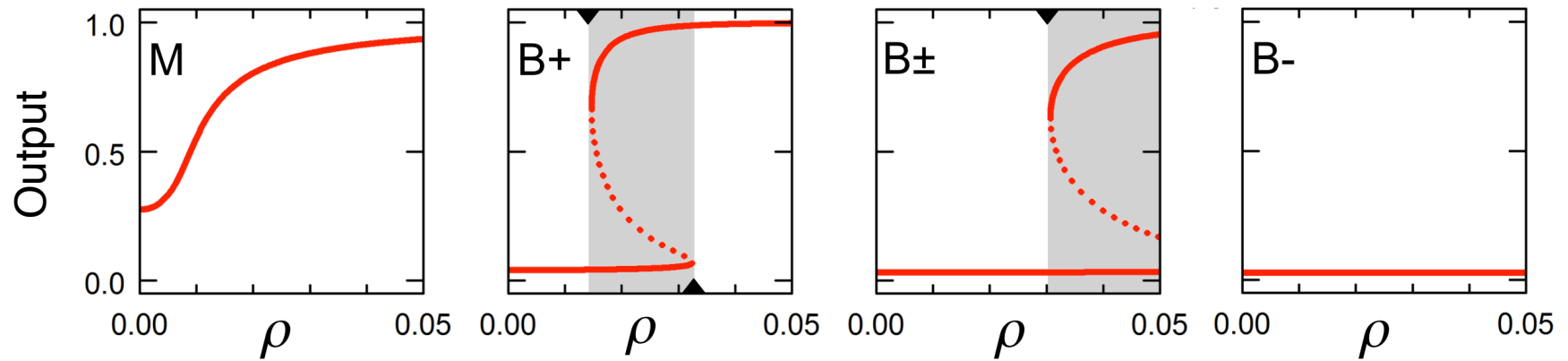
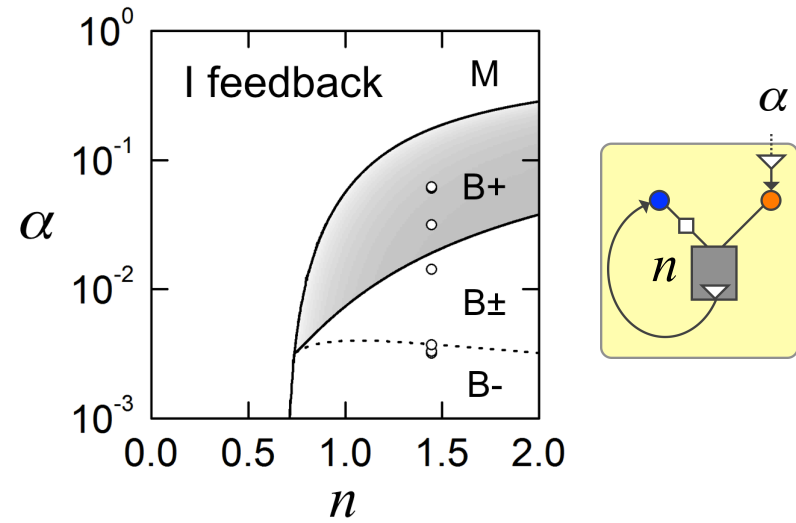
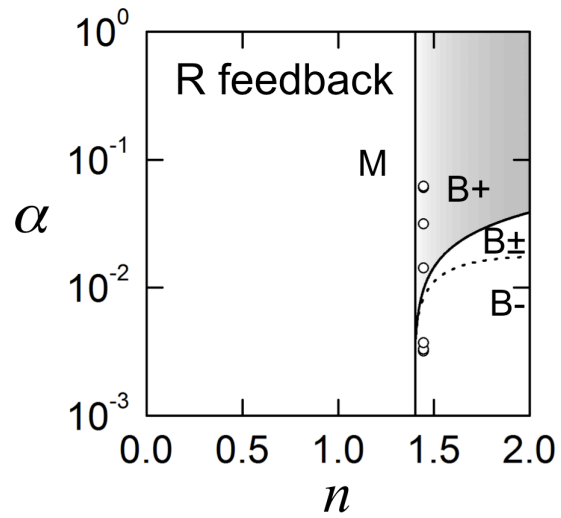
R-feedback response: monostable, smooth

