

Last Time: Ability to oscillate determined by system properties

Description of the System
(Mass Balance/ODEs)

Taylor Approx.

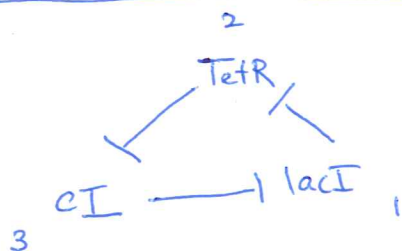
Jacobian

determinant

eigenvalues (λ)



oscillations possible
if imaginary part $\neq 0$
& real part > 0



$M = \text{mRNA}$
 $P = \text{protein}$

TetR Balance

$$\frac{dM}{dt} = \text{trxn} - \text{mRNA decay}$$

$$\frac{dP}{dt} = \text{trln} - \text{protein decay}$$

$$\Rightarrow \frac{dM_2}{dt} = k_2' + \frac{\beta_2 K_2^{n_2}}{K_2^{n_2} + P_1^{n_2}} - \underbrace{(\mu + k_{dm2})}_{\gamma} M_2$$

$$\frac{dP_2}{dt} = \alpha M_2 - \underbrace{(\mu + k_{dp2})}_{\gamma} P_2$$

simplify:

$$\frac{dM_2}{dt} = k_2' + \frac{\beta_2}{1 + \left(\frac{P_1}{K_2}\right)^{n_2}} - \gamma M_2$$

$$\frac{dP_2}{dt} = \alpha M_2 - \gamma P_2$$

Assume parameters for all 3 repressors the same
e.g. $n_1 = n_2 = n_3 = n$

$$\textcircled{1} \quad \frac{dM_2}{dt} = k' + \frac{\beta}{1 + \left(\frac{P_1}{K}\right)^n} - \gamma M_2$$

$$\textcircled{2} \quad \frac{dP_2}{dt} = \alpha M_2 - \gamma P_2$$

Let's non-dimensionalize

$$p_i \equiv \frac{P_i}{K}$$

\Rightarrow

$$P_i = P_i K$$

$$dP_i = K dp_i$$

Sub into ①

$$\frac{dM_2}{dt} = k' + \frac{\beta}{1+p_i^n} - \eta M_2$$

$$\frac{1}{\eta} \frac{dM_2}{dt} = \frac{k'}{\eta} + \frac{\beta}{\eta} \left(\frac{1}{1+p_i^n} \right) - M_2 \quad (1a)$$

$$\tau = \eta t \quad \left. \begin{array}{l} d\tau = \eta dt \end{array} \right\} \text{natural time scale}$$

Sub into ② $\eta \frac{K dp_2}{d\tau} = a M_2 - \gamma K p_2$

$$\frac{dp_2}{d\tau} = \frac{a}{\eta K} M_2 - \frac{\gamma}{\eta} p_2$$

$b \equiv$ ratio
of
protein to
mRNA decay

$$\frac{dp_2}{d\tau} = \left[\frac{a}{\eta K} \right] M_2 - b p_2$$

$a[\tau] = \frac{\text{protein}}{\text{mRNA}} \text{ s}$

$K[\tau] = \text{protein}$
 $\gamma[\tau] = \text{s}^{-1}$

$$m_2 = \left[\frac{a}{\eta K} \right] M_2$$

$\left[\frac{a}{\eta K} \right]$ translation efficiency

$$\Rightarrow \frac{dp_2}{d\tau} = b m_2 - b p_2$$

$$\left[\frac{dp_2}{d\tau} = -b(p_2 - m_2) \right]$$

$$dM_2 = \frac{K\gamma}{a} dm_2$$

sub into 1a

$$\Leftrightarrow K\gamma \frac{dm_2}{d\tau} = \frac{k'}{\eta} + \frac{\beta}{\eta} \left(\frac{1}{1+p_i^n} \right) - \frac{K\gamma}{a} m_2$$

$$\Rightarrow \frac{dm_2}{d\tau} = \alpha_0 + \frac{\alpha}{1+p_i^n} - m_2$$

where $\alpha_0 = \frac{k' a}{\eta K\gamma}$ $\alpha = \frac{\beta a}{\eta K\gamma}$

let $i = \begin{bmatrix} 1 = \text{lacI} \\ 2 = \text{tetR} \\ 3 = \text{cI} \end{bmatrix}$

