

Mathematical Cell Biology Graduate Summer Course
University of British Columbia, May 1-31, 2012
Leah Edelstein-Keshet

Simple biochemical motifs (1)



www.math.ubc.ca/~keshet/MCB2012/

Biochemical (and gene) circuits

Switches, oscillators, adaptation, and
amplification circuits

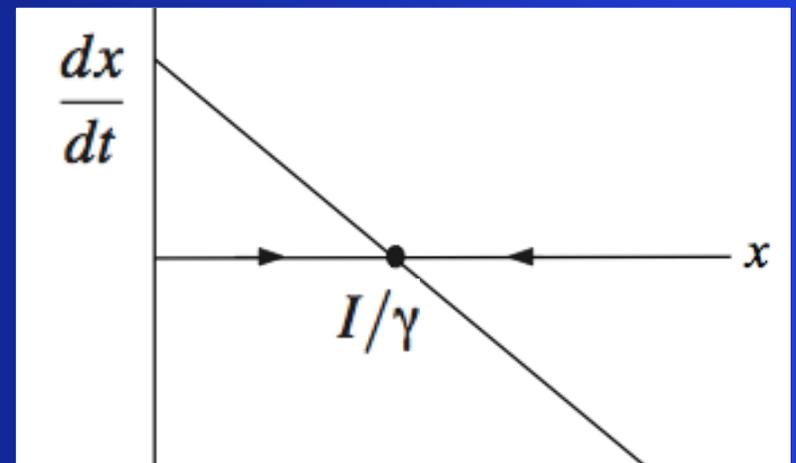
Production-decay at constant rates



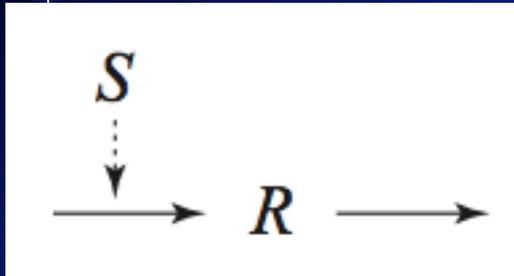
$$\frac{dx}{dt} = I - \gamma x$$

$I, \gamma > 0$ constants.

