Chemical reaction networks

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Overview

In biochemistry, 2+ species, or "reactants" can react if they come toegether and collide.

Alternatively, one species can degrade.

More is needed, though: correct orientation, enough energy, etc.

Examples

$$CH_4 + 2O_2 \longrightarrow CO_2 + 2H_2O$$
 (burning of methane)
 $H^+ + OH^- \longrightarrow H_2O$
unfolded protein \longrightarrow folded protein
 $2SO_2 + O_2 \Longrightarrow 2SO_3$
 $H_2O + CO_2 \Longrightarrow H_2CO_3$ (carbonic acid synthesis)
 $O_3 \longrightarrow O_2 + O$
 $2O_3 \longrightarrow 3O_2$

Classification of reactions:

- $\blacksquare A \longrightarrow P$: "uni-molecular"
- $A + B \longrightarrow P$: "bi-molecular"
- $A + B + C \longrightarrow P$: "tri-molecular"

Law of mass-action kinetics

A reaction rate is proportional to the probability of collision of reactants involved.

The probability of collisions is proportional to the concentration of each reactant R, denoted $\lceil R \rceil$.

If x is proportional to y, then this means that they differ by a constant k, i.e.,

$$x = ky$$
.

If x is proportional to y and z, then for some constant k,

$$x = kyz$$
.

Law of mass-action kinetics

A reaction rate is proportional to the concentrations of the the reactants.

ODE model

$$\blacksquare A \xrightarrow{k} P: \qquad \frac{d[P]}{dt} = k[A]$$

$$A + B \xrightarrow{k} P: \qquad \frac{d[P]}{dt} = k[A][B]$$

$$2A \xrightarrow{k} P: \frac{d[P]}{dt} = k[A]^2$$

■
$$2A \xrightarrow{k} P$$
:
$$\frac{d[P]}{dt} = k[A]^2$$
■ $A + B \xrightarrow{k_1} P$:
$$\frac{d[P]}{dt} = k_1[A][B] - k_2[P]$$

An example

Consider the following chemical reaction network:

$$A + B \rightleftharpoons_{k_2}^{k_1} C, \qquad A \xrightarrow{k_3} 2B$$

Let $x_1(t)$, $x_2(t)$, and $x_3(t)$ denote the concentrations of A, B, and C. Then we get the system

$$\frac{dx_1}{dt} = -k_1 x_1 x_2 + k_2 x_3 - k_3 x_1$$

$$\frac{dx_2}{dt} = -k_1 x_1 x_2 + k_2 x_3 + 2k_3 x_1$$

$$\frac{dx_3}{dt} = k_1 x_1 x_2 - k_2 x_3.$$

To find the steady-states, set each $x_i'=0$ and solve the system.

This can be found by computing a Gröbner basis of the ideal

$$I = (-k_1x_1x_2 + k_2x_3 - k_3x_1, -k_1x_1x_2 + k_2x_3 + 2k_3x_1, k_1x_1x_2 - k_2x_3).$$

An example

Consider the following chemical reaction network:

$$A + B \rightleftharpoons_{k_2} C, \qquad A \xrightarrow{k_3} 2B,$$

and the resulting system of nonlinear ODEs:

$$\begin{aligned} \frac{dx_1}{dt} &= -k_1 x_1 x_2 + k_2 x_3 - k_3 x_1 \\ \frac{dx_2}{dt} &= -k_1 x_1 x_2 + k_2 x_3 + 2k_3 x_1 \\ \frac{dx_3}{dt} &= k_1 x_1 x_2 - k_2 x_3. \end{aligned}$$

Questions

- Does the system have a positive or non-negative equilibrium?
- Does the system have *multiple* positive equlibria?
- Does the system have a *stable* positive equillbrium?
- Does the system have an unstable positive equilibrium?
- Do all positive species concentrations admit a positive cyclic composition trajectory?
- How do the answers to these questions depend on the rate constants?

Enzymes are proteins that catalyze reactions (up to 10¹²-fold!)

