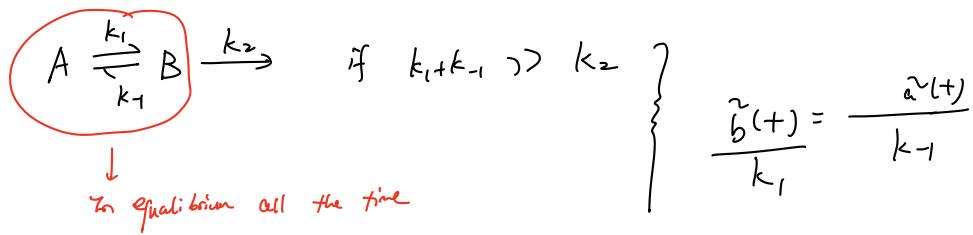




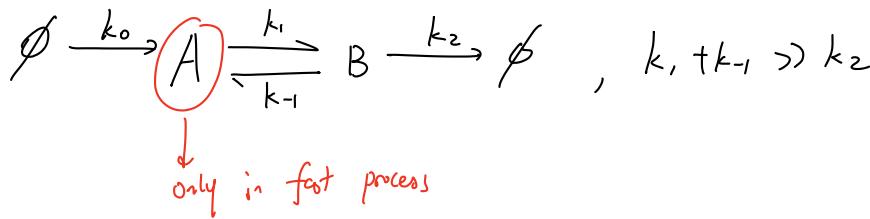
Rapid Equilibrium Assumption



$$\tilde{a}(+) + \tilde{b}(+) = \tilde{c}(+)$$

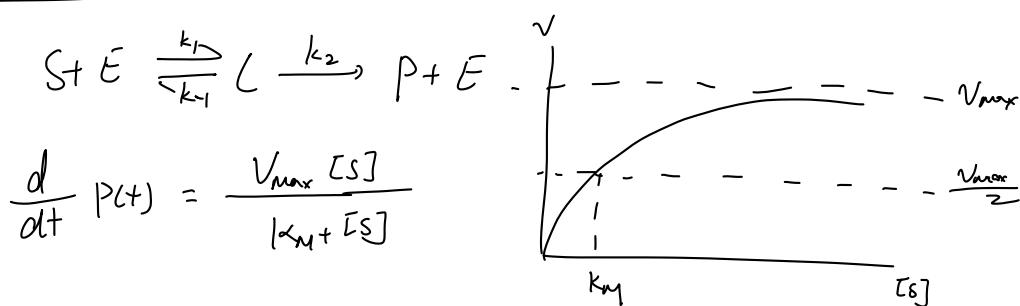
$$\Rightarrow \tilde{a}(+) = \left(\frac{k_1}{k_{-1} + k_1} \right) \tilde{c}(+) \quad \& \quad \tilde{b}(+) = \left(\frac{k_1}{k_{-1} + k_1} \right) \tilde{c}(+)$$

Quasi-Steady State Approximation



$$0 = k_0 - k_1 a^{ss} + k_{-1} b(+) \rightarrow \text{reduce } a(+) \text{ to } a^{ss}$$

Michaelis-Menton Kinetics



two substrates:

$$V = \frac{V_{max} [A] [B]}{K_{AB} + k_B [A] + k_A [B] + [A] [B]} \quad * V_{max} = k_2 e_T$$

Activation

Use the signalling molecule as enzyme:

$$V = \frac{k_1 [Signal] [S]}{k_2 + [S]}$$

Competitive Inhibition

$$V = \frac{V_{max} [S]}{K_m \left(1 + \frac{[I]}{K_i} \right) + [S]}$$

Allosteric Inhibition

$$V = \frac{V_{max}}{1 + \frac{[I]}{K_i}} \cdot \frac{\frac{S}{K_m + S}}{M1 - \text{kinetic}}$$

↓ inhibition

Cooperativity by Hill Function

$$T = \frac{[X]^h}{K^h + [X]^h} = \frac{1}{1 + \left(\frac{K}{[X]}\right)^h} \Rightarrow \begin{aligned} &\text{more cooperative for larger } h \\ &\hookrightarrow \text{also more sigmoidal} \end{aligned}$$

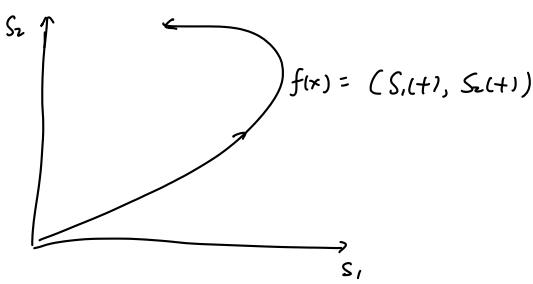
Diffusion

$$V = D ([S]_1 - [S]_2)$$

Facilitated Diffusion

$$V = \frac{\alpha_1 [S_1]/K_1 - \alpha_2 [S_2]/K_2}{1 + [S_1]/K_1 + [S_2]/K_2}$$

Phase plane



Nullclines

$$\begin{cases} 0 = f(S_1, S_2) = \frac{d}{dt} S_1(t) \\ 0 = g(S_1, S_2) = \frac{d}{dt} S_2(t) \end{cases}$$

