BIOMENG 261

TISSUE AND BIOMOLECULAR ENGINEERING

Oliver Maclaren oliver.maclaren@auckland.ac.nz

MODULE OVERVIEW

Reaction kinetics and systems biology (Oliver Maclaren)

[11 lectures/3 tutorials/2 labs]

- 1. Basic principles: modelling with reaction kinetics [4 lectures]
 - Conservation, directional and constitutive principles. Mass action. Enzyme kinetics. Enzyme regulation. Mathematical/graphical tools for analysis and fitting.
- 2. Systems biology I: signalling and metabolic systems [2 lectures]

Overview of systems biology. Modelling signalling systems using reaction kinetics. Introduction to parameter estimation. Modelling metabolic systems using reaction kinetics. Flux balance analysis and constraint-based methods.

- 3. Systems biology II: genetic systems [3 lectures]
 - Modelling genes and gene regulation using reaction kinetics. Gene regulatory networks, transcriptomics and analysis of microarray data.

LECTURE 2 ENZYME KINETICS

- Enzymes
- The Michaelis-Menten model
- Quasi-equilibrium and quasi-steady-state analysis
- Fitting to data

ENZYMES: MOTIVATION/IDEA

$$A + B \longrightarrow C$$

might be

Possible (thermodynamically favourable)

but still

Too slow (kinetic 'path' takes too long)

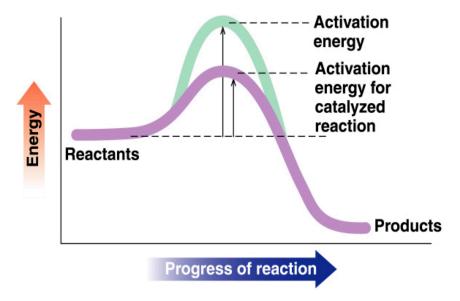
Q: Can we speed it up, without changing the start and end points?

CATALYSTS

A *catalyst* is a substance that

- Can speed up a reaction ('help along the reaction path')
- Doesn't change the start and end points (overall free energy is the same)
- Are not used up in the reaction themselves

CATALYSTS



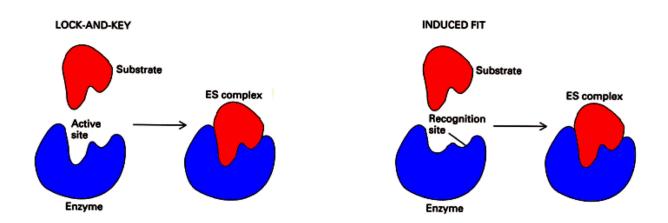
ENZYMES

An enzyme is a biological catalyst

- Usually proteins/large macromolecules
- Think: 'helper machines'
- Usually end in '-ase'
 - permease
 - kinase
 - etcase

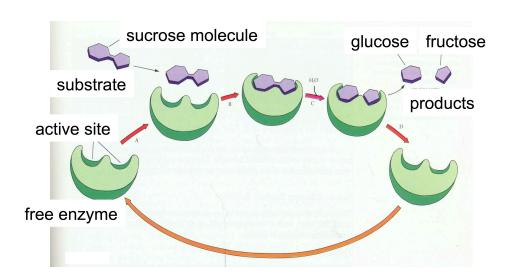
ENZYMES: MECHANISM(S)

- 'Lock and key': rigid enzyme
- 'Induced fit': same basic idea but deformable enzyme

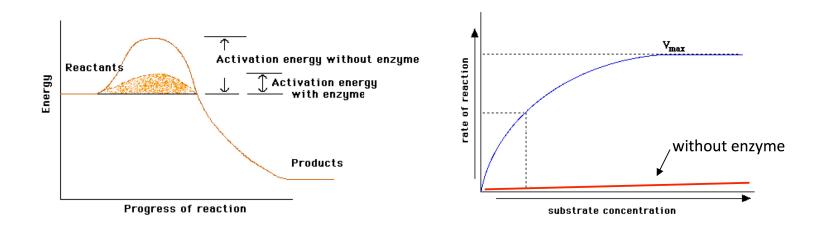


ENZYMES: EXAMPLE

Hydrolysis ('splitting') of sucrose into glucose and fructose



ENZYMES: EFFECT



Problem: *nonlinear* in 'substrate' (reactants)

But mass action would be rate = k[S], i.e. linear!