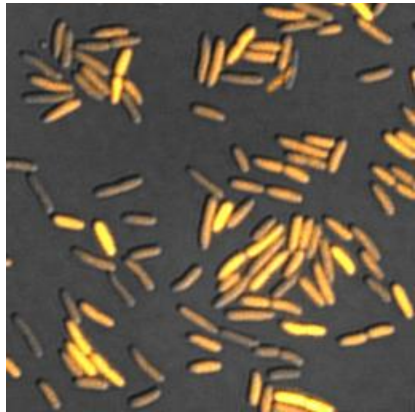


I-feedback response: hysteretic, switch-like



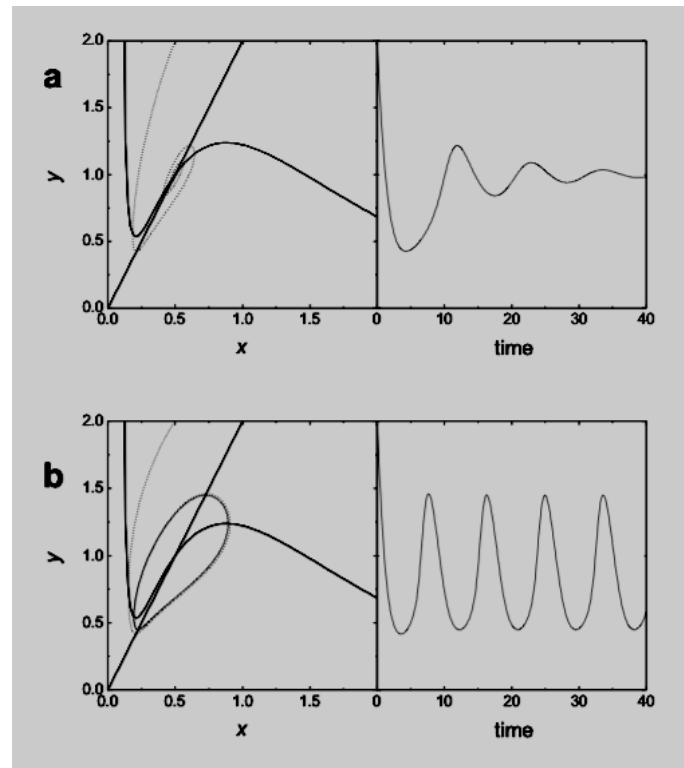
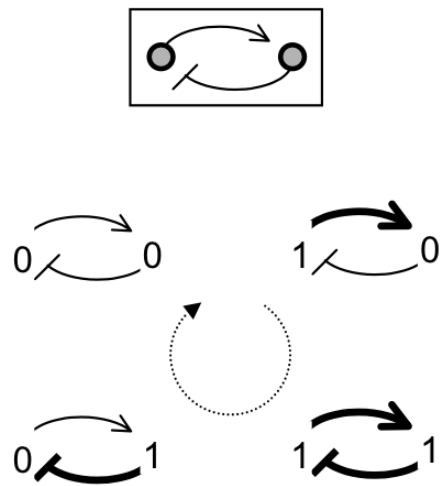
Phase diagram of density-dependent responses