↪ separate entire phase plane

↪ intersections are steady states

↪ stable or unstable

Stability Analysis

$$\text{For } \begin{cases} \frac{d}{dt} S_1 = f(S_1, S_2) \\ \frac{d}{dt} S_2 = g(S_1, S_2) \end{cases} \Rightarrow J = \begin{bmatrix} \frac{\partial f}{\partial S_1} & \frac{\partial f}{\partial S_2} \\ \frac{\partial g}{\partial S_1} & \frac{\partial g}{\partial S_2} \end{bmatrix}$$

If real parts of eigenvalues of J is:

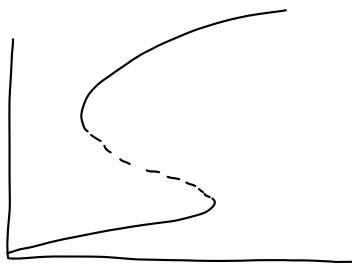
① all negative \Rightarrow stable steady states

② at least one positive \Rightarrow unstable steady states

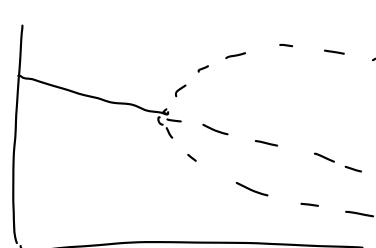
Bifurcation Analysis

plot: S.S. concentration vs. parameter value

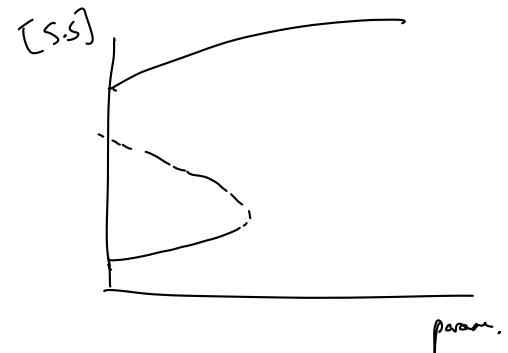
Saddle-node Bifurcation



Hopf Bifurcation



Inversible switch



Near bifurcation point: fragile; far from bifurcation point: robust

To a hysteretic switching is robust

Local Sensitivity Analysis

Absolute (local) parametric sensitivity:

$$\frac{d[\bar{s}]}{dp} \approx \frac{s(p+\Delta p) - s(p)}{\Delta p}$$

relative (local) sensitivity

$$\frac{p}{s} \frac{ds}{dp} \approx \left(\frac{p}{s}\right) \frac{s(p+\Delta p) - s(p)}{\Delta p}$$

Flux (J)

- rxn flux: steady state rxn. rate
- pathway flux: S.S. rxn rate through an unbranched chain
- flux control coefficient:

$$C_{ej}^J = \frac{e_j}{J_k} \frac{\partial J_k}{\partial e_j}$$

Summation Theorem

$$\sum_i C_{e_i}^J = 1$$

Ultrasensitivity

10% - 90% activation for < 8-fold increase in input

mRNA Expression Model

$$\frac{d}{dt} p(t) = \frac{k_i k_o}{g_m} - \delta_p p(t)$$

Rate of Activated Expression

$$V = \alpha_0 + \alpha \frac{[P]/k}{1 + [P]/k}$$

↑ transcription factor

Rate of Repressible Expression

$$V = \alpha_0 + \alpha \frac{1}{1 + [P]/k} \quad ; \quad V = \alpha_0 + \frac{\alpha}{1 + [P]/k}$$

for cooperativity

Multiple transcription factors

$$V = \alpha \left(\frac{\text{States with positive transcription}}{\text{total states}} \right)$$

$$\begin{aligned} O: & \frac{1}{1 + \frac{[A]}{K_A} + \frac{[B]}{K_B} + \frac{[AB]}{K_A K_B}} \\ OA: & \frac{[A]}{K_A} / \dots \\ OB: & \frac{[B]}{K_B} / \dots \\ OAB: & \frac{[AB]}{K_A K_B} / \dots \end{aligned}$$

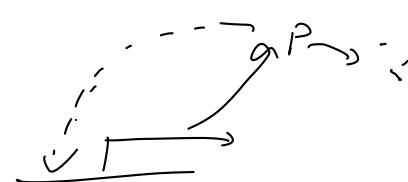
Autoinhibition

$$\frac{d}{dt} p(t) = \alpha \frac{1}{1 + p(t)/k} - \delta_p p(t)$$



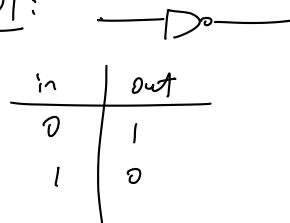
Autorepressor

$$\frac{d}{dt} p(t) = \alpha \frac{p(t)/k}{1 + p(t)/k} - \delta_p p(t)$$



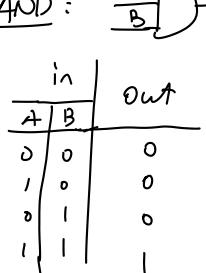
Logic Gates

NOT:



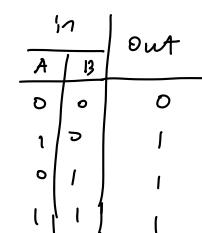
Repressor binding

AND:



Both activators binding
for transcription
(OAB state)

OR:



At least one activator binding
for transcription
(OA, OB, OAB)