

Synthesizing Units

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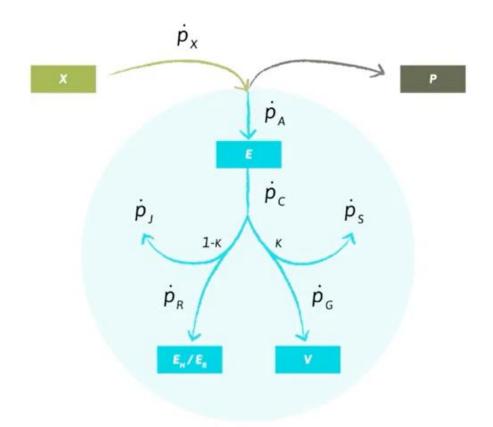






Conservation

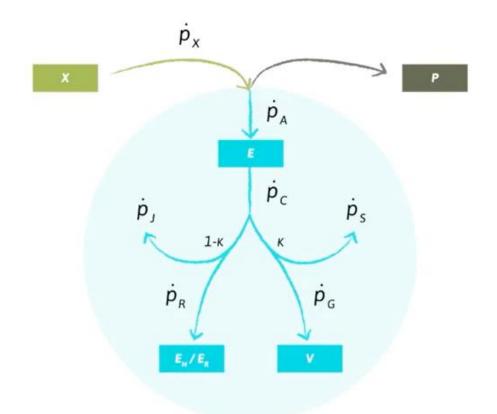
- Energy





Conservation

- Energy
- Mass





Conservation

- Energy
- Mass
- Time



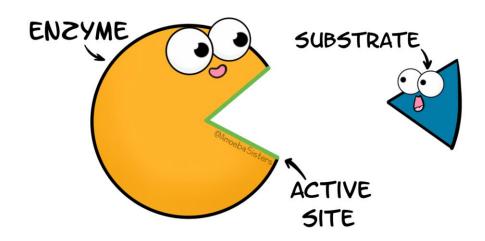
An entity that gets substrate(s) and processes them into product(s)

- Enzyme
- Organism
- Factory



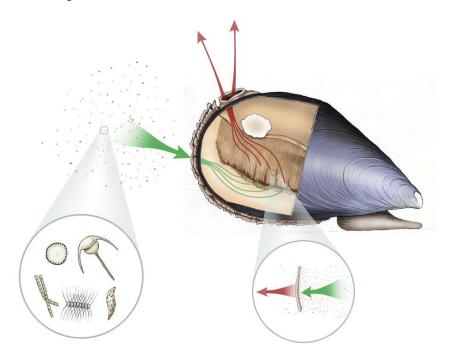
An entity that **gets** substrate(s) and processes them into product(s)

Passive





An entity that **gets** substrate(s) and processes them into product(s)



Passive

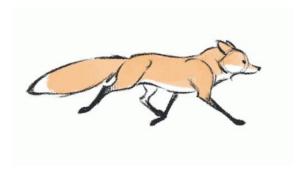




An entity that **gets** substrate(s) and processes them into product(s)

Passive



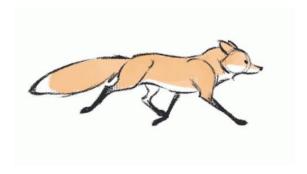




An entity that **gets** substrate(s) and processes them into product(s)

Passive



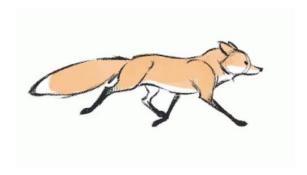




An entity that **gets** substrate(s) and processes them into product(s)

Passive

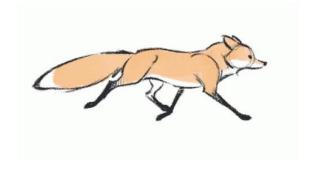






An entity that **gets** substrate(s) and processes them into product(s)







An entity that **gets** substrate(s) and processes them into product(s)





Unconstrained



An entity that **gets** substrate(s) and processes them into product(s)