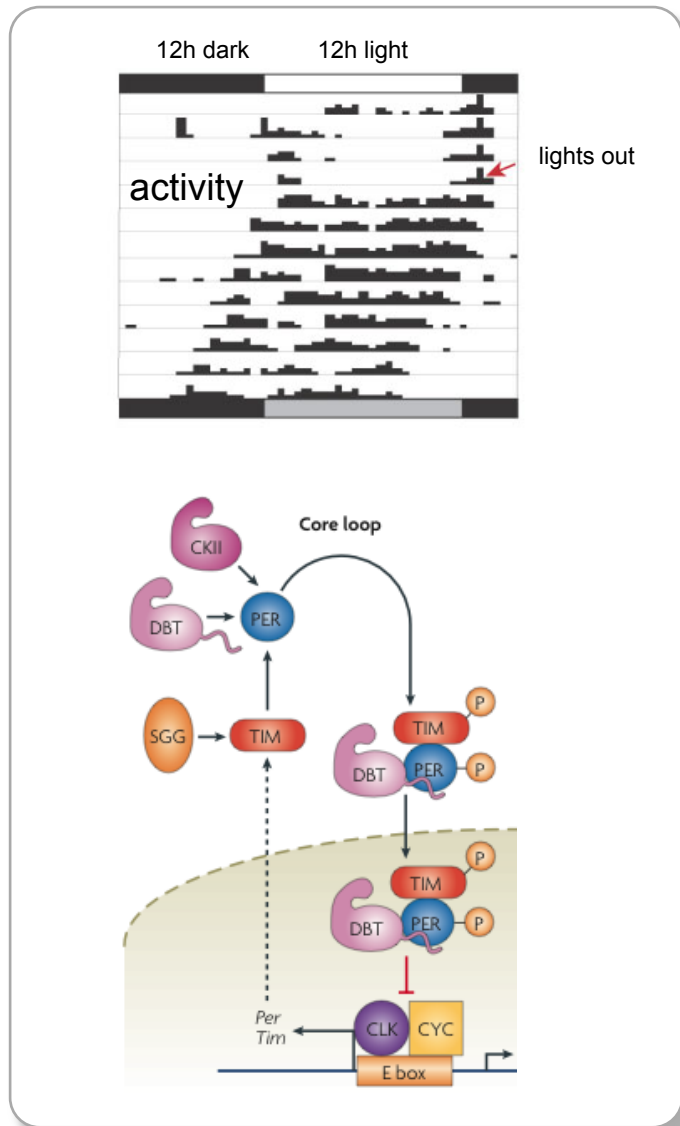


Endgame:
Building an oscillator from
the bottom up

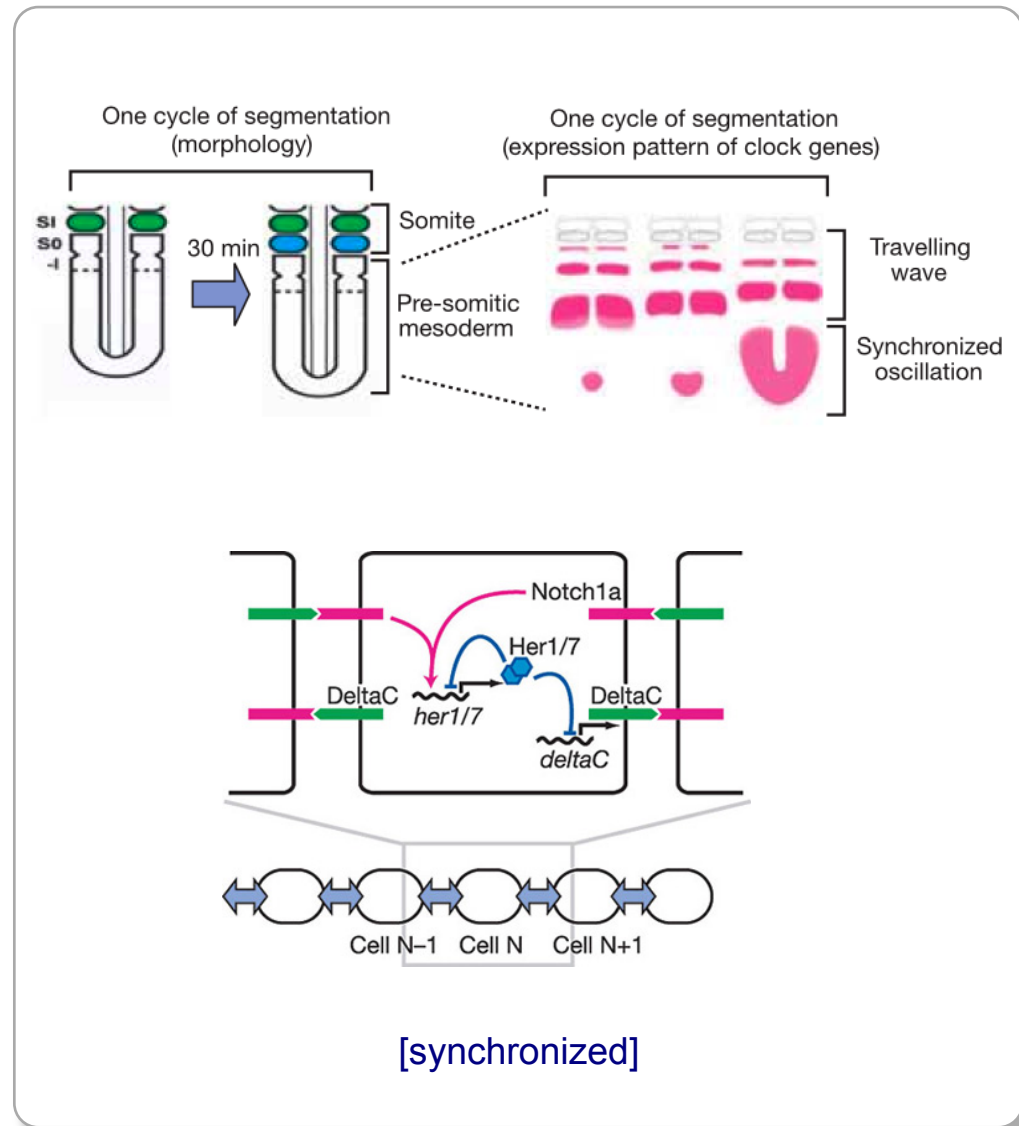
The “hysteretic oscillator” motif: negative feedback with delay