Unique positive Steady state

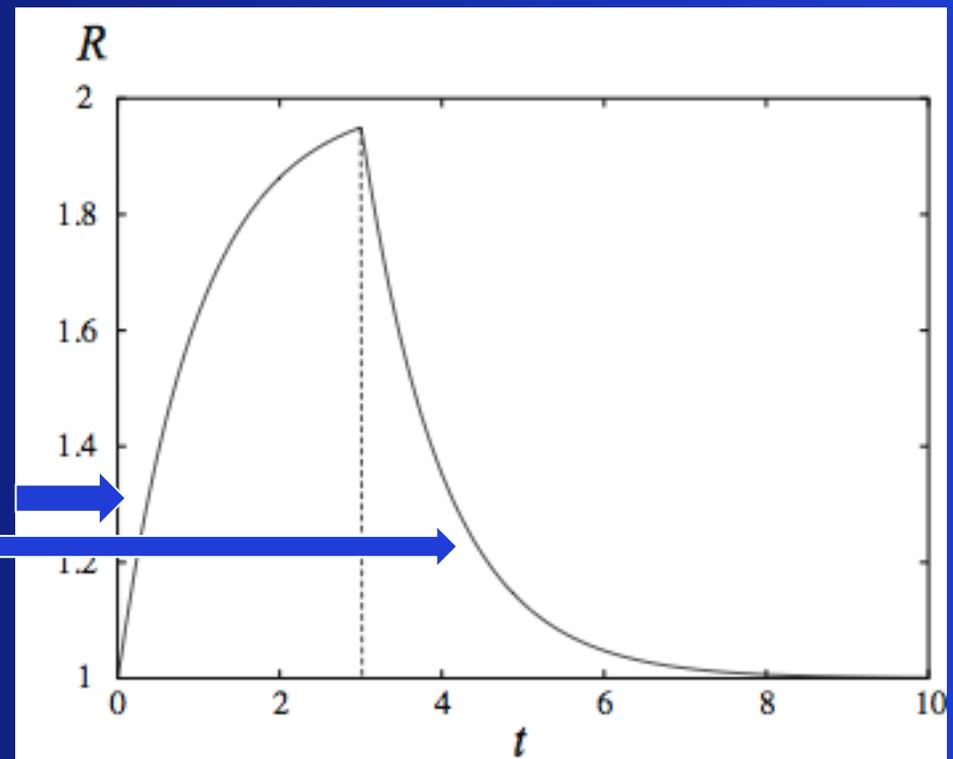


Signal-induced Production

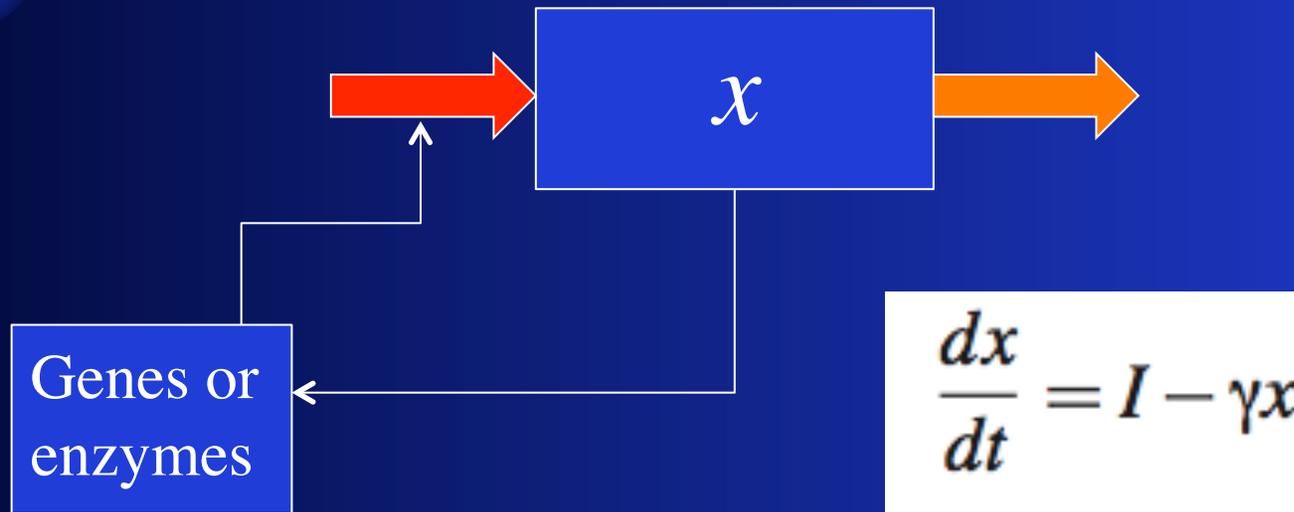


$$\frac{dR}{dt} = k_0 + k_1 S - k_2 R.$$

Note typical “ $1 - \exp(-k_2 t)$ ” rise and exponential decay tail



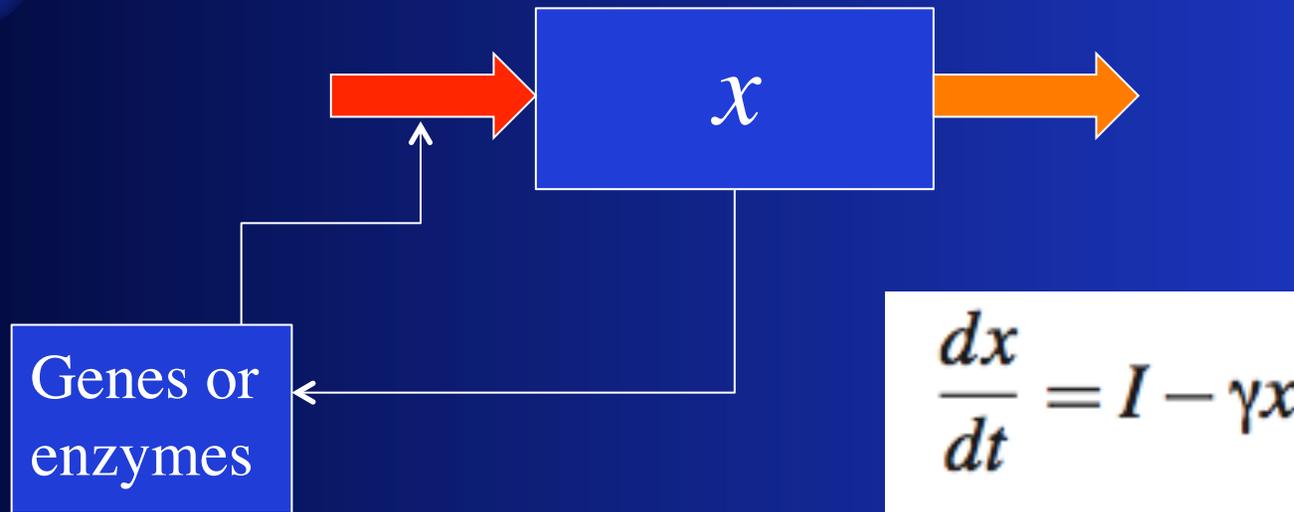
Feedback to production



$$\frac{dx}{dt} = I - \gamma x$$

I is now a function of x

Michaelian Feedback to production

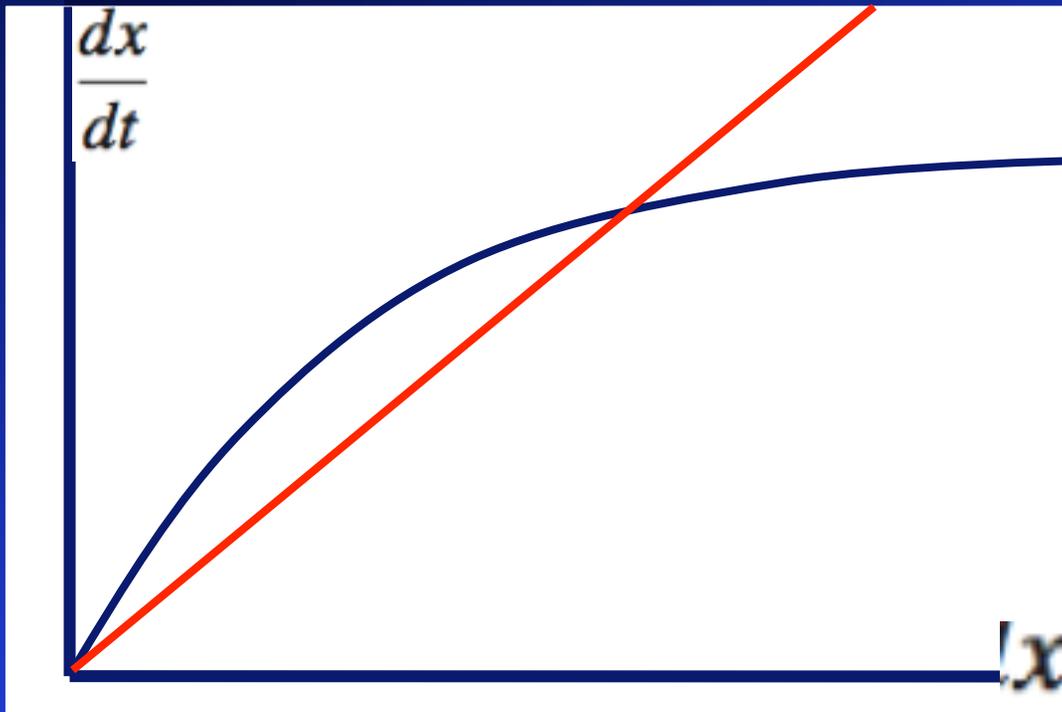


$$I(x) = I_0 + \frac{I_{max} x}{k_n + x}$$

Michaelian Feedback to production

$$I(x) = I_0 + \frac{I_{max} x}{k_n + x}$$

$$\frac{dx}{dt} = I - \gamma x$$

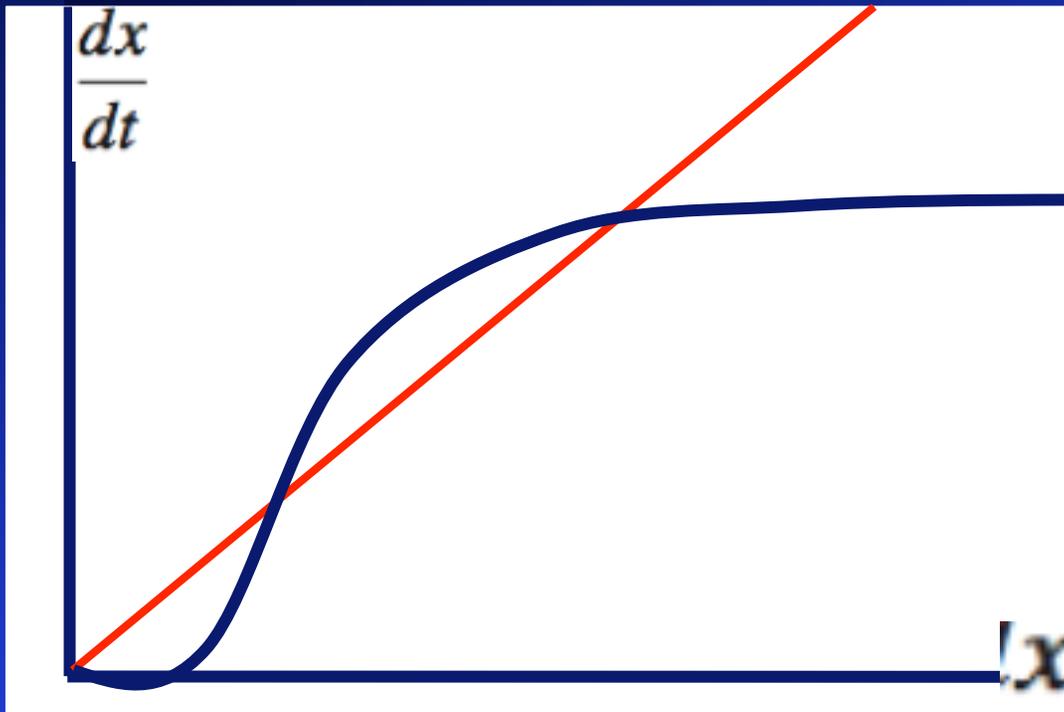


At most 2 steady states, one stable.

Sigmoidal Feedback to production

$$I(x) = I_0 + \frac{I_{max} x^2}{k_n^2 + x^2}$$

$$\frac{dx}{dt} = I - \gamma x$$



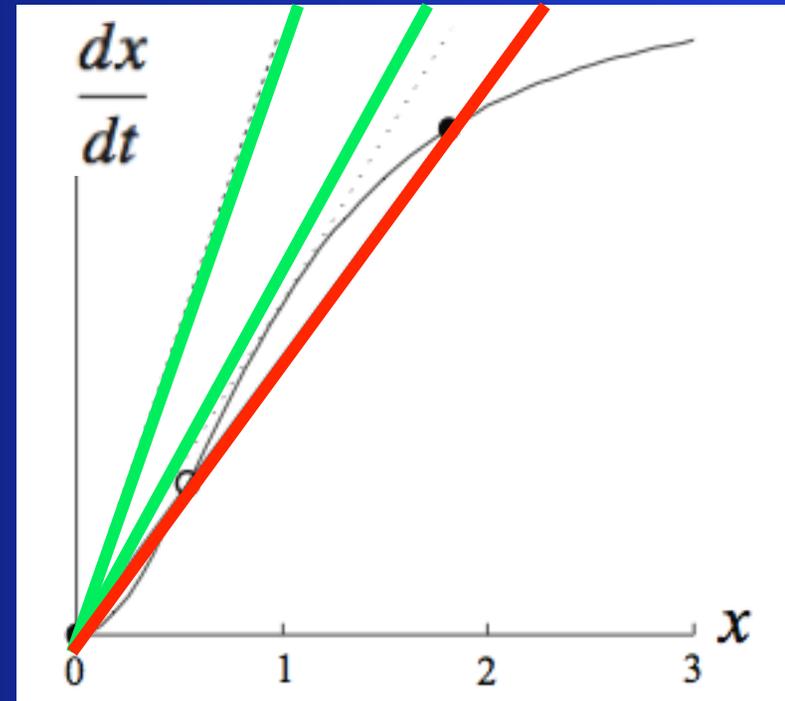
Up to 3 steady states, two stable.

“bistability”

Sigmoidal cont'd

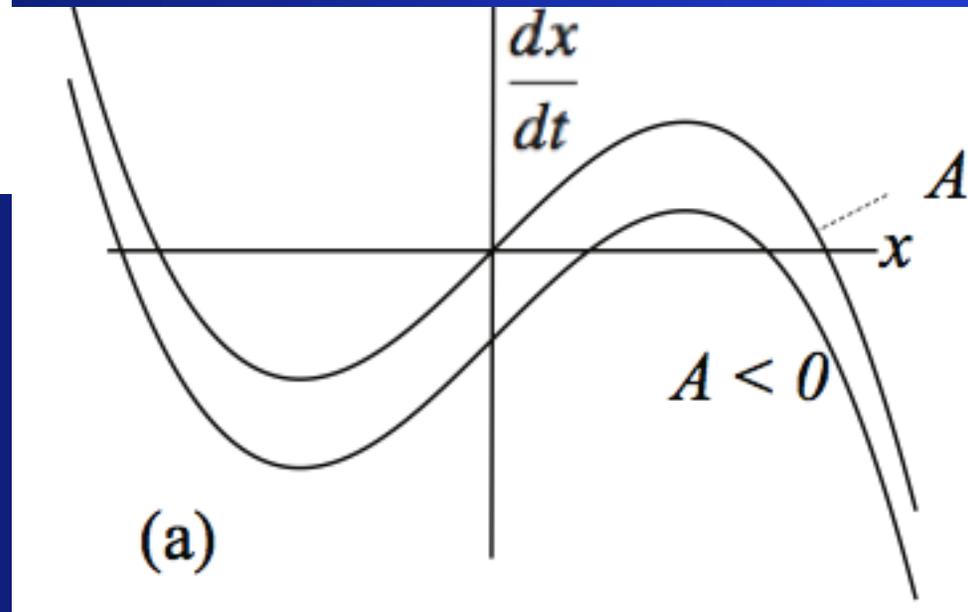
$$\frac{dx}{dt} = f(x) = \frac{x^2}{1+x^2} - mx + b$$