$j = \begin{bmatrix} 1 = \text{cI} \\ 2 = \text{lacI} \\ 3 = \text{tetR} \end{bmatrix}$

(3)

$$\frac{dm_i}{dt} = d_0 + \frac{\alpha}{1+p_j^n} - m_i$$

$$\frac{dp_i}{dt} = -b(p_i - m_i)$$

Same as in paper

For simplicity, neglect mRNA dynamics

(much faster than protein)

$\Rightarrow \frac{dm_i}{dt} = 0 \Rightarrow m_i = d_0 + \frac{\alpha}{1+p_j^n}$

* Since parameters are all equal, steady state conc of p_i & m_i ~~then~~ same for all

$f = \frac{dp_1}{dt} = -bp_1 + b\alpha_0 + \frac{b\alpha}{1+p_3^n}$

$g = \frac{dp_2}{dt} = -bp_2 + b\alpha_0 + \frac{b\alpha}{1+p_1^n}$

$h = \frac{dp_3}{dt} = -bp_3 + b\alpha_0 + \frac{b\alpha}{1+p_2^n}$

Jacobian \Rightarrow

	p_1	p_2	p_3
f	$-b$	0	$b\alpha$
g	$b\alpha$	$-b$	0
h	0	$b\alpha$	$-b$

$\frac{\partial f}{\partial p_3} \bigg|_{ss} = \frac{-b\alpha n p_3^{n-1}}{(1+p_3^{ss})^2}$

$p_3^{ss} = p_2^{ss} = p_1^{ss} = p$

$X = \frac{-\alpha n p^{n-1}}{(1+p^n)^2}$

Stability determined by eigenvalues given by

(4)

$$\det \begin{bmatrix} -b-\lambda & 0 & bx \\ bx & -b-\lambda & 0 \\ 0 & bx & -b-\lambda \end{bmatrix} = 0$$

$$(-b-\lambda) [(-b-\lambda)^2 - 0] - 0 + (bx)^3 = 0$$

$$\lambda_1 = x - b$$

$$\lambda_2 = -b - \frac{1}{2}x - i\frac{\sqrt{3}}{2}x$$

$$\lambda_3 = -b - \frac{1}{2}x + i\frac{\sqrt{3}}{2}x$$

oscillates if
 $\lambda_I \neq 0$
 $\lambda_{Re} > 0$

$$x_1 < 0$$

$$\text{Re}(\lambda_2) \neq \text{Re}(\lambda_3)$$

$$\Rightarrow -b - \frac{1}{2}x > 0$$

$$x < -2b$$

system unstable & oscillations possible

when

$$\boxed{\frac{\alpha n p^{n-1}}{(1+p^n)^2} > 2b}$$

where
 $p = \frac{\alpha}{1+p^n} + \alpha_0$

Assume protein & mRNA decay rates comparable

$$\Rightarrow b = 1$$

$$\frac{\alpha n p^{n-1}}{(1+p^n)^2} > 2$$

Assume no leaky expression ($\alpha_0 = 0$)

$$p = \frac{\alpha}{1+p^n} \Rightarrow \alpha = p(1+p^n)$$

$$\begin{aligned} \frac{\alpha n p^{n-1}}{(1+p^n)^2} &= \frac{(p + p^{n+1}) n p^{n-1}}{(1+p^n)^2} \\ &= \frac{n p^n (1+p^n)}{(1+p^n)^2} \\ &= \frac{n p^n}{1+p^n} \end{aligned}$$

$$\Rightarrow \frac{np^n}{1+p^n} > 2$$

$$\frac{p^n}{1+p^n} \text{ is bounded b/c } 0 \leq 1$$

$$\therefore n(1) > 2 \Rightarrow n > 2$$

system must have
cooperativity, with no
leaky expression &
comparable decay
rates

What values of p will give this regime?

$$\frac{np^n}{1+p^n} > 2$$

$$\frac{p^n}{1+p^n} > \frac{2}{n}$$

$$p^n > \frac{2}{n} + \frac{2}{n} p^n$$

$$p^n > \frac{\frac{2}{n}}{1 - \frac{2}{n}} = \frac{2}{n-2}$$

$$p > \sqrt[n]{\frac{2}{n-2}} \gg 1 \quad [\text{remember } n \sim 2-4]$$

\therefore instability (no possible limit cycles)
for large p [high trxn/trln , strong
promoter & efficient
RBS]