

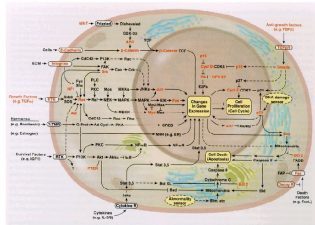
Mechanisms for Noise Attenuation in Molecular Biology Signaling Pathways

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University of California, Irvine

May 25, 2011
On the occasion of Eduardo's 60th birthday

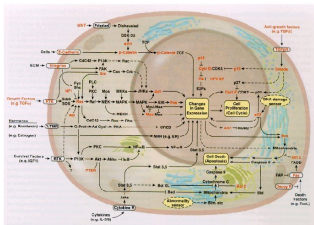
Feedback and noise in biological systems

- "Redundantly" many positive or negative feedback loops

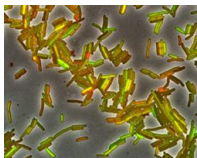


Feedback and noise in biological systems

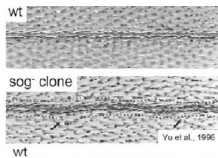
- "Redundantly" many positive or negative feedback loops



- noise (transcription, thermal fluctuation, volume changing, etc.)



noisy gene expressions



zig-zag vein

How does feedback affect a system's noise property?

- positive feedback **amplifies** noise and negative feedback **attenuates** noise (A. Becskei and L. Serrano, 2000; U. Alon, 2007)

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- no strong correlations between the sign of feedbacks and their noise properties (S. Hooshangi and R. Weiss, 2006)

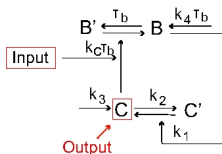
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is there a quantity (rather than the sign of FD) to unify these results?

One-loop and two-loop systems

one-loop system

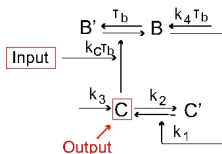


$$c' = k_1 b(1 - c) - k_2 c + k_3$$

$$b' = (k_c s(t)c(1 - b) - b + k_4) T_b$$

One-loop and two-loop systems

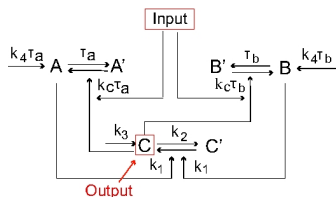
one-loop system



$$c' = k_1 b(1 - c) - k_2 c + k_3$$

$$b' = (k_c s(t)c(1 - b) - b + k_4) T_b$$

two-loop system

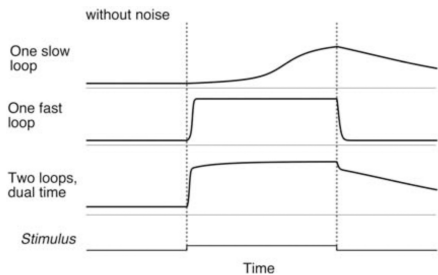


$$c' = k_1(b + a)(1 - c) - k_2 c + k_3$$

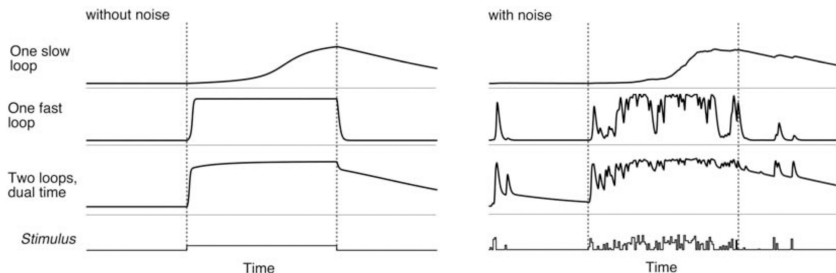
$$b' = (k_c s(t)c(1 - b) - b + k_4) T_b$$

$$a' = (k_c s(t)c(1 - a) - a + k_4) T_a$$

Dynamical and noise properties



Dynamical and noise properties

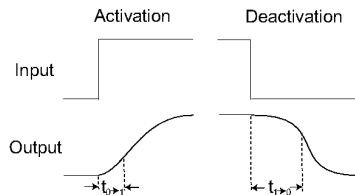


O. Brandman et al., *Science*, 2005

system's intrinsic time scales are crucial to noise attenuation

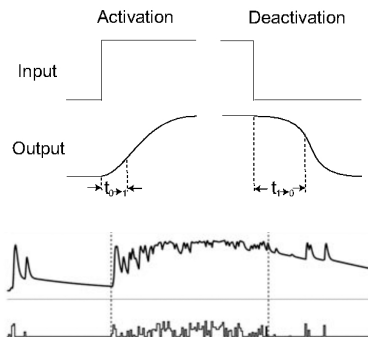
A conjecture

define **activation** and **deactivation** time scales.



