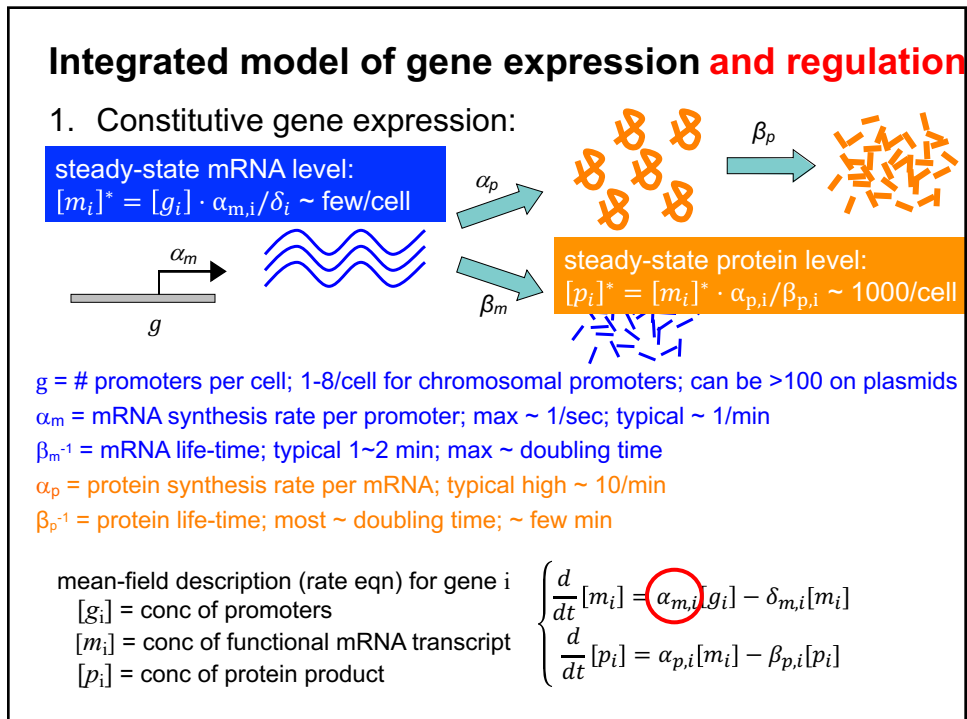


1



2

### transcriptional initiation and termination in bacteria

The diagram illustrates the steps of transcription in bacteria. It starts with RNA polymerase binding to the promoter region (upstream DNA) at the -35 region. This is followed by promoter melting to form an open complex. Initial transcription then begins, leading to elongation as the polymerase moves along the DNA. Termination occurs when the polymerase reaches a specific sequence, releasing the RNA transcript.

**tsx init control by activators, repressors**

**a** basal level of transcription: RNA polymerase binds to the promoter region (operator and promoter).

**b** no transcription: a repressor binds to the operator, blocking RNA polymerase.

**c** spontaneous isomerization leading to activated level of transcription: an activator binds to the activator binding site, facilitating RNA polymerase binding to the promoter.

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### Transcription regulation set molecularly by TF-TF and TF-RNAP interaction

$$\frac{d}{dt} [m_i] = \alpha_{m,i} [g_i] - \delta [m_i]$$

$$\frac{d}{dt} [p_i] = \alpha_{p,i} [m_i] - \beta [p_i]$$

steady state (exponential growth):

$$[m]^* = \alpha_{m,i} [g_i] / \beta \quad [p]^* = \frac{\alpha_{p,i} \alpha_{m,i} [g_i]}{\delta \beta}$$

$\alpha_{m,i} \propto \frac{1 + \omega ([A]/K_A)^n}{1 + ([A]/K_A)^n}$

(Hill function)

$$[p]^* = p_0 \frac{1 + \omega ([A]/K_A)^n}{1 + ([A]/K_A)^n}$$

growth-dependent

log-log slope ("sensitivity", n)