Actual number of steady states depends on parameters, e.g. on slope m (decay rate of x)

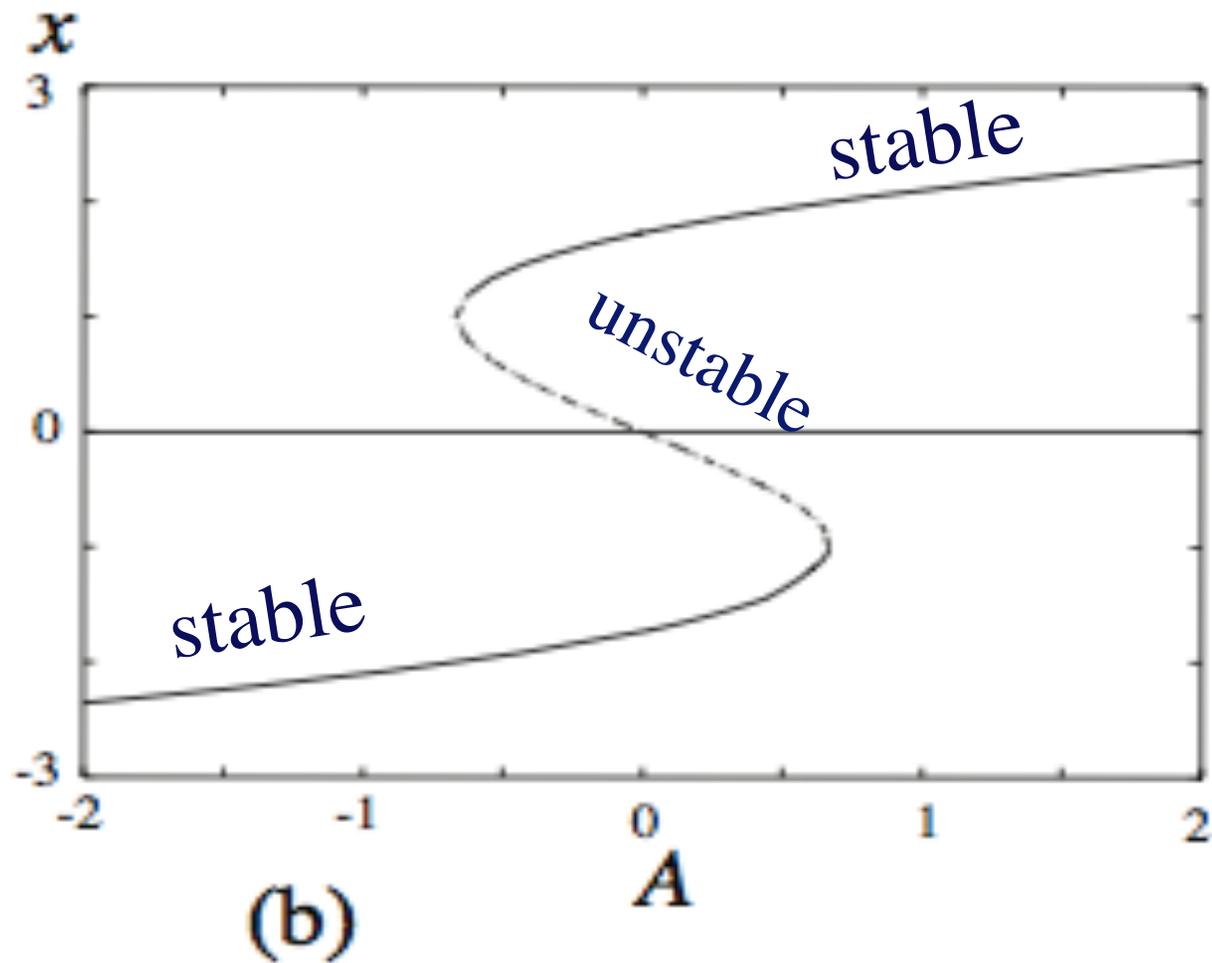


Generic bistability

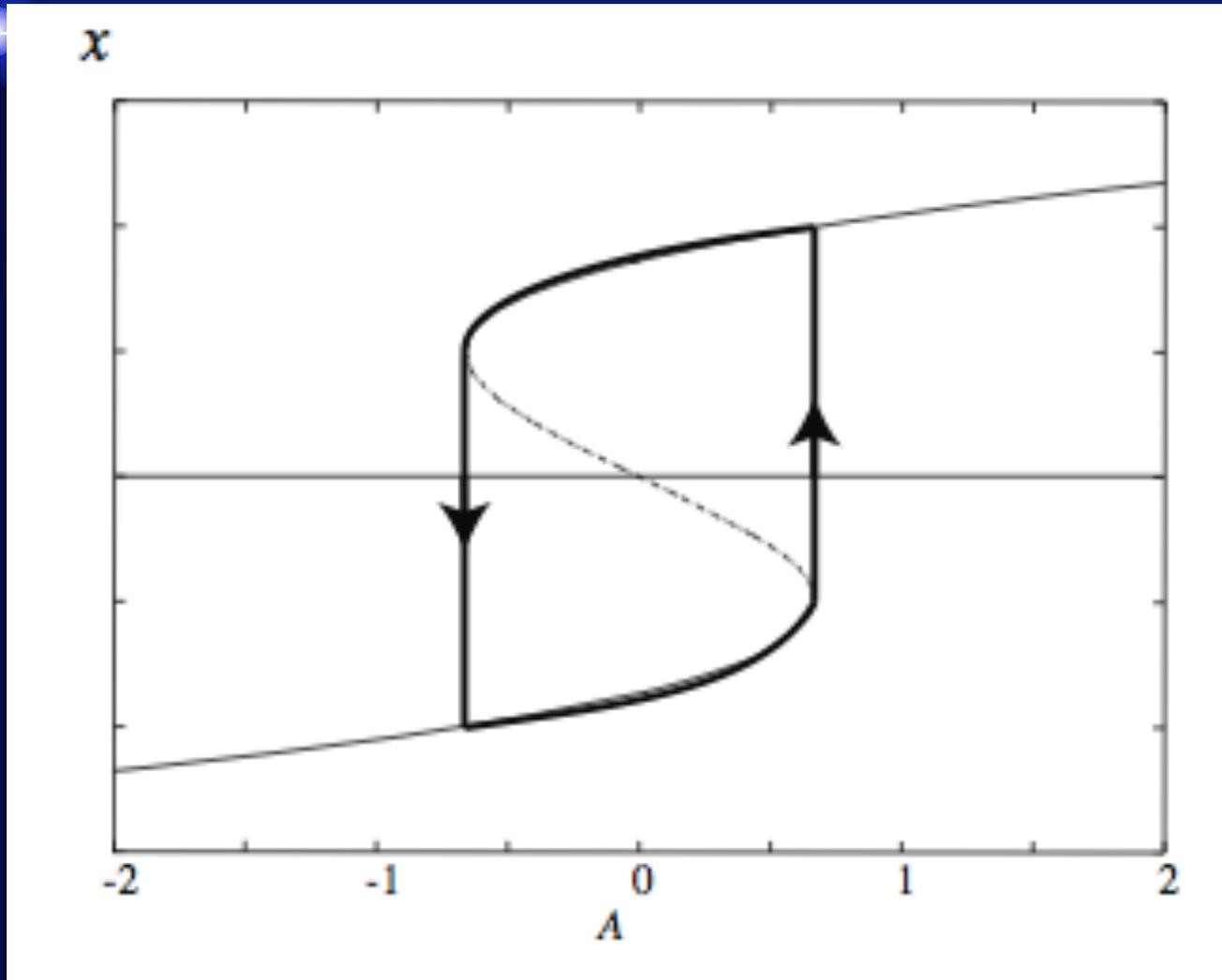
$$\frac{dx}{dt} = c \left(x - \frac{1}{3}x^3 + A \right)$$