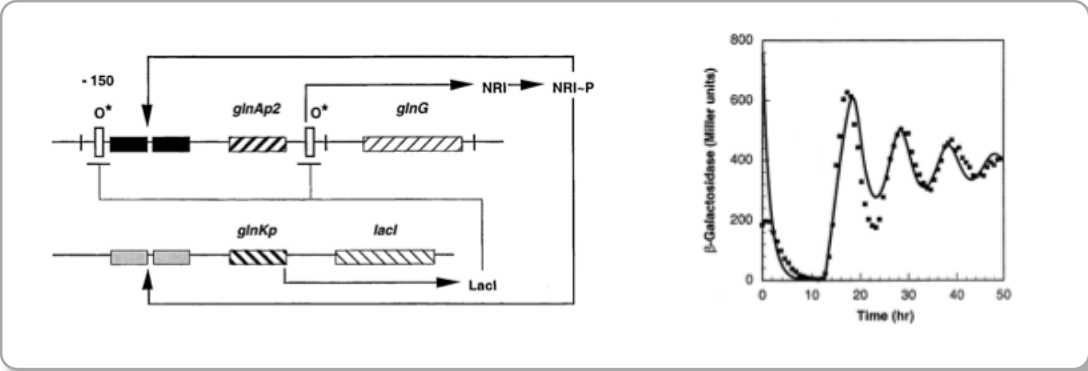
Drosophila circadian clock



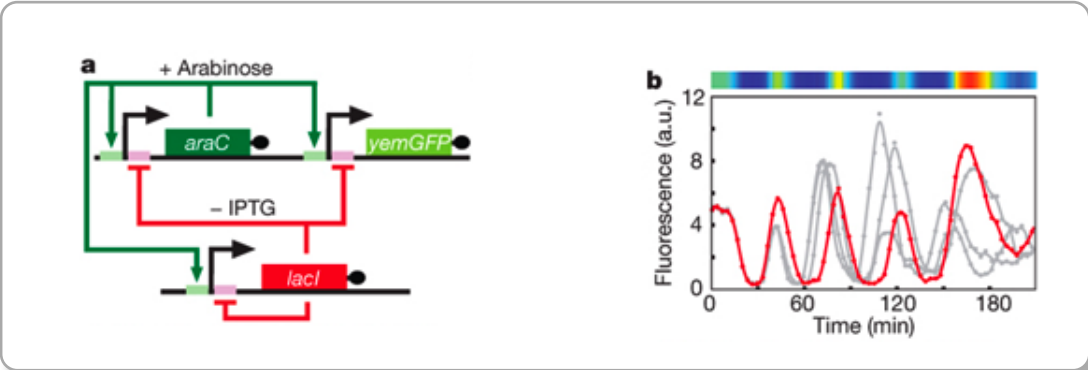
Zebrafish segmentation clock



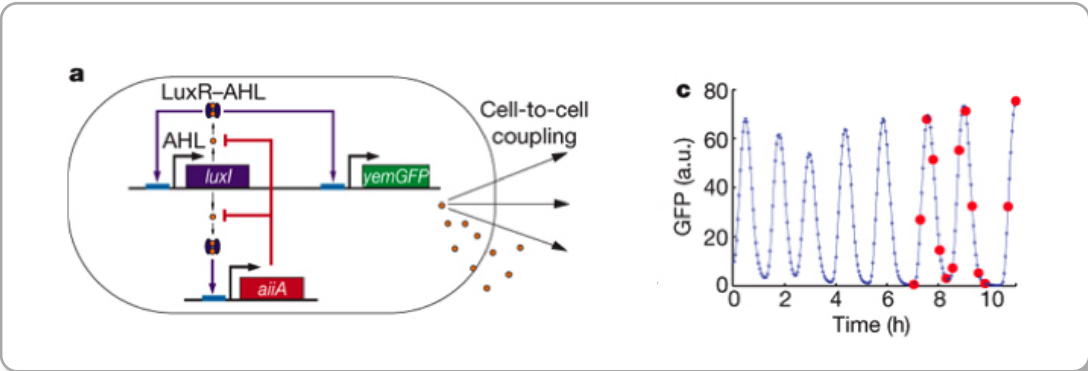
Synthetic oscillators



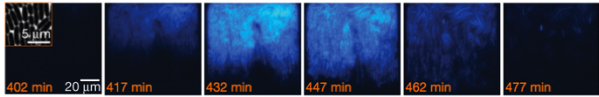
Atkinson et al., Cell, 2003



Stricker et al., Nature, 2008

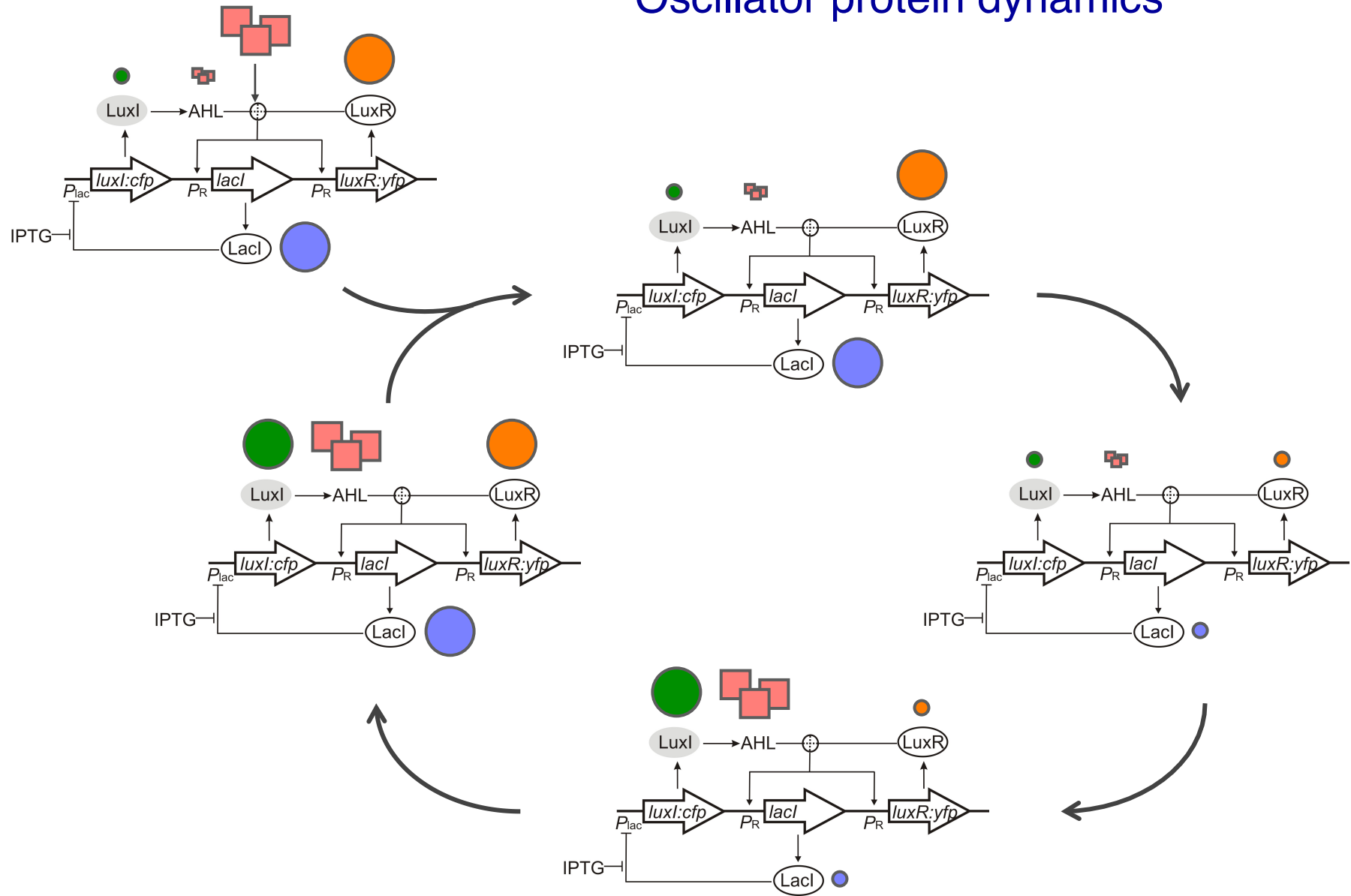


Danino et al., Nature, 2010