THE MICHAELIS-MENTEN MODEL

Michaelis-Menten (1913) introduced one of the first, and simplest, mathematical models of enzyme activity

Each step of full system obeys mass action

But result is

The overall, effective $S \rightarrow P$ reaction does not obey mass action

THE MICHAELIS-MENTEN MODEL: DERVIATION

Assumed reaction mechanism:

$$S+E \xrightarrow{k_1} C \xrightarrow{k_2} P+E$$

THE MICHAELIS-MENTEN MODEL: FULL MODEL

Full system of equations:

$$\frac{d[S]}{dt} = -k_1[E][S] + k_{-1}[C]$$

$$\frac{d[E]}{dt} = -k_1[E][S] + (k_{-1} + k_2)[C]$$

$$\frac{d[C]}{dt} = k_1[E][S] - (k_{-1} + k_2)[C]$$

$$\frac{d[P]}{dt} = k_2[C]$$

Initial conditions:

$$[S](0) = S_0, \quad [E](0) = E_0, \quad [C](0) = 0, \quad [P](0) = 0$$

THE MICHAELIS-MENTEN MODEL: REDUCED MODEL

Goal: reduction to 'effective' constitutive equation for

$$S \longrightarrow P$$

i.e.

$$J_P = \frac{d[P]}{dt} = f([S])$$

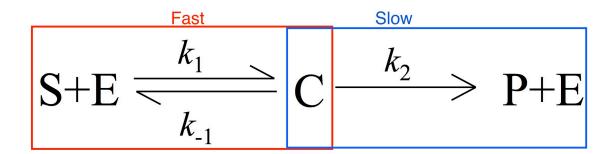
Note: people often use v instead of J in this context.

ANALYSIS METHODS

Recall: quasi-equilibrium vs quasi-steady-state

- Equilibrium: forward and backward components of a single reaction balanced
- Steady state: concentrations constant in time
 - multiple reactions into a particular compartment balance each other; may be unbalanced elsewhere

QUASI-EQUILIBRIUM ANALYSIS



• Assume fast reaction quickly reaches equilibrium.

QUASI-STEADY-STATE ANALYSIS

$$S+E \xrightarrow{linput \approx Output} k_2 \rightarrow P+E$$

Assume 'inputs and outputs' to [C] quickly reach balance.

UPSHOT: THE MICHAELIS-MENTEN CONSTITUTIVE EQUATION

Both result in the same form:

$$v([S]) := J_P([S]) = \frac{d[P]}{dt} = \frac{V_{max}[S]}{K_M + [S]}$$

with different K_M in terms of elementary steps - often just treat *empirically*, i.e. fit K_M . Notes:

• A *nonlinear* constitutive equation for what would usually be a linear mass action reaction.

NOTE: LARGE VS SMALL?

- Always compare quantities with the *same units*
- *Ratio* is then independent of units i.e. *dimensionless*

e.g. quasi-equilibrium compare:

$$\frac{k_2}{k_{-1}} \ll 1?$$

quasi-steady state compare:

$$\frac{E_0}{S_0} \ll 1?$$

Justification.

USE FOR FITTING DATA: LINEWEAVER-BURK PLOTS

To fit to data we can rewrite the Michaelis-Menten relation

$$v = \frac{V_{\mathsf{max}}[S]}{K_M + [S]}$$

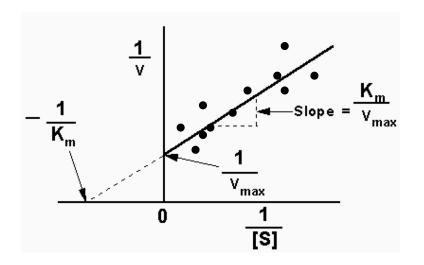
as

$$\frac{1}{v} = \frac{K_M}{V_{max}} \frac{1}{[S]} + \frac{1}{V_{max}}$$

i.e.
$$y = mx + c$$
 for $y = \frac{1}{v}$ and $x = \frac{1}{[S]}$.

USE FOR FITTING DATA: LINEWEAVER-BURK PLOTS

Called a Lineweaver-Burk plot.



Can also just fit original equation for a range of initial conditions.