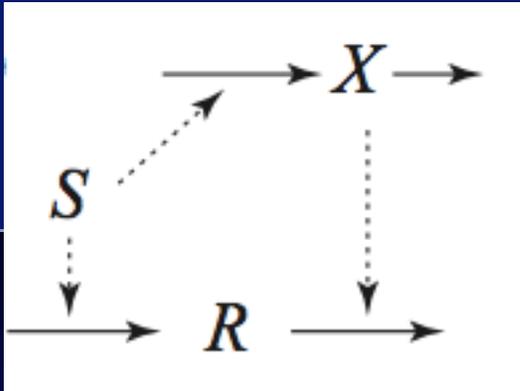
Bifurcation Diagram



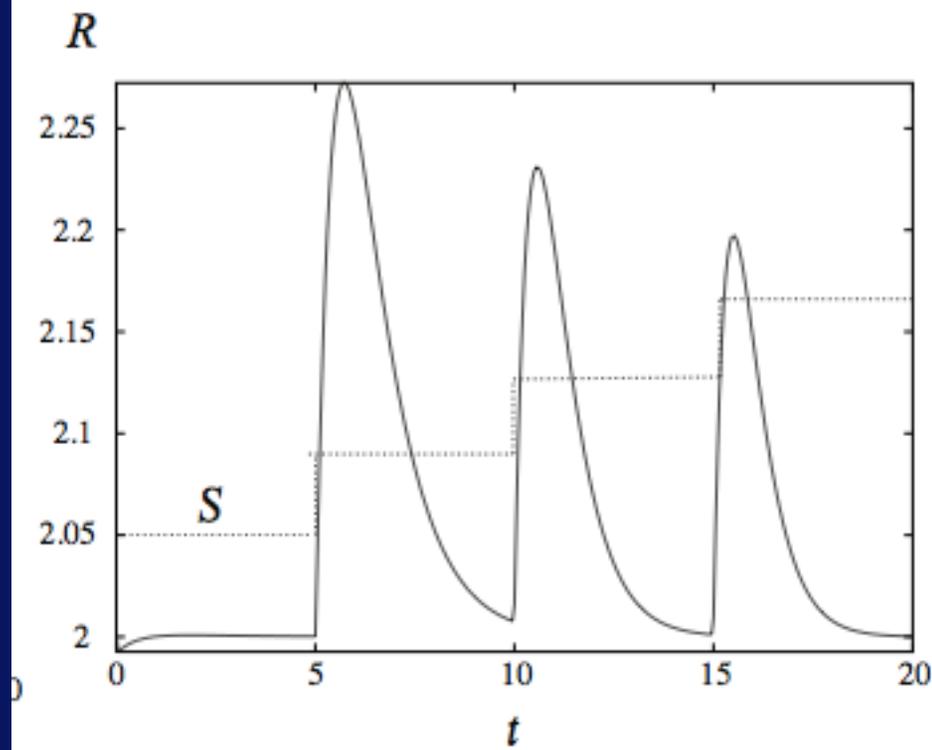
Hysteresis



Adaptation



$$\frac{dR}{dt} = k_1 S - k_2 X R,$$
$$\frac{dX}{dt} = k_3 S - k_4 X.$$



Mathematical Cell Biology Graduate Summer Course
University of British Columbia, May 1-31, 2012
Leah Edelstein-Keshet

Simple biochemical motifs (2)



www.math.ubc.ca/~keshet/MCB2012/

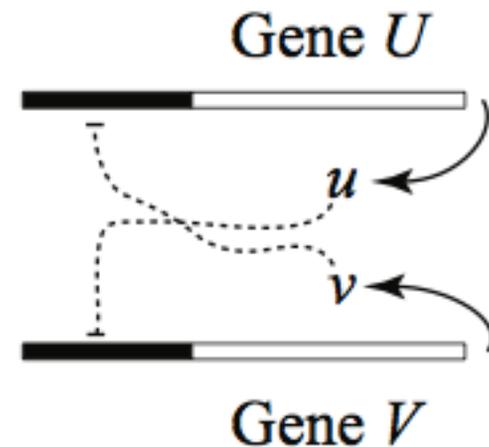
Genetic toggle switch

Construction of a genetic toggle switch in *Escherichia coli*

Timothy S. Gardner^{*†}, Charles R. Cantor^{* &} James J. Collins^{*†}

NATURE | VOL 403 | 20 JANUARY 2000 | www.nature.com

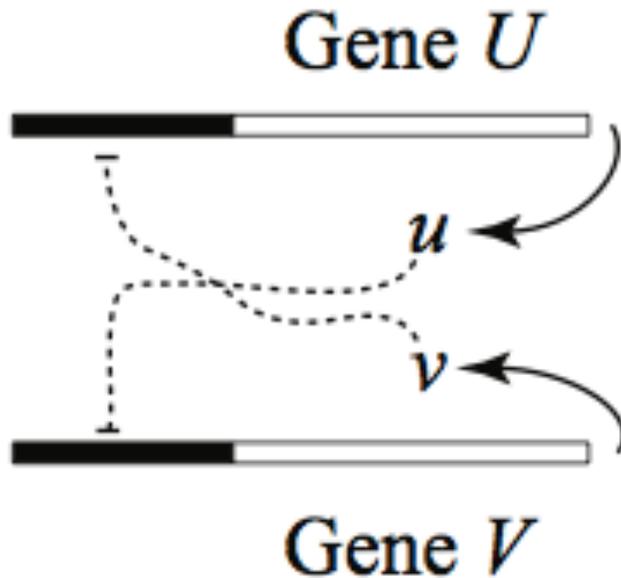
An actual “engineered genetic circuit” based on the concepts and models of biochemical switches.



Genetic toggle switch

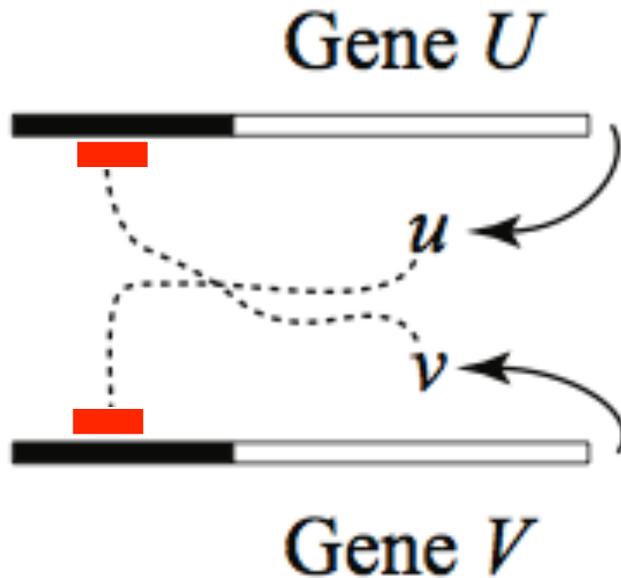
“Here we present the construction of a genetic toggle switch: a synthetic, bistable gene-regulatory network in *E. coli* and provide .. theory that predicts conditions for bistability.”

Production-decay of two proteins



$$\frac{du}{dt} = I_u - d_u u,$$
$$\frac{dv}{dt} = I_v - d_v v.$$

Negative feedback



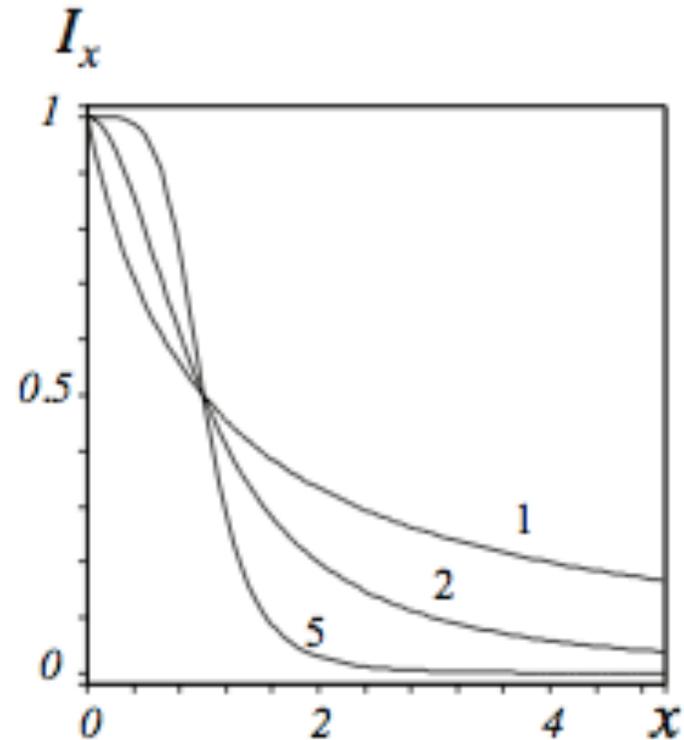
$$\frac{du}{dt} = I_u - d_u u,$$
$$\frac{dv}{dt} = I_v - d_v v.$$