Handshake protocols

- Open
- Closed



See section 7.1

Unconstrained



An entity that **gets** substrate(s) and **processes** them into product(s)

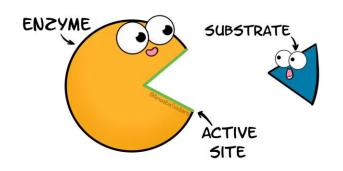
Handshake protocols for carriers

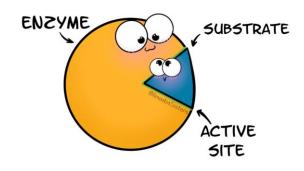
- Open
- Closed

See section 7.1



An entity that **gets** substrate(s) and **processes** them into product(s)





binding phase

prossessing phase



 $\theta_{.}$ $\theta_{.}$ proportion in binding

 $heta_S$ proportion in processing



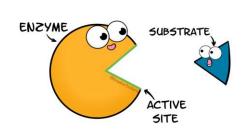
 $\theta_{.}$ $\theta_{.}$ proportion in binding

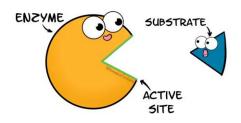
 $heta_{\mathcal{S}} \qquad \qquad heta_{\mathcal{S}} \ ext{proportion in processing}$

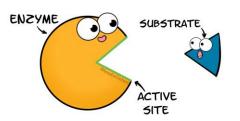
proportion of time of one SU in state of number of SUs in state

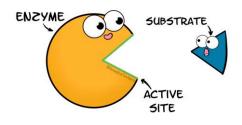


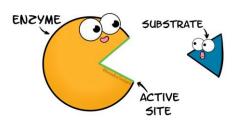
Examples of SUs



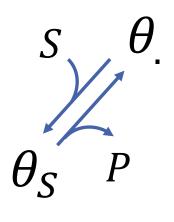












 $\theta_{.}$ proportion in binding

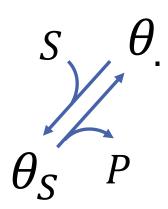
 $heta_{\mathcal{S}}$ proportion in processing

proportion

of time of one SU in state

of number of SUs in state

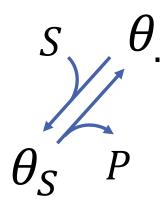




Conservation

$$\theta_{.} + \theta_{S} = 1$$



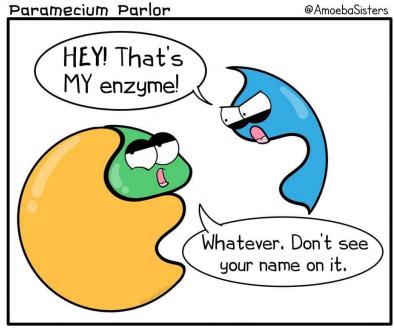


Conservation

$$\theta_1 + \theta_S = 1$$

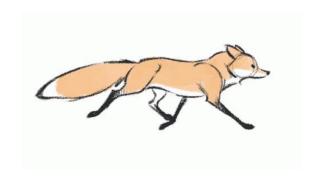
$$j_P = \left(\frac{1}{\dot{k}} + \frac{1}{\dot{b}S}\right)^{-1}$$



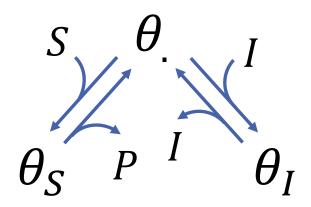


Competitive Inhibitors: If it fits, it sits.