A conjecture

define **activation** and **deactivation** time scales.



..... guess: at the "on" state,

$$t_{1 \rightarrow 0} \gg 1/\omega, t_{0 \rightarrow 1} \ll 1/\omega \Rightarrow \text{better noise attenuation}$$

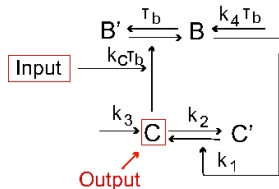
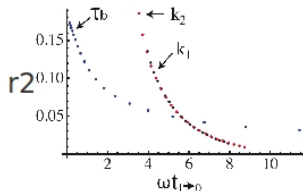
ω : the frequency of the input noise.

Testing the conjecture

define **noise amplification rate**: $r_2 = \frac{\text{std}(\text{output})/\langle \text{output} \rangle}{\text{std}(\text{input})/\langle \text{input} \rangle}$

G. Hornung and N. Barkai, *PLoS Comp. Bio.*, 2008

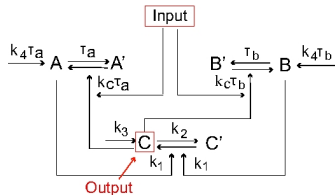
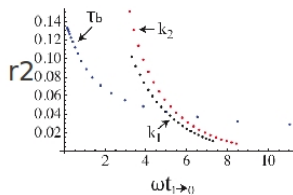
testing in the one-loop system:



$t_{1 \rightarrow 0} \gg 1/\omega \Rightarrow$ better noise attenuation

Testing the conjecture

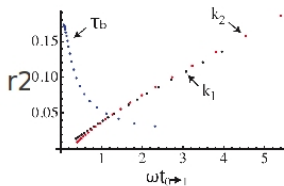
in the two-loop system:



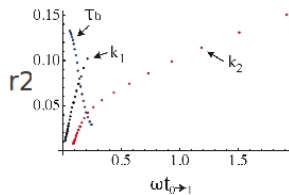
$t_{1 \rightarrow 0} \gg 1/\omega \Rightarrow$ better noise attenuation

Testing the conjecture

one-loop system



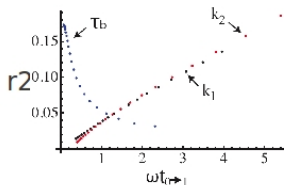
two-loop system



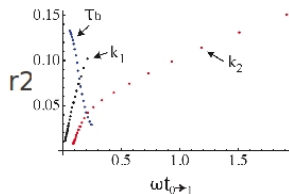
why is τ_b inconsistent?

Testing the conjecture

one-loop system

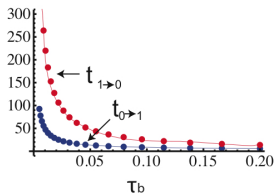


two-loop system



why is τ_b inconsistent?

... a closer look



is there a simple way to take into account both changes?

A critical quantity: signed activation time

- signed activation time (SAT) = $t_{1 \rightarrow 0} - t_{0 \rightarrow 1}$
- SAT has a **negative** relation with the noise amplification rate



Analytical studies using the Fluctuation Dissipation Thm

$$r_2^2 \approx \frac{\tau_b/\omega}{\langle s \rangle (k_1 k_c/k_2 - 1)(k_1/k_2 + 1) \frac{k_c}{k_c+1}}$$

key observation:

- r_2 negatively depends on k_c and k_1/k_2 .

linear analysis of the noise-free ODE:

- SAT positively depends on k_c and k_1/k_2

⇒ r_2 negatively depends on SAT = $t_{1 \rightarrow 0} - t_{0 \rightarrow 1}$

Analytical studies - two-time-scale analysis

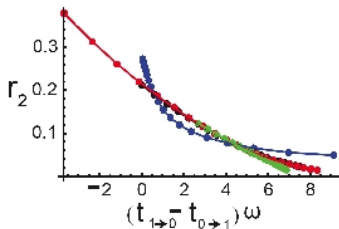
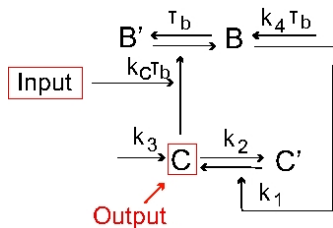
$$\begin{aligned}c' &= k_1 b(1 - c) - k_2 c + k_3 \\b' &= (k_c s(t)c(1 - b) - b + k_4)\tau_b\end{aligned}$$