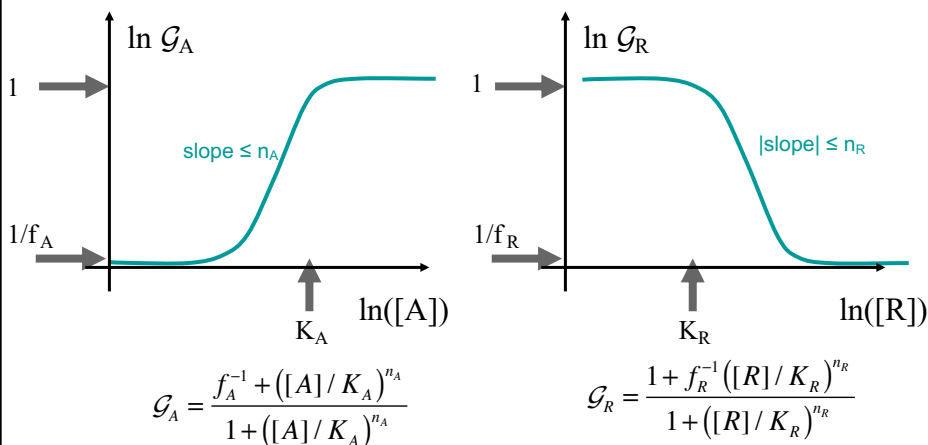
max fold change  $f$  ("capacity") --> set by  $\omega$

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### Simple circuit using transcriptional control

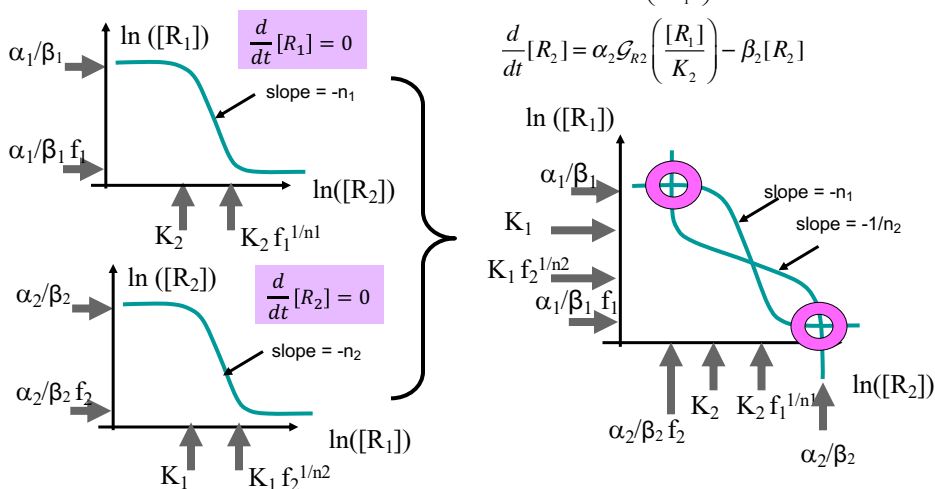
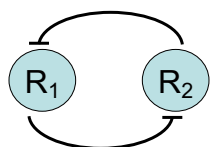
consider txs init control only (for simplicity)

$$\frac{d}{dt} [p_i] = \alpha_0 \cdot \mathcal{G} - \beta [p_i]$$



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### 3. Toggle switch



• qualitatively, two stable states:

- R1 on and R2 off
- R1 off and R2 on

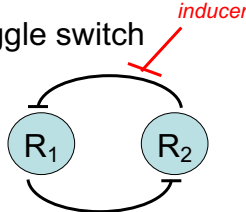
• quantitatively:

$$\frac{d}{dt} [R_1] = \alpha_1 \mathcal{G}_{R1} \left( \frac{[R_2]}{K_1} \right) - \beta_1 [R_1]$$

$$\frac{d}{dt} [R_2] = \alpha_2 \mathcal{G}_{R2} \left( \frac{[R_1]}{K_2} \right) - \beta_2 [R_2]$$