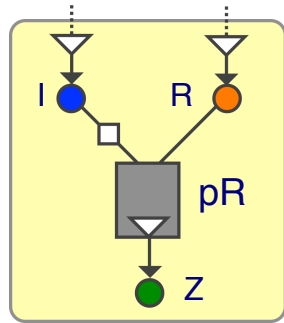


[synchronized]

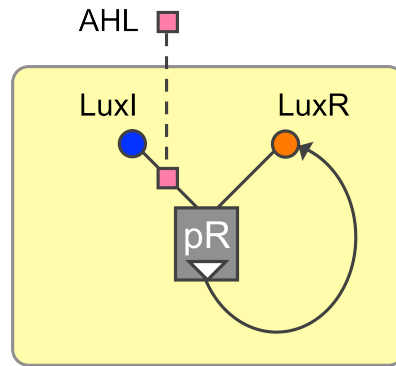
Oscillator protein dynamics



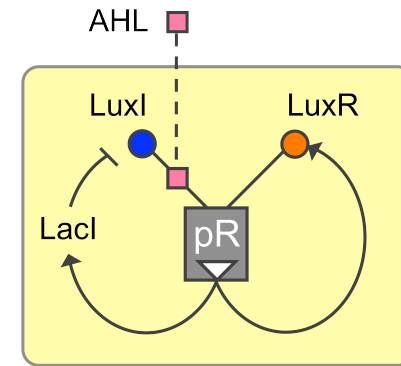
Roadmap: putting parts together to make the oscillator



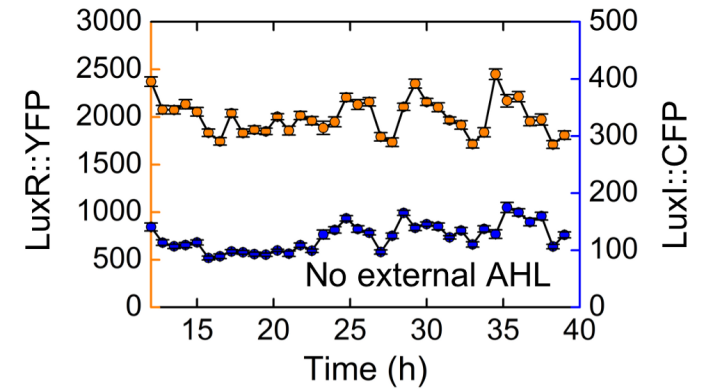
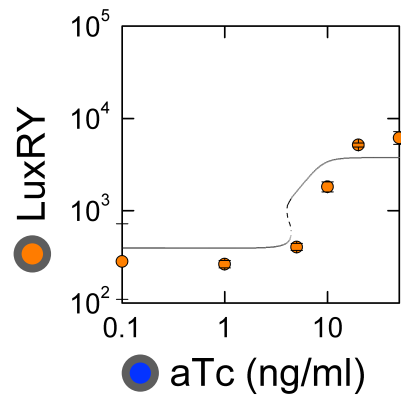
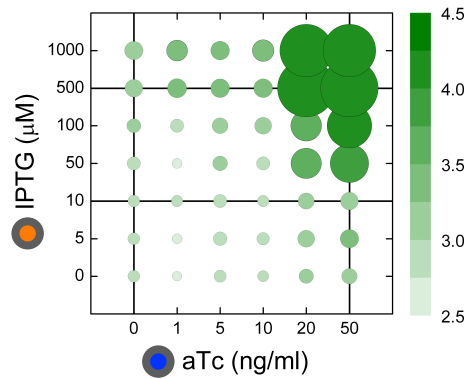
2 inputs