An example

Consider the following chemical reaction

$$E + S \xrightarrow{k_1} ES \xrightarrow{k_3} E + P$$

 $E= {\sf enzyme}, \ S= {\sf substrate}, \ ES= {\sf enzyme}{\sf -substrate} \ {\sf complex}, \ {\sf and} \ P= {\sf product}.$

$$\begin{cases} \frac{d[ES]}{dt} = k_1[E][S] - (k_2 + k_3)[ES] \\ \\ \frac{d[P]}{dt} = k_3[ES] \\ \\ E_0 = [E] + [ES], \qquad E_0 = \text{ initial enzyme concentration} \end{cases}$$

Assumptions

- \blacksquare E_0 is constant.
- Enzyme-substrate complex reaches equilibrium much earlier than the product does, so $\frac{d[ES]}{dt} \approx 0$.

Goal

Write the differential equation $\frac{d[P]}{dt} = k_3[ES]$ in terms of [S], not [ES].

Since $\frac{d[ES]}{dt} \approx 0$, we can simplify the ODE for [ES]:

$$\frac{d[ES]}{dt} = k_1[E][S] - (k_2 + k_3)[ES] = 0.$$

Upon solving for [E], we get

$$[E] = \frac{(k_2 + k_3)[ES]}{k_1[S]}.$$

Plugging this into $E_0 = [E] + [ES]$ and solving for [ES]:

$$[ES] = \frac{E_0[S]}{\frac{k_2 + k_3}{k_3} + [S]}.$$

Alas, we can write

$$\frac{d[P]}{dt} = k_3[ES] = \frac{k_3 E_0[S]}{\frac{k_2 + k_3}{k_1} + [S]} = \frac{V_{\text{max}}[S]}{K_m + [S]}.$$

Michaelis-Menten equation

Recall the following chemical reaction:

$$E + S \xrightarrow{k_1} ES \xrightarrow{k_3} E + P$$

E= enzyme, S= substrate, ES= enzyme-substrate complex, and P= product.

Definition

The Michaelis-Menten equation is one of the best-known models of enzyme kinetics.

$$\frac{d[P]}{dt} = \underbrace{\frac{V_{\text{max}}[S]}{K_m + [S]}}_{f([S])}, \quad \text{where } V_{\text{max}} = k_3 E_0, \quad \text{and } K_m = \frac{k_2 + k_3}{k_1}$$

Remarks

- The "reaction rate", f([S]), is a strictly increasing function of [S].
- $\lim_{|S| \to \infty} f([S]) = V_{\max}$, (biologically, the maximum reaction rate)
- $f(K_m) = \frac{1}{2}V_{\text{max}}.$
- The reaction rate f([S]) is proportional to E_0 .

Michaelis-Menten equation

Recall the following chemical reaction:

$$E + S \xrightarrow{k_2} ES \xrightarrow{k_3} E + P$$

E= enzyme, S= substrate, ES= enzyme-substrate complex, and P= product.

Further assumptions

- Substrate concentration is conserved: $S_0 = [S] + [ES] + [P]$.
- $E_0 \ll S_0$, so $[ES] \ll [S]$ and [P].

Together, this means $S_0 \approx [S] + [P]$. Taking $\frac{d}{dt}$ of both sides yields

$$\frac{d[S]}{dt} = -\frac{d[P]}{dt} = -\frac{V_{\text{max}}[S]}{K_m + [S]} \; . \label{eq:definition}$$

Usually, V_{max} , K_m , and S_0 are known quanities. This is now something we can easily solve, graph, analyze, etc.

