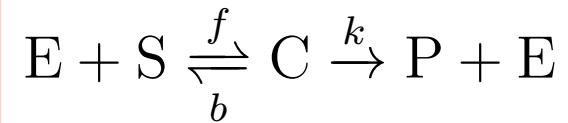


The Michaelis-Menten model of catalysis by enzymes

The standard way to model enzymes is with the Michaelis-Menten equation

In this model, an enzyme catalyses the *irreversible* conversion of substrate S to product P



Using the law of mass action

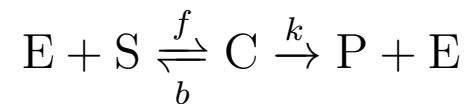
$$\frac{d[E]}{dt} = -f[E][S] + (b + k)[C]$$

$$\frac{d[S]}{dt} = -f[E][S] + b[C]$$

$$\frac{d[C]}{dt} = f[E][S] - (b + k)[C]$$

$$\frac{d[P]}{dt} = k[C].$$

We can simplify the equations because the reaction conserves enzymes



$$\frac{d[E]}{dt} = -f[E][S] + (b+k)[C]$$

$$\frac{d[S]}{dt} = -f[E][S] + b[C]$$

$$\frac{d[C]}{dt} = f[E][S] - (b+k)[C]$$

$$\frac{d[P]}{dt} = k[C].$$

implying

$$\frac{d[E]}{dt} + \frac{d[C]}{dt} = 0$$

and so

$$[E] + [C] = [E]_0 + [C]_0 = E_{\text{tot}}$$

the amount
of enzyme is
conserved

Most of the enzyme is bound up with substrate because there is so much more substrate compared to enzyme

quasi-steady state $\frac{d[C]}{dt} \simeq 0$ because the amount of enzyme is conserved

but

$$\frac{d[C]}{dt} = f[E][S] - (b + k)[C]$$

and so

$$f[E][S] = (b + k)[C] \quad \text{at quasi-steady state}$$

Using

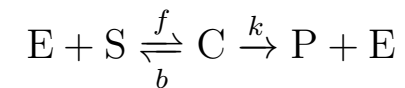
$$[E] = E_{\text{tot}} - [C]$$

we have

$$[C] \simeq \frac{E_{\text{tot}}[S]}{\frac{b+k}{f} + [S]} \quad \text{at quasi-steady state}$$

The Michaelis-Menten equation describes the rate of formation of the product

$$\frac{d[P]}{dt} = k[C] \quad \text{or} \quad \frac{d[P]}{dt} \simeq \frac{kE_{\text{tot}}[S]}{\frac{b+k}{f} + [S]} \quad \text{at quasi-steady state}$$



which is usually written as

$$\frac{d[P]}{dt} \simeq \frac{V_{\text{max}}[S]}{K_m + [S]}$$

with $V_{\text{max}} = kE_{\text{tot}}$; $K_m = \frac{b+k}{f}$

A more formal analysis shows that the assumption of quasi-steady state requires

$$\frac{E_{\text{tot}}}{[S]_0 + K_m} \ll 1$$

The Michaelis-Menten equation also describes the rate of consumption of the substrate