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### 3. Toggle switch



- qualitatively, two stable states:
  - R1 on and R2 off
  - R1 off and R1 on
- quantitatively:
 
$$\frac{d}{dt}[R_1] = \alpha_1 \mathcal{G}_{R_1} \left( \frac{[R_2]}{K_1} \right) - \beta_1 [R_1]$$

$$\frac{d}{dt}[R_2] = \alpha_2 \mathcal{G}_{R_2} \left( \frac{[R_1]}{K_2} \right) - \beta_2 [R_2]$$

• bistability favored by

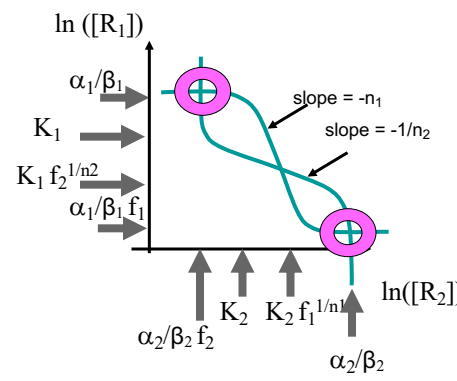
- $n_1 \cdot n_2 \gg 1$
- large  $f_1$  and  $f_2$
- appropriate ranges of  $K, \alpha, \beta$

• defining feature of bistability:

- remain in a state for a “long” time after initialized to it

• initialization by changing

- $K$  (via inducer)
- $\alpha$  (via activator/repressor)
- $\beta$  (via proteolysis, temperature)



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### Induction of TF

$$X + I \xrightleftharpoons[k_-]{k_+} XI$$

dissociation constant  $K_I = \frac{[X] \cdot [I]}{[XI]} = \frac{k_-}{k_+}$

$$[X]_{tot} = [X] + [XI] \quad [XI] = [X]_{tot} \frac{[I]}{[I] + K_I} \approx [X]_{tot} \frac{[I]_{tot}}{[I]_{tot} + K_I}$$

usually  $[I]_{tot} \gg [X]_{tot}$ , so  $[I] \approx [I]_{tot}$   
will drop the subscript "tot" from here on

“activated TF”  $X^*$  = form of TF able to bind specifically to DNA  
or able to activate RNAP

if  $X^* = XI$ , then  $[X^*] = [X]_{tot} \frac{[I]}{[I] + K_I}$  more generally, Hill form

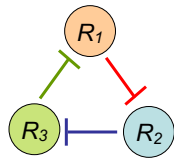
if  $X^* = X$ , then  $[X^*] = [X]_{tot} \frac{K_I}{[I] + K_I}$

$$[X^*] = [X]_{tot} / \left( 1 + \left( \frac{[I]}{K_I} \right)^{\pm n} \right)$$

or  $[X^*] = [X]_{tot} / \left( 1 + \left( \frac{[I]}{K_I} \right)^{\pm 1} \right)$

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### 4. Oscillators



### A synthetic oscillatory network of transcriptional regulators

Michael B. Elowitz & Stanislas Leibler  
NATURE | VOL 403 | 20 JANUARY 2000

“Repressilator”

- a.k.a. ring-oscillator
- uses only transcriptional repressors (with protein degradation tags)
- modeling gives oscillation for sufficiently cooperative repression

$$\frac{d[R_1]}{dt} = \alpha_1 \cdot \mathcal{G}_{R_1}([R_3]) - \beta_1 \cdot [R_1]$$

$$\frac{d[R_2]}{dt} = \alpha_2 \cdot \mathcal{G}_{R_2}([R_1]) - \beta_2 \cdot [R_2]$$

$$\frac{d[R_3]}{dt} = \alpha_3 \cdot \mathcal{G}_{R_3}([R_2]) - \beta_3 \cdot [R_3]$$