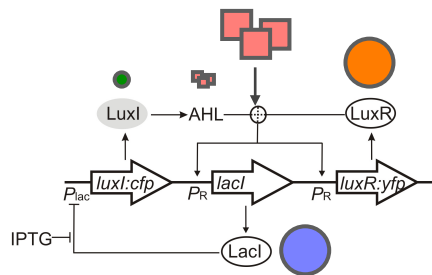
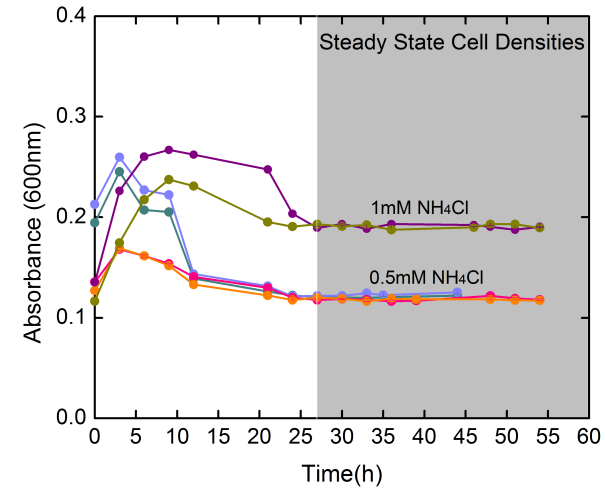
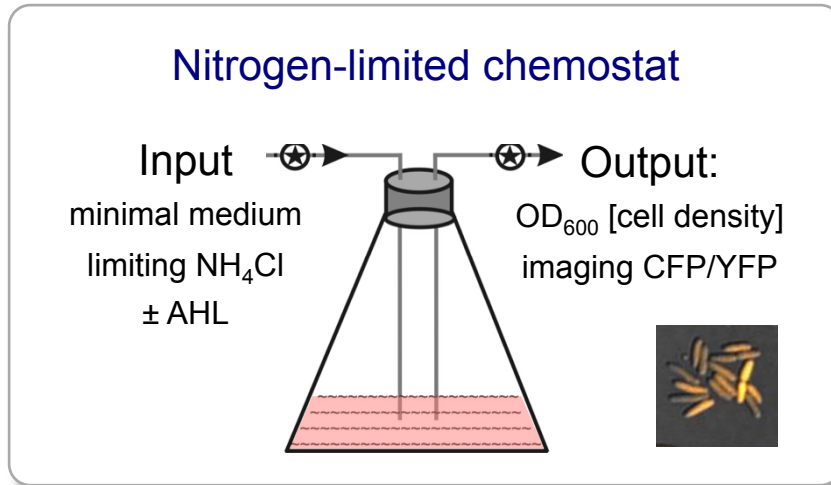
1 input