The substrate can be in only three states

$$\frac{d[S]}{dt} + \frac{d[C]}{dt} + \frac{d[P]}{dt} = 0$$

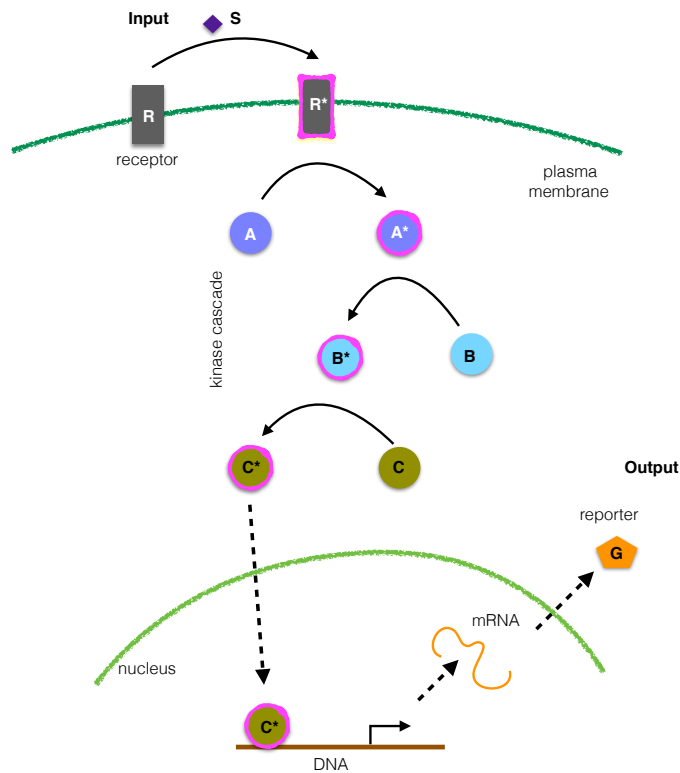
At quasi-steady state

$$\frac{d[S]}{dt} \simeq -\frac{d[P]}{dt}$$

implying

$$\frac{d[S]}{dt} \simeq -\frac{V_{\max}[S]}{K_m + [S]}$$

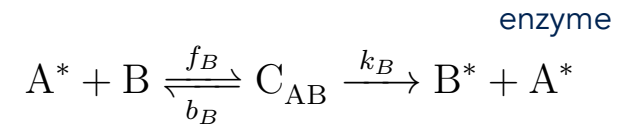
Modelling signal transduction IV.i



The rate equation for $[B^*]$

$$\frac{d[B^*]}{dt} \approx \frac{k_B[A^*][B]}{\frac{b_B+k_B}{f_B} + [B]} - \frac{k'_B[P][B^*]}{\frac{b'_B+k'_B}{f'_B} + [B^*]}$$

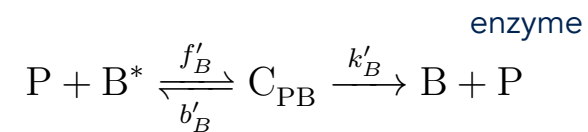
phosphorylation



The rate of change of $[B^*]$ has a positive term

$$\frac{k_B[A^*][B]}{\frac{b_B+k_B}{f_B} + [B]}$$

de-phosphorylation



The rate of change of $[B^*]$ has a negative term

$$-\frac{k'_B[P][B^*]}{\frac{b'_B+k'_B}{f'_B} + [B^*]}$$

Modelling signal transduction IV.ii

We will simplify the rate equation for $[B^*]$

$$\frac{d[B^*]}{dt} \simeq \frac{k_B[A^*](B_0 - [B^*])}{\frac{b_B+k_B}{f_B} + B_0 - [B^*]} - \frac{k'_B[P][B^*]}{\frac{b'_B+k'_B}{f'_B} + [B^*]} \quad \text{because} \quad B_0 = [B] + [B^*].$$

by assuming that the phosphatase is far from being saturated

$$[B^*] \ll \frac{b'_B+k'_B}{f'_B}$$

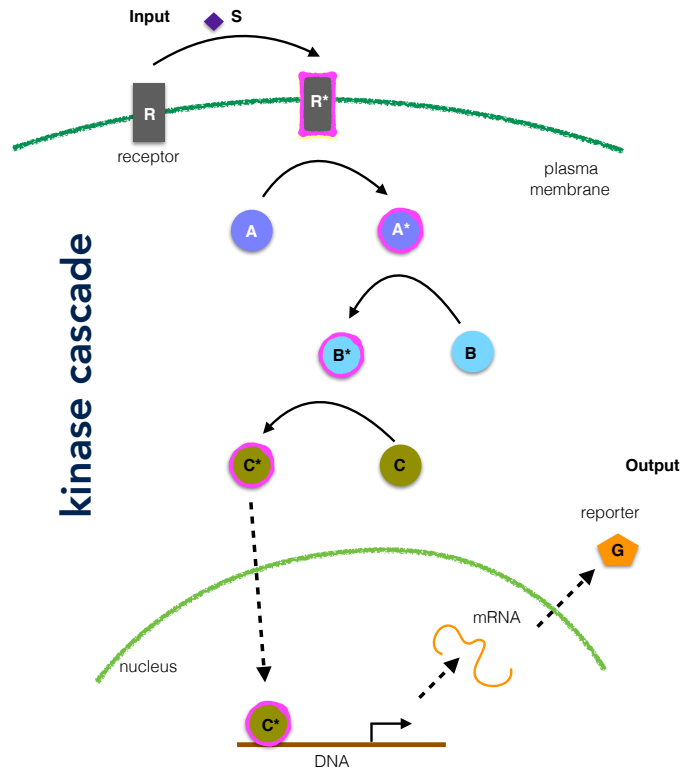
so that

$$\frac{d[B^*]}{dt} \simeq \frac{k_B[A^*](B_0 - [B^*])}{\frac{b_B+k_B}{f_B} + B_0 - [B^*]} - d_B[B^*]$$