When $\varepsilon := \tau_b \ll k_2$, \exists two time scales: $t_f = t$ and $t_s = \varepsilon t$,

$$\begin{aligned}c &= c_0(t_s, t_f) + \varepsilon c_1(t_s, t_f) + \varepsilon^2 c_2(t_s, t_f) + \dots \\b &= b_0(t_s, t_f) + \varepsilon b_1(t_s, t_f) + \varepsilon^2 b_2(t_s, t_f) + \dots\end{aligned}$$

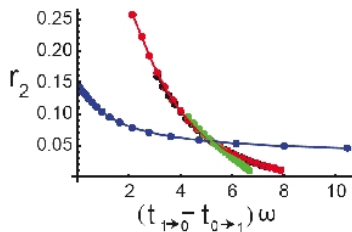
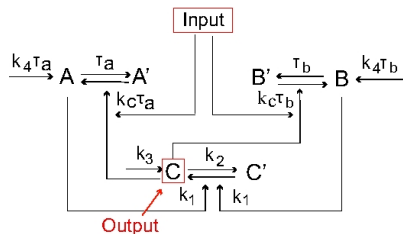
- $s(t)$ varies on the time scale of $t_f \Rightarrow$ noise is filtered out in c_0 .
- $s(t)$ varies on the time scale of $t_s \Rightarrow$ noise persists in c_0 .

SAT in one-loop systems



r_2 decreases in SAT

SAT in two-loop systems



r_2 decreases in SAT

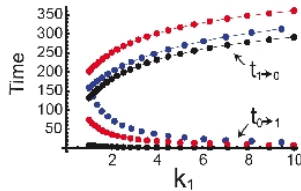
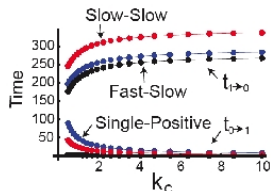
How to achieve large SAT?

- linear stability analysis

	single	slow-slow	fast-slow
activation	$\frac{k_c+1}{(K_a+1)k_c}$	$\frac{k_c+1}{(2K_a+1)^2 k_c}$	$\frac{k_c+1}{2(2K_a+1)^2 k_c}$
deactivation	$\frac{(K_a+1)k_c}{1+k_c}$	$\frac{(2K_a+1)k_c}{1+k_c}$	$\frac{(K_a+1/2)k_c}{1+k_c}$

\Rightarrow large k_c and $K_a := k_1/k_2$

- simulations



Why multiple loops?

- faster activation

	single	slow-slow	fast-slow
activation	$\frac{k_c+1}{(K_a+1)k_c}$	$\frac{k_c+1}{(2K_a+1)^2 k_c}$	$\frac{k_c+1}{2(2K_a+1)^2 k_c}$

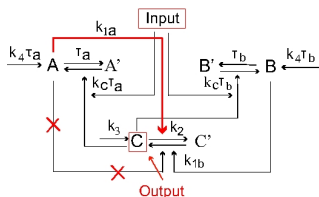
- more robust (w.r.t. parameter changes)

$k_c \in (0.5, 10)$	single	slow-slow	fast-slow
activation	(8.2, 89.9)	(0.8, 3.9)	(4.5, 43.7)

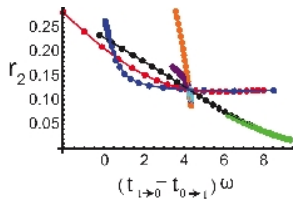
$k_1 \in (1, 10)$	single	slow-slow	fast-slow
activation	(15.6, 158.2)	(0.9, 8.4)	(8.9, 75)

Does SAT apply to negative feedback systems?

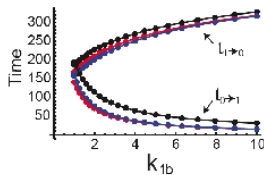
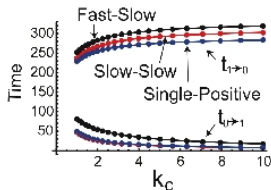
a system with negative feedback



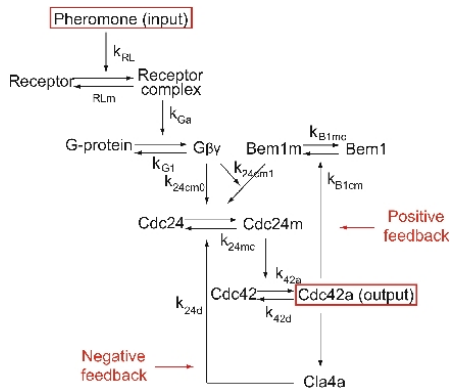
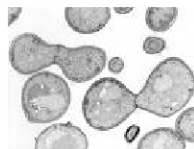
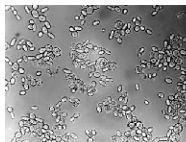
r_2 decreases in SAT



to achieve large SAT:

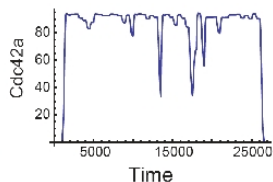
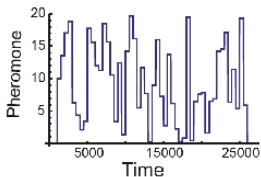
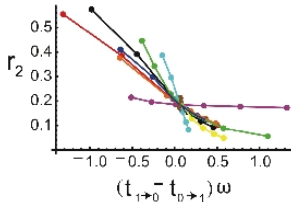
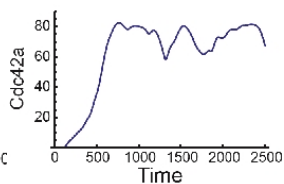
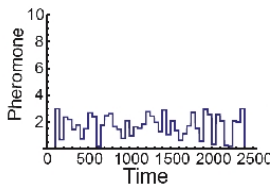
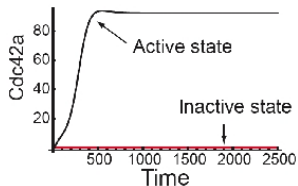


The yeast polarization system

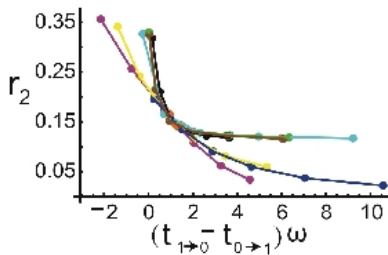
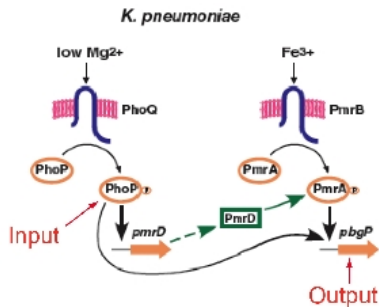


A non-spatial model,
simplified from C.S. Chou et al., 2008

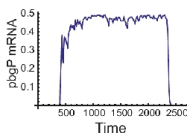
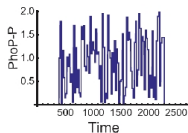
SAT in the yeast cell polarization system



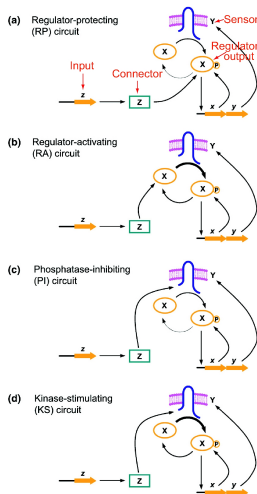
SAT in a Polymyxin B resistance model



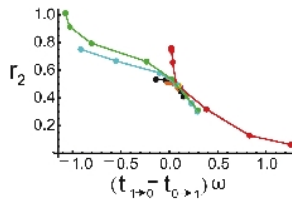
13 parameters are varied in ± 3 range.



SAT in connector-mediated models



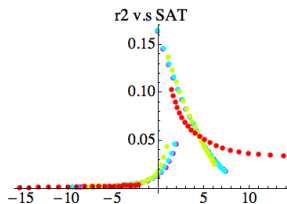
	RP	RA	KS	PI
activation	30.1	30.4	4.5	62.4
deactivation	45.2	37.2	5.9	6.4
SAT	0.76	0.34	0.07	-2.8
r_2	0.14	0.34	0.5	0.85



A.Y. Mitrophanov and
E.A. Groisman, 2010

Summary and future work

- proposed a new quantity $SAT = t_{1 \rightarrow 0} - t_{0 \rightarrow 1}$
- at ON state, r_2 (noise amplification rate) decreases in SAT.
- SAT is the intrinsic time scale determined by network structure and parameters
- additional positive feedback drastically reduces the activation time and makes the system more robust to parameter variations
- what is the prediction for OFF state? bistable system? PDE?



Acknowledgements

Qing Nie

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(Dept. of Mathematics, UCI)

Tau-Mu Yi

(Dept. of Developmental and Cell Biology, UCI)

Congratulations on your achievements!

Happy Birthday!