$$I_x = \frac{\alpha}{1 + x^n}.$$

Negative feedback function

$$I_x = \frac{\alpha}{1 + x^n}$$

Higher n means
sharper response
with increasing x



Mutual inhibition

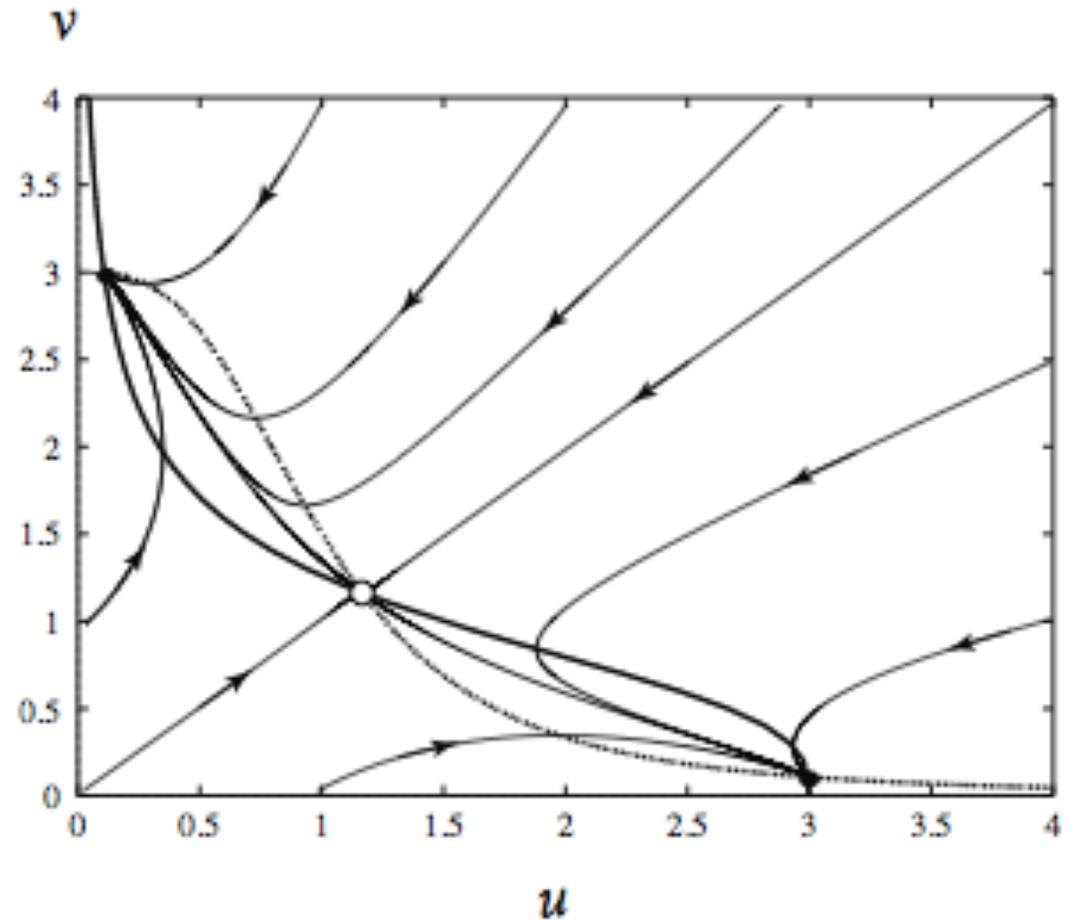
$$\frac{du}{dt} = \frac{\alpha_1}{1 + v^n} - u,$$
$$\frac{dv}{dt} = \frac{\alpha_2}{1 + u^m} - v.$$

Each gene product inhibits the other gene.

“... the toggle equations have 2 fundamental aspects: cooperative repression and degradation .. of the repressors”

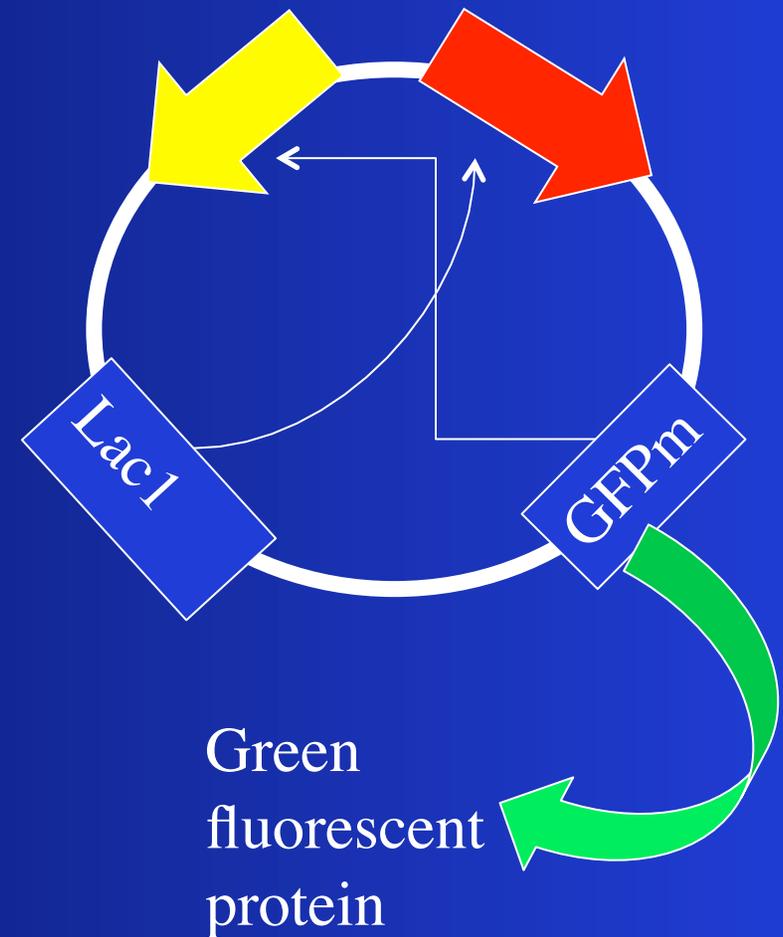
Switch-like behaviour

$$\frac{du}{dt} = \frac{\alpha_1}{1+v^n} - u,$$
$$\frac{dv}{dt} = \frac{\alpha_2}{1+u^m} - v.$$



Plasmid circuit

a synthetic, bistable
gene-regulatory network
in *E. coli*



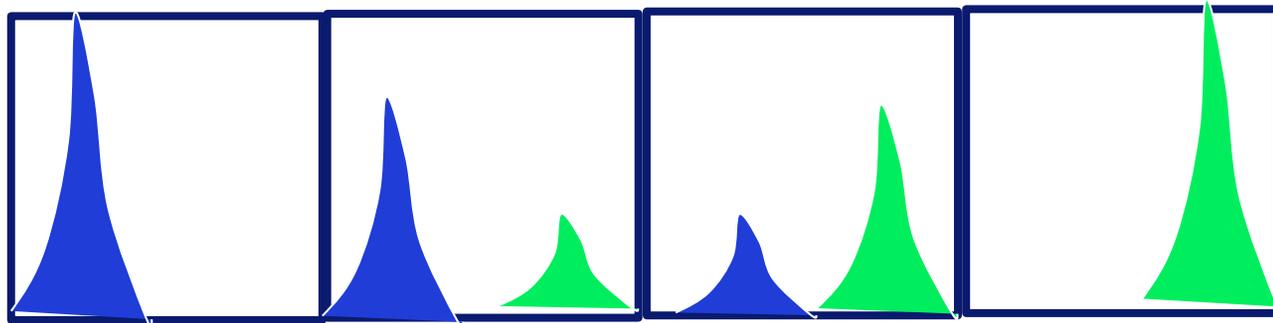
Cells switching can be induced

3hrs

4hrs

5hrs

6hrs



fluorescence

Mathematical Cell Biology Graduate Summer Course
University of British Columbia, May 1-31, 2012
Leah Edelstein-Keshet

Simple biochemical motifs (2.5)