$$\frac{d\theta}{dt} =$$

$$S \theta_{I} = \frac{d\theta_{S}}{dt} = \theta_{S} P \theta_{I} \frac{d\theta_{I}}{dt} = 0$$



$$\frac{d\theta}{dt} = -\dot{b}_S S\theta$$

$$\theta_{S} = \theta_{I}$$
 $\theta_{I} = \theta_{I}$

$$\frac{d\theta_S}{dt} = \dot{b}_S S \theta_S$$

$$\frac{d\theta_I}{dt} =$$



$$\frac{d\theta}{dt} = -\dot{b}_S S\theta + \dot{k}_S \theta_S$$

$$\theta_{S} = \frac{1}{P} \left[\frac{1}{P} \right]$$

$$\frac{d\theta_S}{dt} = \dot{b}_S S \theta_{\cdot} - \dot{k}_S \theta_S$$

$$\frac{d\theta_I}{dt} =$$



$$\frac{d\theta}{dt} = -\dot{b}_S S\theta + \dot{k}_S \theta_S - \dot{b}_I I\theta$$

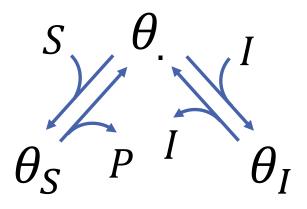
$$\theta_{S} = \theta_{I} = \theta_{I}$$

$$\frac{d\theta_S}{dt} = \dot{b}_S S \theta_{.} - \dot{k}_S \theta_S$$

$$\frac{d\theta_I}{dt} = \dot{b}_I I \theta$$



$$\frac{d\theta}{dt} = -\dot{b}_S S\theta + \dot{k}_S \theta_S - \dot{b}_I I\theta + \dot{k}_I \theta_I$$



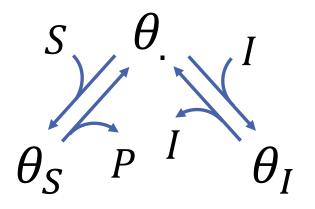
$$\frac{d\theta_S}{dt} = \dot{b}_S S \theta_{.} - \dot{k}_S \theta_S$$

$$\frac{d\theta_I}{dt} = \dot{b}_I I \theta_I - \dot{k}_I \theta_I$$



Fast dynamics assumption

$$0 = -\dot{b}_S S\theta_I + \dot{k}_S \theta_S - \dot{b}_I I\theta_I + \dot{k}_I \theta_I$$

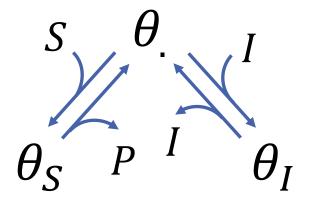


$$0 = \dot{b}_S S \theta_1 - \dot{k}_S \theta_S$$

$$0 = \dot{b}_I I \theta_I - \dot{k}_I \theta_I$$



$$\theta_I + \theta_S + \theta_I = 1$$

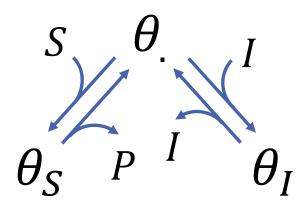


$$0 = \dot{b}_S S \theta_{\cdot} - \dot{k}_S \theta_S$$

$$0 = \dot{b}_I I \theta_I - \dot{k}_I \theta_I$$

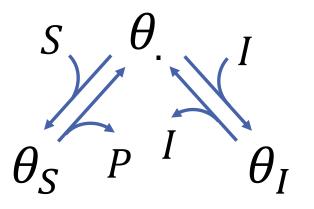


$$j_P = \dot{k}_S \theta_S$$





$$j_P = \dot{k}_S \theta_S = \left(\frac{1}{\dot{k}_S} + \frac{1}{\dot{b}S} \left(1 + \frac{\dot{b}_I}{\dot{k}_I}I\right)\right)^{-1}$$



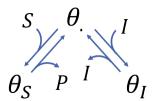
$$I=0$$

$$j_P = \left(\frac{1}{\dot{k}} + \frac{1}{\dot{b}S}\right)^{-1}$$



Step-by-step summary

1. Draw the scheme



- 2. Write the equations
- 3. Use the fast dynamics assumption
- 4. Add the conservation constraint
- 5. Compute the flux
- 6. Check consistency of the solution