0 inputs



Experiments: The synchronization protocol



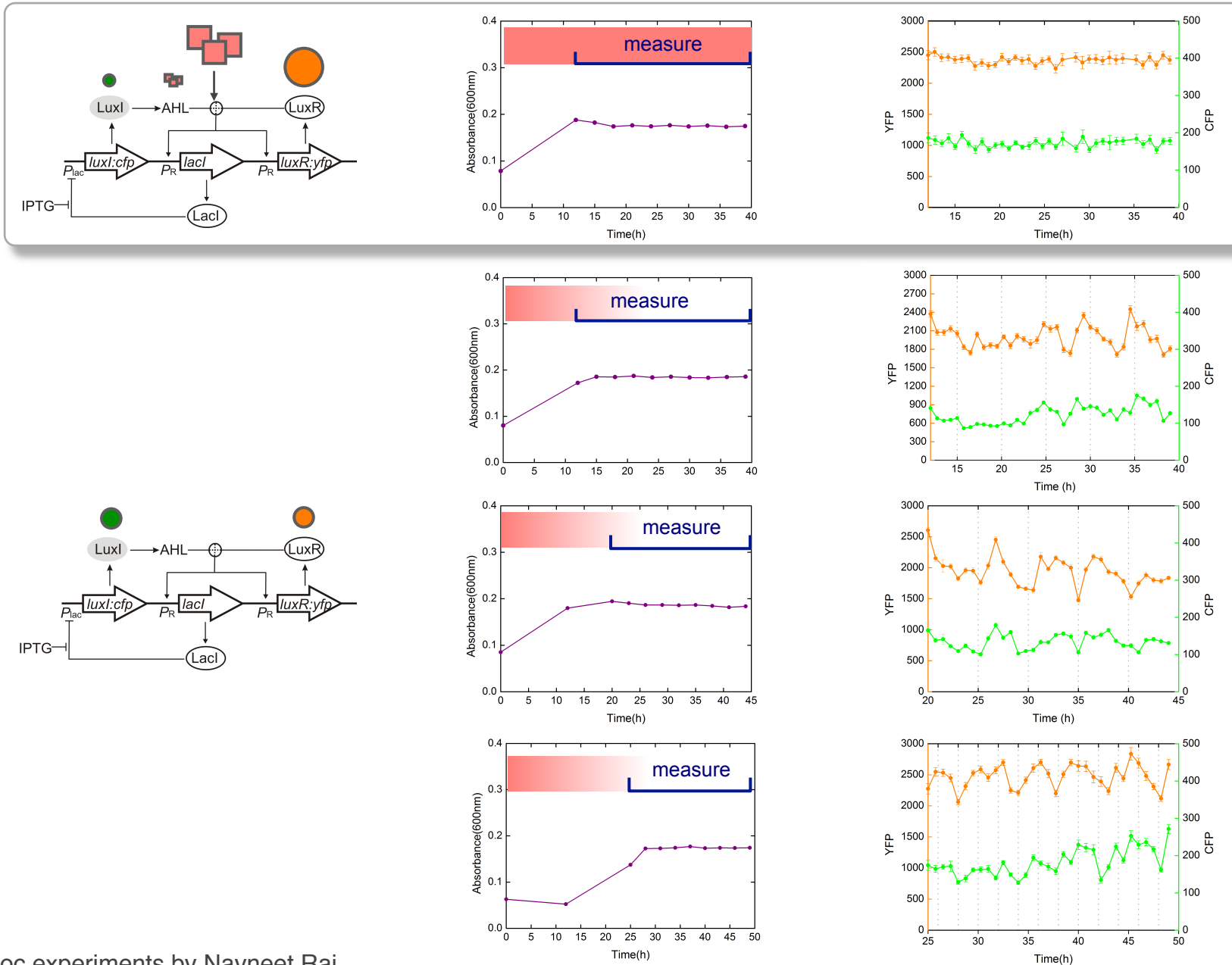
Control: continuous AHL



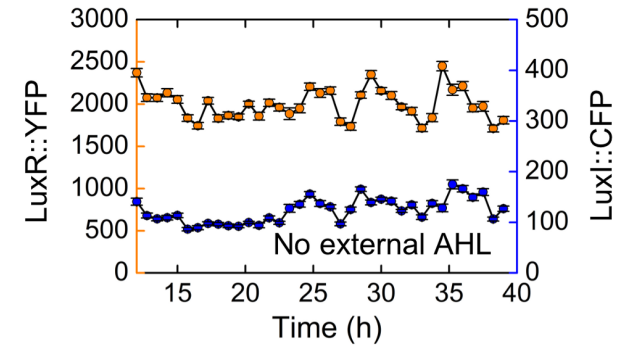
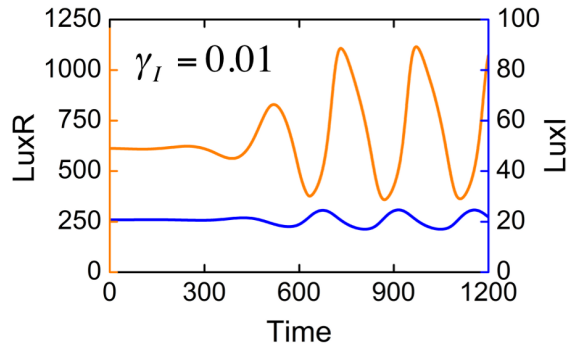
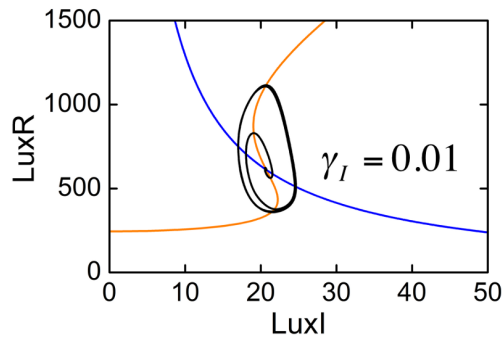
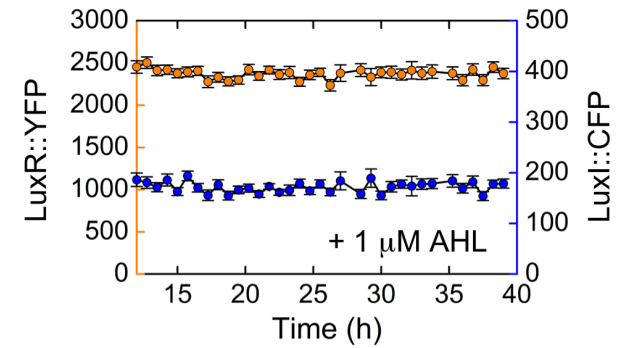
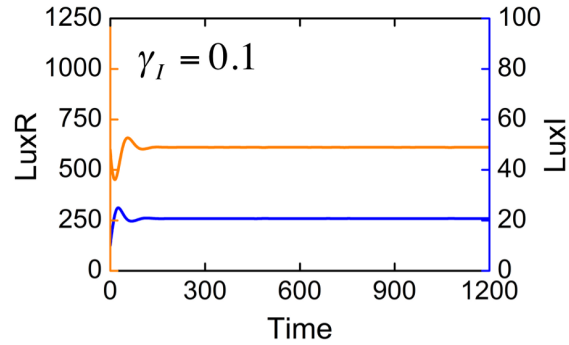
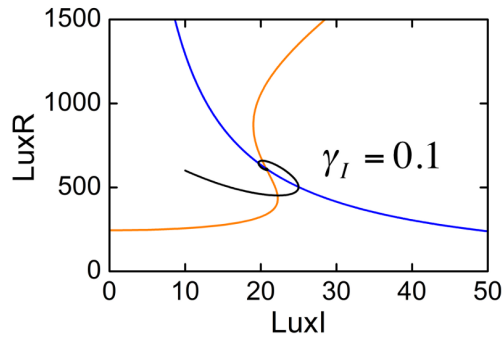
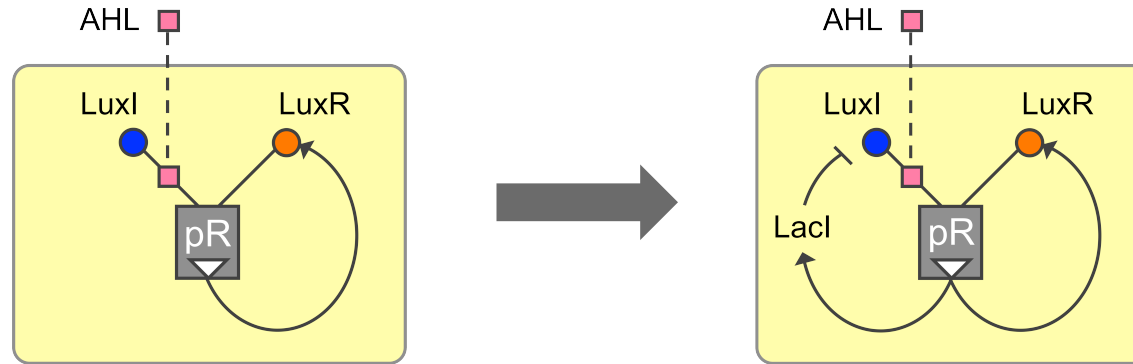
Test: Synchronize with 12h AHL