with

$$d_B = \frac{f'_B k'_B [P]}{b'_B + k'_B}$$

Modelling signal transduction IV.iii



We have a complete model for the cytoplasmic part of the signal transduction

allosteric receptor

phosphatase far from saturation

$$\frac{d[A^*]}{dt} = \frac{k_A R_0 (1 + K^*[S])}{1 + K^*[S] + L(1 + K[S])} (A_0 - [A^*]) - d_A [A^*]$$

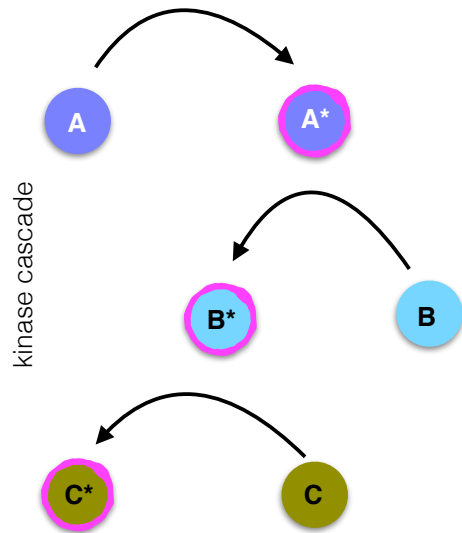
$$\frac{d[B^*]}{dt} = \frac{k_B [A^*] (B_0 - [B^*])}{\frac{b_B + k_B}{f_B} + B_0 - [B^*]} - d_B [B^*]$$

$$\frac{d[C^*]}{dt} = \frac{k_C [B^*] (C_0 - [C^*])}{\frac{b_C + k_C}{f_C} + C_0 - [C^*]} - d_C [C^*]$$

kinase cascade

Enzymatic cascades

Why have a cascade of kinases? Enzymatic cascades can increase ultrasensitivity



Active B is a Hill function of active A

$$[B^*] = [B^*]_{\max} \cdot \frac{[A^*]^{n_B}}{K_B^{n_B} + [A^*]^{n_B}} \quad \text{steady state}$$

Active C is a Hill function of active B

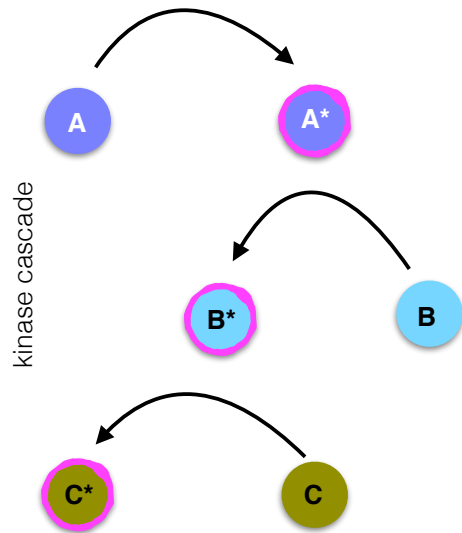
$$[C^*] = [C^*]_{\max} \cdot \frac{[B^*]^{n_C}}{K_C^{n_C} + [B^*]^{n_C}}$$

How does active C depend on active A ?

$$[C^*] = [C^*]_{\max} \cdot \frac{\left([B^*]_{\max} \frac{[A^*]^{n_B}}{K_B^{n_B} + [A^*]^{n_B}} \right)^{n_C}}{K_C^{n_C} + \left([B^*]_{\max} \frac{[A^*]^{n_B}}{K_B^{n_B} + [A^*]^{n_B}} \right)^{n_C}}$$

can we make this function approximate a Hill function?