www.math.ubc.ca/~keshet/MCB2012/

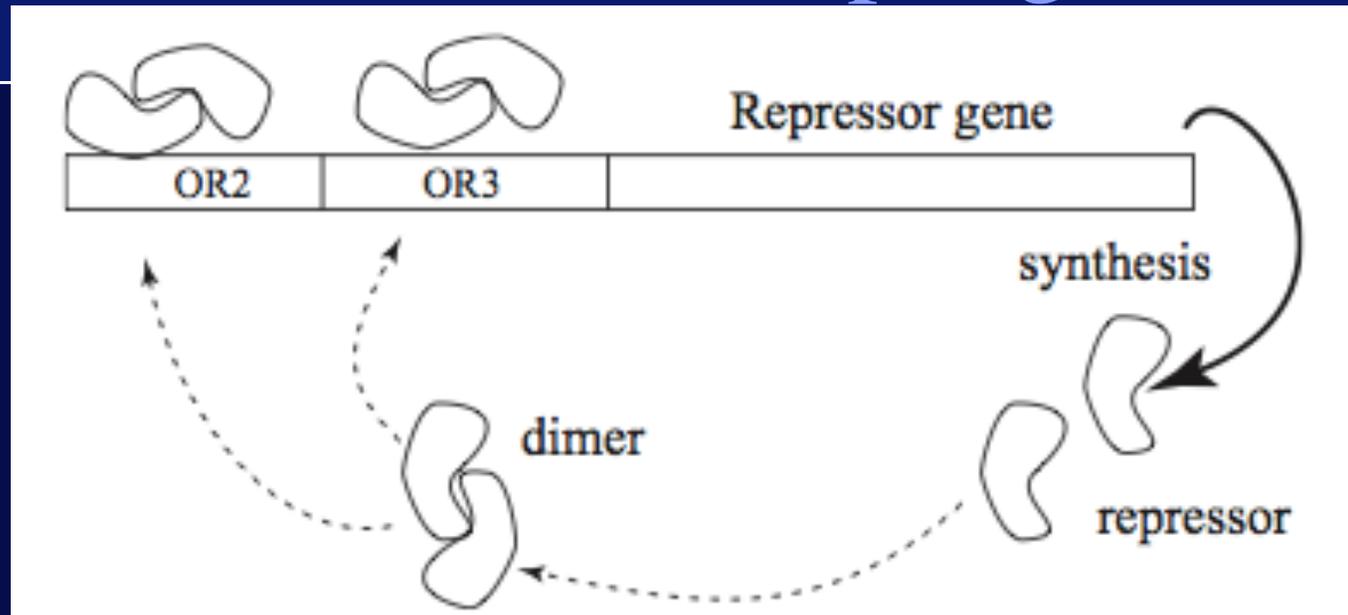


Noise-based switches and amplifiers for gene expression

Jeff Hasty^{*†}, Joel Pradines^{*}, Milos Dolnik^{**‡}, and J. J. Collins^{*}

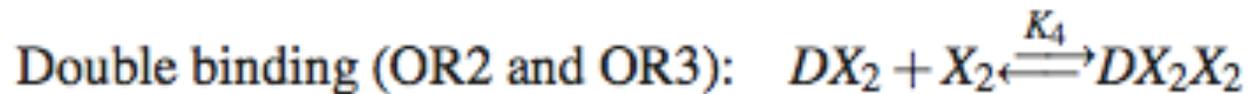
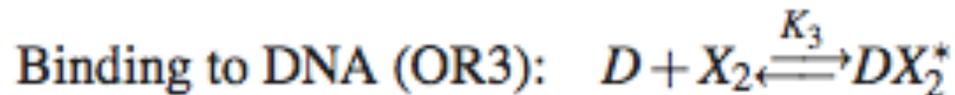
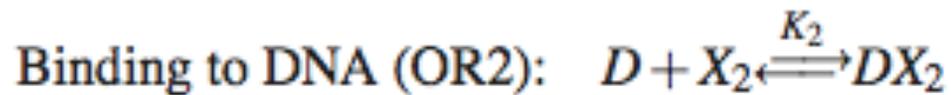
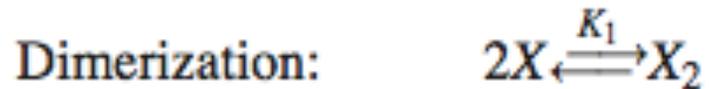
PNAS | February 29, 2000 | vol. 97 | no. 5 | 2075–2080

Dimerization and the phage lambda



- The phage λ gene encodes for protein (conc x)
- Protein dimerizes (conc of dimers y).
- Dimers bind to regulatory sites on the gene.
- Binding to OR2 activates transcription.
- Binding to OR3 inhibits transcription.

Reaction scheme



DX_2 = the dimerized repressor bound to site OR2

DX_2^ = the dimerized repressor bound to site OR3,*

DX_2X_2 = both OR2 and OR3 are bound by dimers

QSS

$$y = K_1 x^2,$$

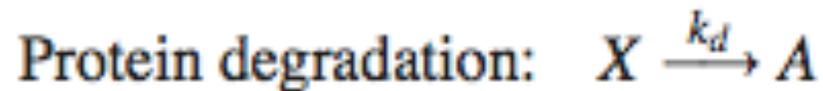
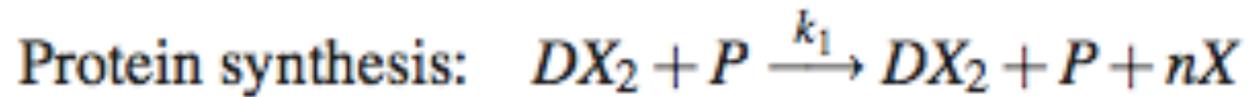
$$u = K_2 dy = K_1 K_2 dx^2,$$

$$v = \sigma_1 K_2 dy = \sigma_1 K_1 K_2 dx^2,$$

$$z = \sigma_2 K_2 uy = \sigma_2 (K_1 K_2)^2 dx^4.$$

The “fast variables” assumed to equilibrate rapidly with the variable x .

Slower timescale

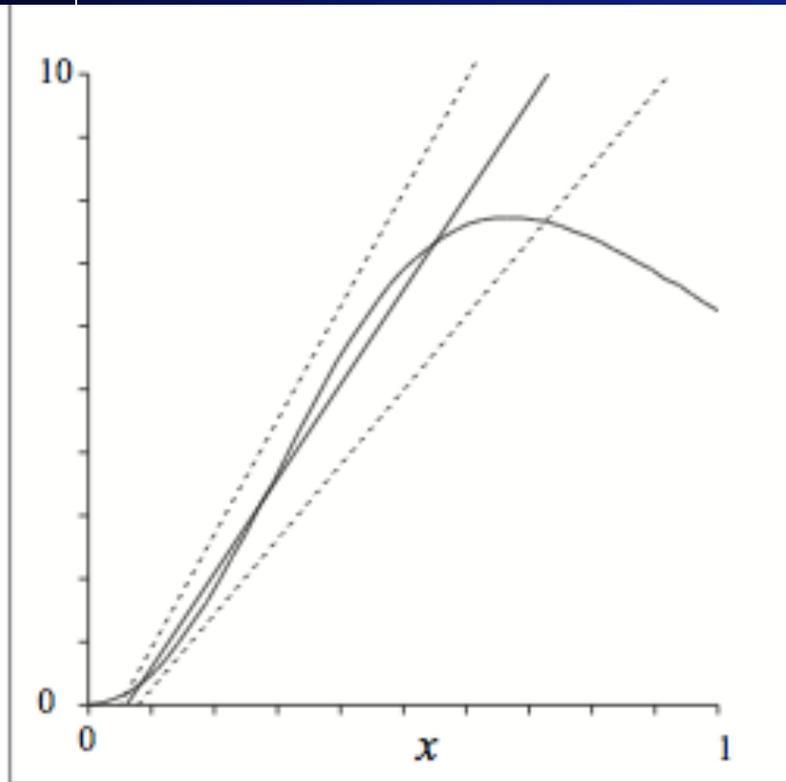


QSS and scaling the equations: system collapses to one variable, amt of synthesized protein, x :

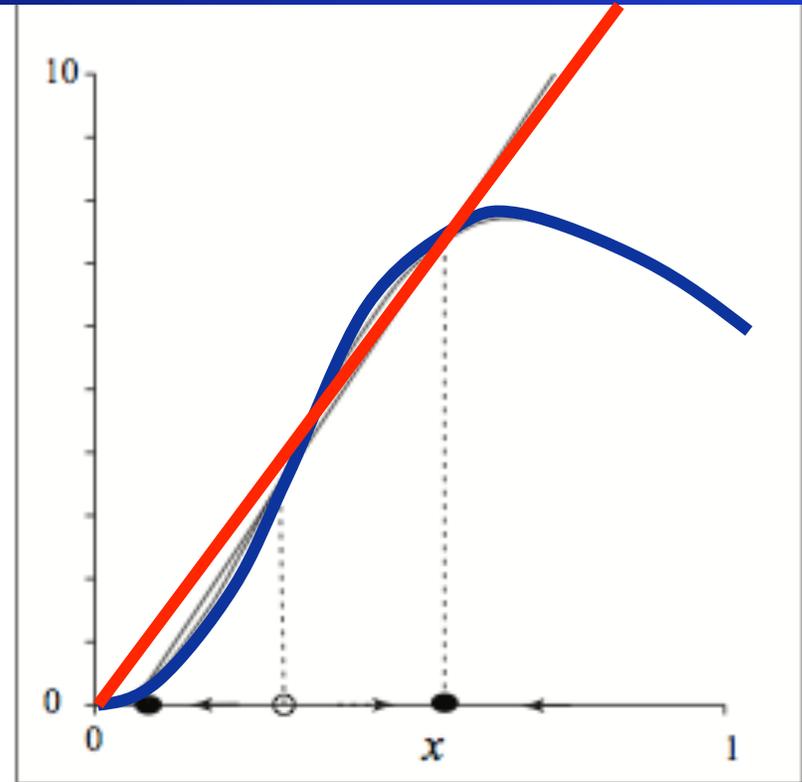
$$\frac{dx}{dt} = \frac{\alpha x^2}{1 + (1 + \sigma_1)x^2 + \sigma_2 x^4} - \gamma x + 1.$$

bistability

$$\frac{dx}{dt} = \frac{\alpha x^2}{1 + (1 + \sigma_1)x^2 + \sigma_2 x^4} - \gamma x + 1.$$