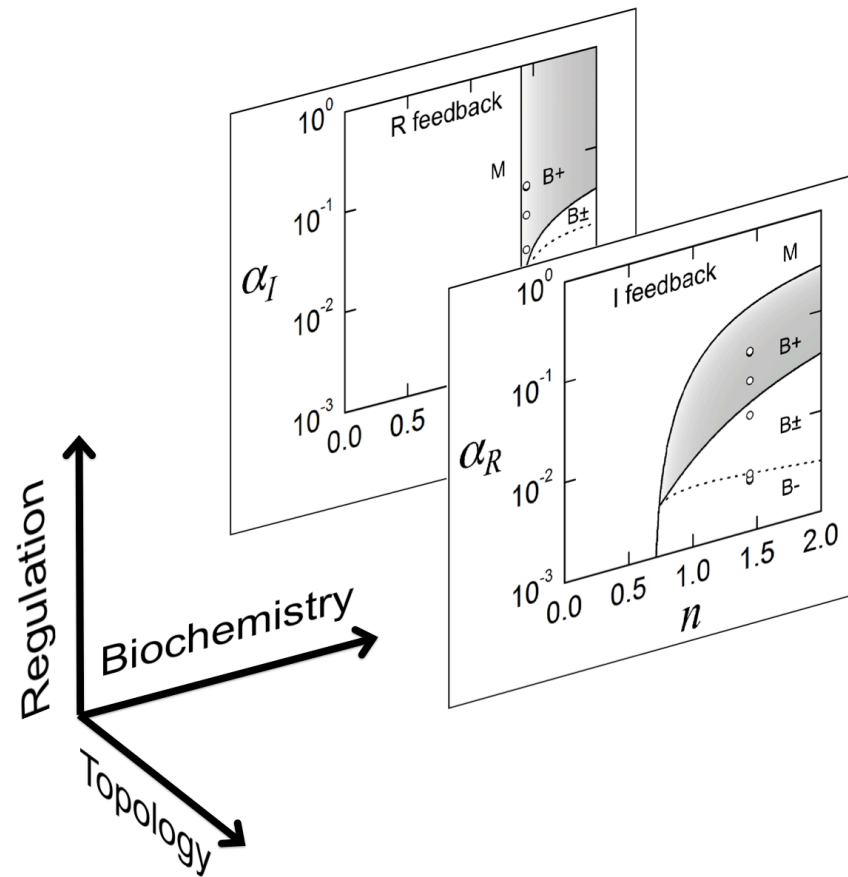
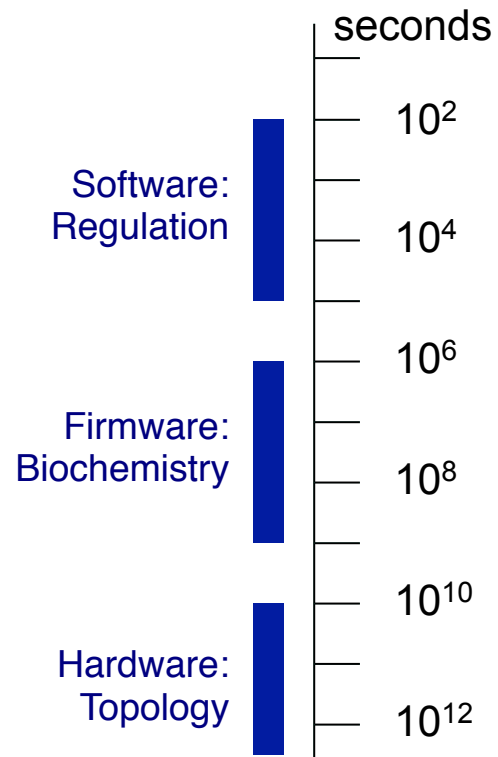
Experiments: Synchronized oscillations over 24h

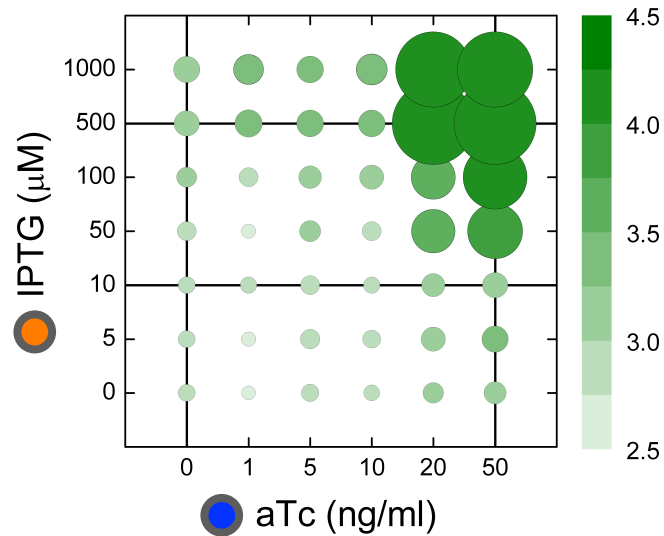


Promoter logic explains oscillations in dual feedback system



The regulation / biochemistry / topology hierarchy





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Stephen's), Vini Gautam (Delhi University), Senthil Kumar (PSG
Coimbatore)

Further reading:

> Rai, N. et al., Prediction by promoter logic in bacterial quorum sensing. *PLoS Comp. Biol.* 8, e1002361 (2012).

> Thattai, M. Using topology to tame the complex biochemistry of genetic networks. *Phil. Trans. Roy. Soc. A* 20110548 (2012).



thattai@ncbs.res.in