Hill numbers multiply in a cascade: $n_{\text{final}} = n_B n_C$



Active B is a Hill function of active A

$$[B^*] = [B^*]_{\max} \cdot \frac{[A^*]^{n_B}}{K_B^{n_B} + [A^*]^{n_B}}$$

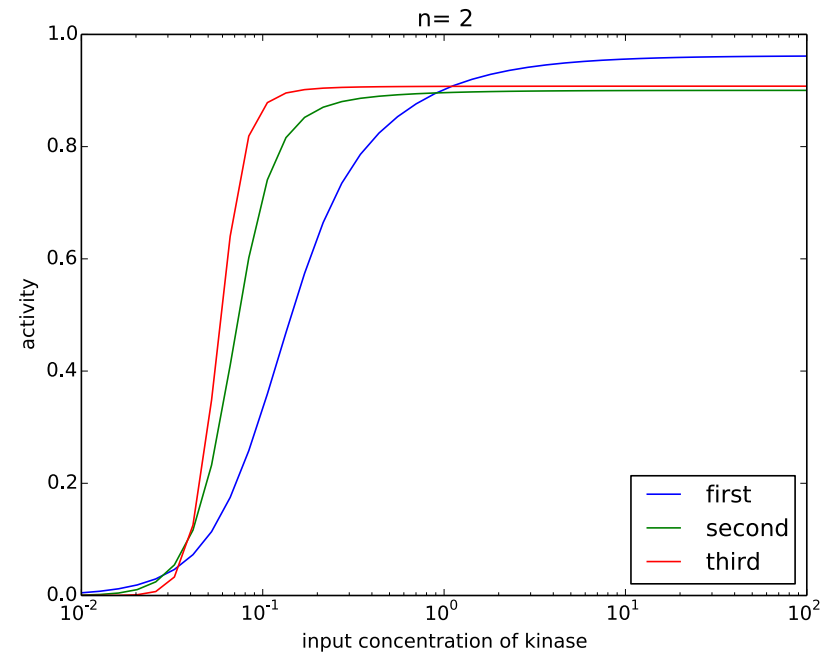
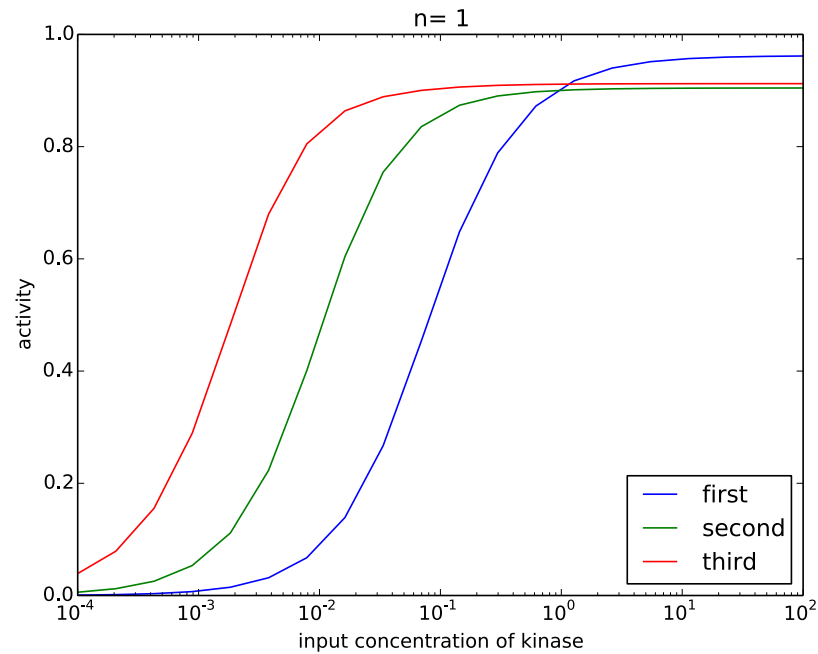
Active C is a Hill function of active B

$$[C^*] = [C^*]_{\max} \cdot \frac{[B^*]^{n_C}}{K_C^{n_C} + [B^*]^{n_C}}$$

How does active C depend on active A ?

$$[C^*] \simeq [C^*]_{\max} \cdot \frac{[A^*]^{n_B n_C}}{\frac{K_B^{n_B n_C} K_C^{n_C}}{[B^*]_{\max}^{n_C}} + [A^*]^{n_B n_C}} \quad \text{if} \quad [A^*] \ll K_B$$

For the cascade to increase sensitivity, the Hill numbers of intermediate steps must be greater than one



Many kinases require two phosphorylation to activate and so have a Hill number greater than one if the activating enzyme is distributive.



A **distributive** kinase binds, phosphorylates, dissociates, and then binds and phosphorylates again. A **processive** enzyme binds once, phosphorylates twice, and then dissociates.