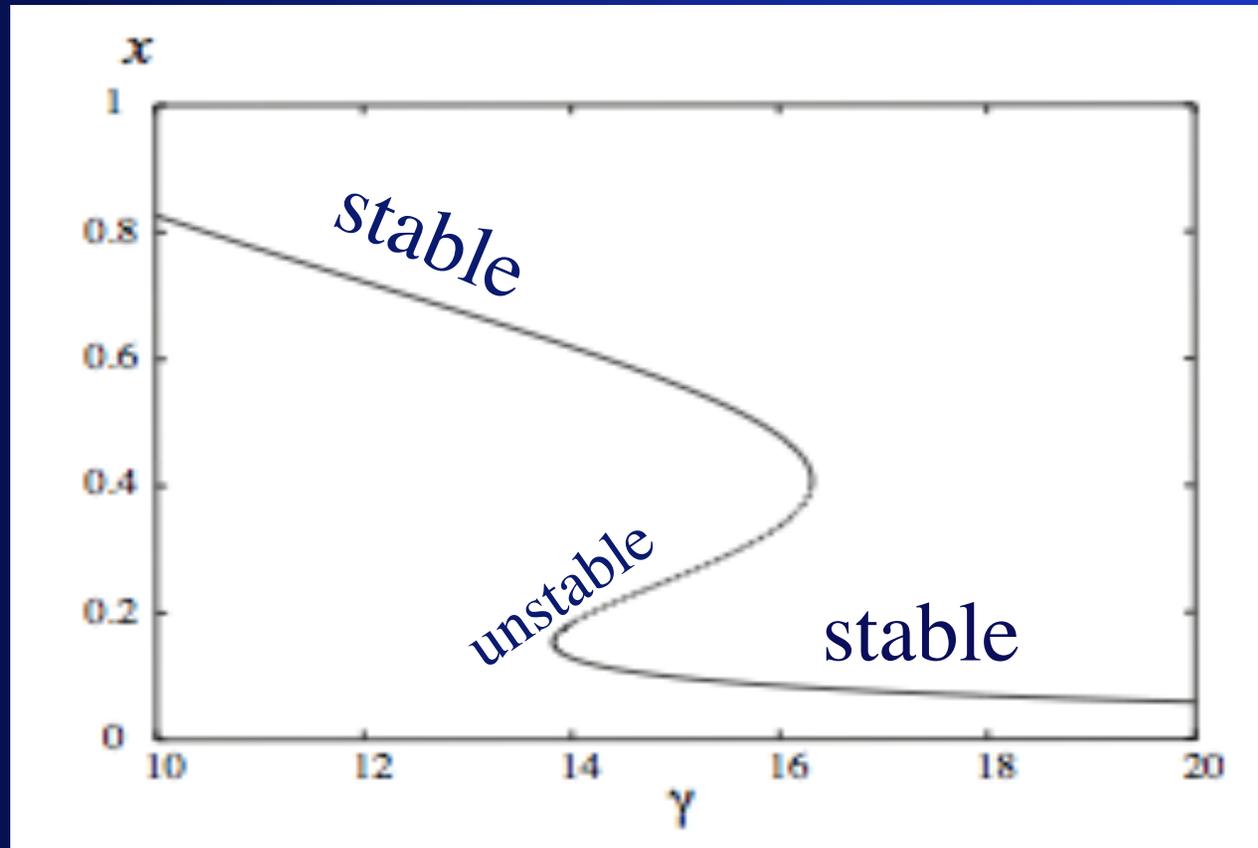
(a)



(b)

Bifurcation:

$$\frac{dx}{dt} = \frac{\alpha x^2}{1 + (1 + \sigma_1)x^2 + \sigma_2 x^4} - \gamma x + 1.$$



Comments

Combination of scaling, time scale considerations, and various simplifications can often reduce larger networks to effective dynamics of simpler systems.

Other examples will be provided.

Mathematical Cell Biology Graduate Summer Course
University of British Columbia, May 1-31, 2012
Leah Edelstein-Keshet

Simple biochemical motifs (3)

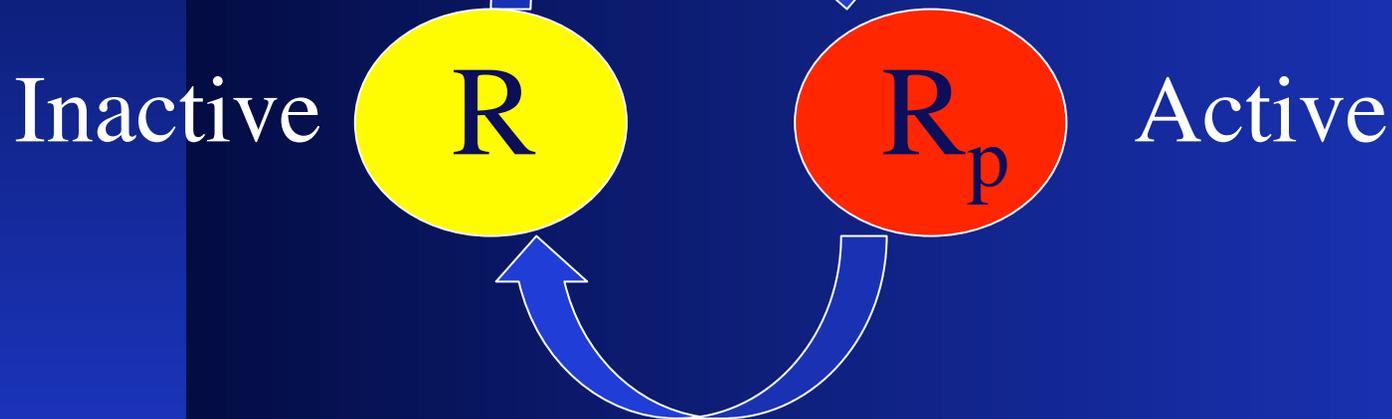


www.math.ubc.ca/~keshet/MCB2012/

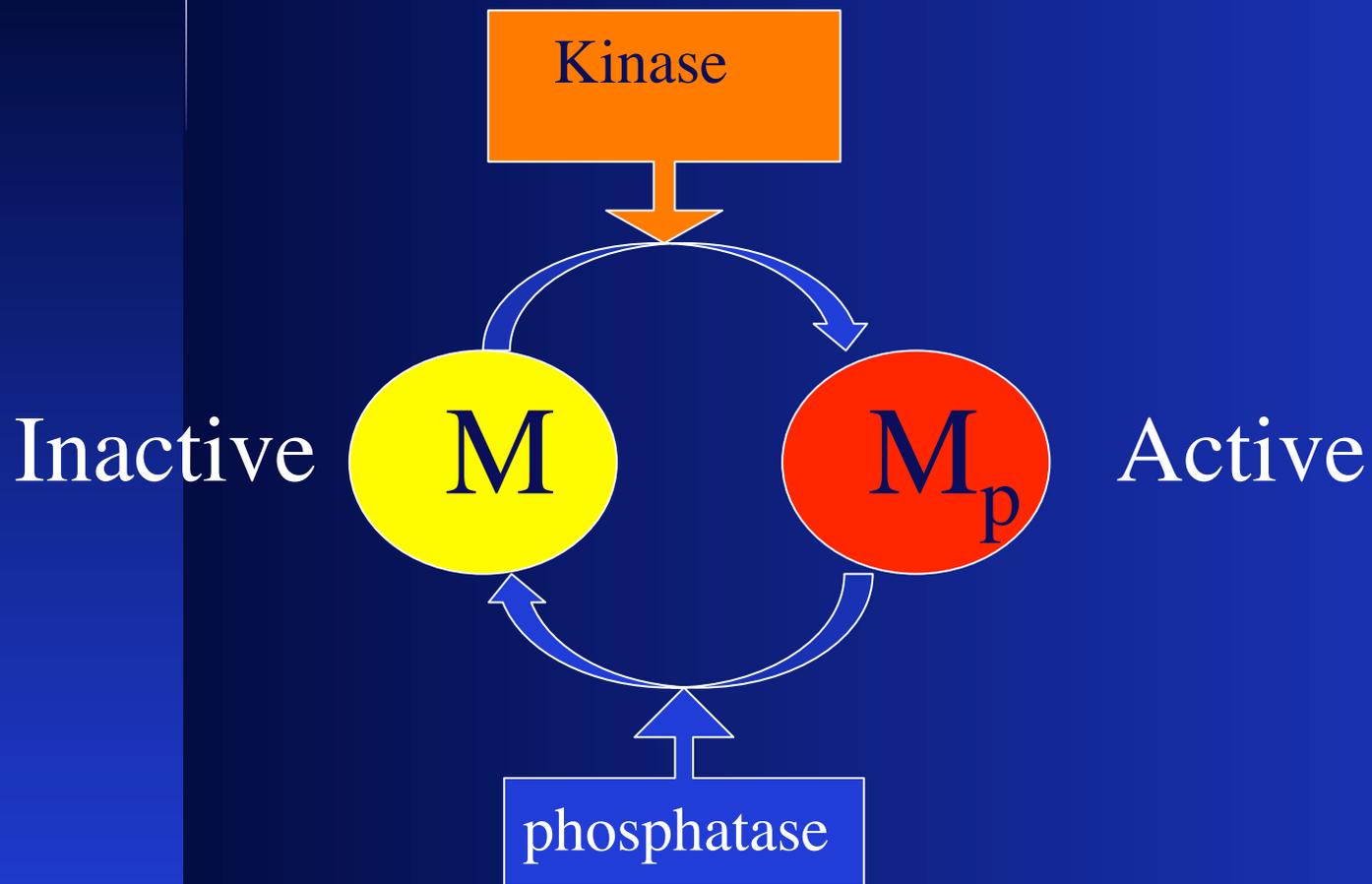
Activation-inactivation

$$\frac{dR_p}{dt} = \frac{k_1SR}{K_{m1} + R} - \frac{k_2R_p}{K_{m2} + R_p}$$

$$R_T = R + R_p = \text{constant.}$$

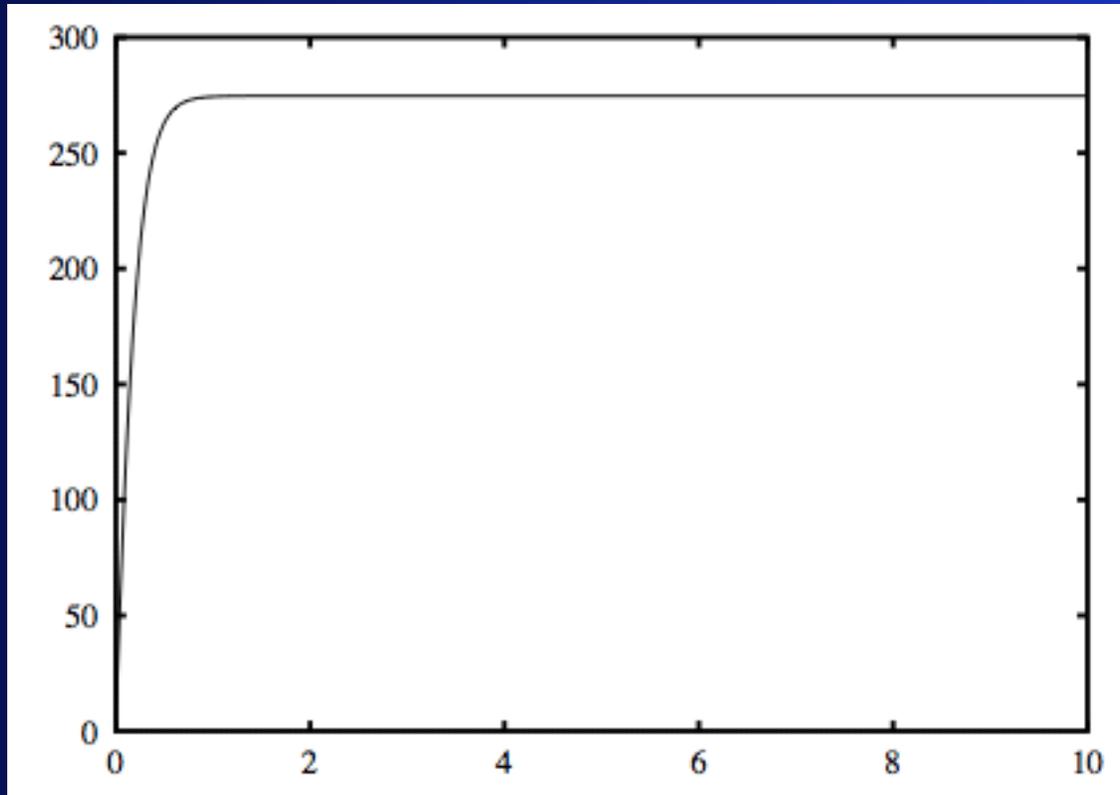


GTPase cycle



Without feedback: Fast equilibration

M_p



Time (seconds)

System has a single biologically relevant steady state

Eliminate R, rescale

$$\frac{dR_p}{dt} = \frac{k_1SR}{K_{m1} + R} - \frac{k_2R_p}{K_{m2} + R_p}$$

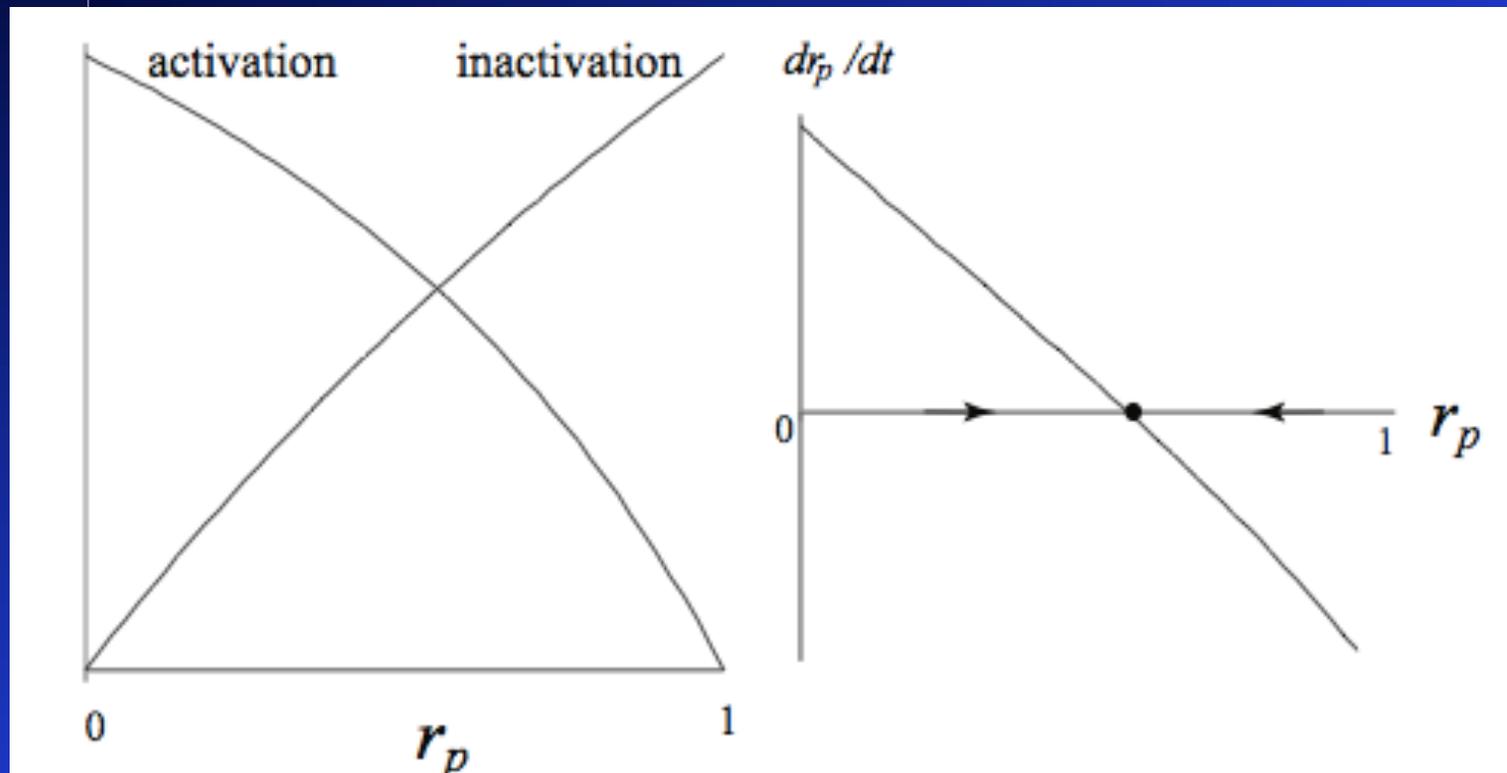
Use

$$R_T = R + R_p = \text{constant.}$$

$$r_p = R_p/R_T$$

Rescaled

$$\frac{dr_p}{dt} = \frac{k_1 S (1 - r_p)}{K'_{m1} + (1 - r_p)} - \frac{k_2 r_p}{K'_{m2} + r_p}$$



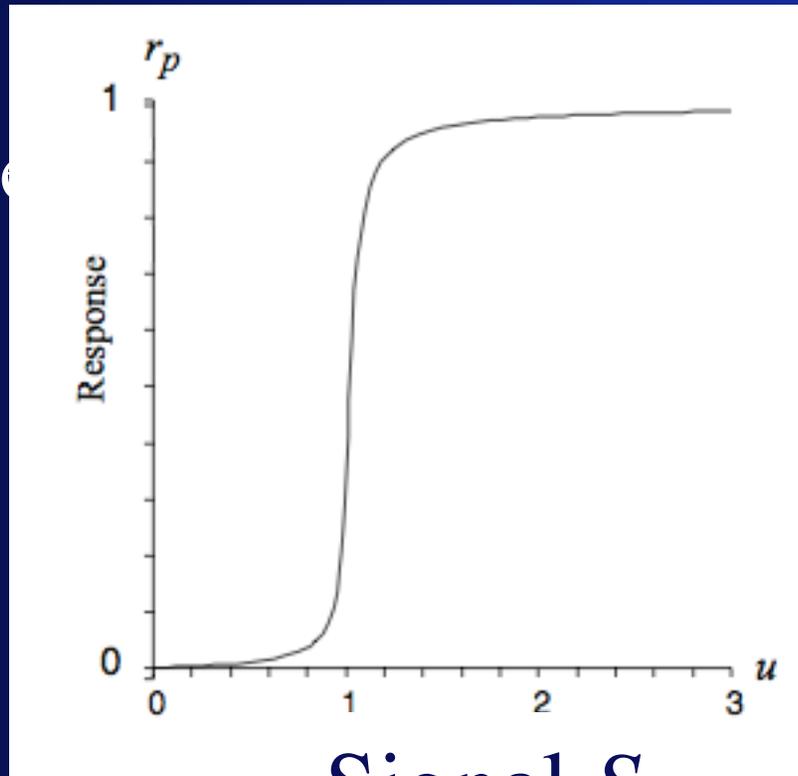
Steady states

$$\frac{dr_p}{dt} = \frac{k_1 S(1 - r_p)}{K'_{m1} + (1 - r_p)} - \frac{k_2 r_p}{K'_{m2} + r_p} = 0$$

The steady states can be shown to be solutions to a quadratic equation. Only one is positive and is called the “Goldbeter-Koshland function” of the stimulus.

“Zero order ultrasensitivity”

Steady
state
response



Signal S

response is minimal for low signal level, until some threshold. Then there is steep rise to full response. – *Goldbeter and Koshland*