Multi-molecule binding

Consider a reaction where n molecules of a substrate S react with an enzyme E:

$$E + nS \stackrel{k_1}{\rightleftharpoons} ES_n \stackrel{k_3}{\longrightarrow} E + P$$

The enzyme-substrate complex here is ES_n . By mass-action kinetics,

$$\begin{cases} \frac{d[ES_n]}{dt} = k_1[E][S]^n - (k_2 + k_3)[ES_n] \\ \\ \frac{d[P]}{dt} = k_3[ES_n] \\ \\ E_0 = [E] + [ES_n], \qquad E_0 = \text{ initial enzyme concentration} \end{cases}$$

As before, assume $[ES_n]$ reaches equilibrium much quicker than [P] and [S]:

$$\frac{d[ES_n]}{dt} = 0 \qquad \Longrightarrow \qquad [E] = \frac{(k_2 + k_3)[ES_n]}{k_1[S]^n}.$$

Plugging this into $E_0 = [E] + [ES_n]$ and solving for $[ES_n]$ yields

$$[ES_n] = \frac{E_0[S]^n}{\frac{k_2 + k_3}{k_1} + [S]^n} \qquad \Longrightarrow \qquad \boxed{\frac{d[P]}{dt} = \frac{V_{\max}[S]^n}{K_m + [S]^n}}.$$

Multi-molecule binding

Hill equation

Given the chemical reaction

$$E + nS \stackrel{k_2}{\underset{k_1}{\rightleftharpoons}} ES_n \stackrel{k_3}{\longrightarrow} E + F$$

We derived the following ODE involving [P] and [S]:

$$\frac{d[P]}{dt} = \underbrace{\frac{V_{\max}[S]^n}{K_m + [S]^n}}_{f([S])}, \qquad \text{where } V_{\max} = k_3 E_0, \quad \text{and } K_m = \frac{k_2 + k_3}{k_1}$$

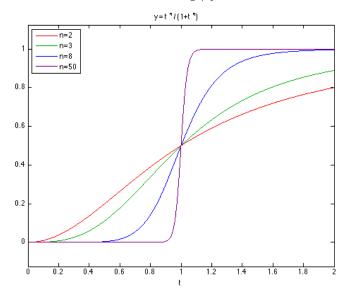
This is called the Hill equation with Hill coefficient n.

Remarks

- The "reaction rate", f([S]), is a strictly increasing function of [S].
- $\blacksquare \lim_{|S| \to \infty} f([S]) = V_{\max}, \quad \text{(biologically, the maximum reaction rate)}$
- $f(K_m^{1/n}) = \frac{1}{2}V_{\text{max}}.$
- The reaction rate f([S]) is proportional to E_0 .
- $\mathbf{n} = 1$ is just the Michaelis-Menden equation.

Hill equations

The following shows several "Hill functions" $y = \frac{t^n}{1 + t^n}$, for various values of n.



Michaelis-Menten variants

We just saw an equation for where 1 molecule of a substrate reacts with an enzyme:

$$E + S \xrightarrow[k_2]{k_1} ES \xrightarrow{k_3} E + P, \qquad \qquad Solution: \quad \frac{d[P]}{dt} = \frac{V_{\text{max}}[S]}{K_m + [S]}.$$

Suppose two substrates compete for the same enzyme:

$$E + S \xrightarrow{\frac{p_1}{p_2}} ES \xrightarrow{p_3} P + E$$

$$E + T \xrightarrow{\frac{q_1}{q_2}} ET \xrightarrow{q_3} Q + E$$

$$E_0 = [E] + [ES] + [ET].$$

Exercise (HW): Solve for $\frac{d[P]}{dt}$ and $\frac{d[Q]}{dt}$.

Question

How does the effects of the competition affect the dynamics of the system?

Michaelis-Menten variants

We just saw an equation for where 2 molecules of a substrate reacts with an enzyme:

$$E + 2S \xrightarrow[k_2]{k_1} ES_2 \xrightarrow{k_3} E + P, \qquad \qquad Solution: \quad \frac{d[P]}{dt} = \frac{V_{\text{max}}[S]^2}{K_m + [S]^2}.$$

But what if the binding is sequential?

$$E + S \rightleftharpoons_{k_2} ES + S \rightleftharpoons_{k_4} ES_2 \xrightarrow{k_5} P + E.$$

This is really two separate reactions:

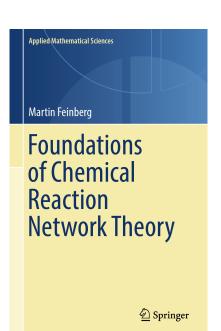
$$E+S \stackrel{k_1}{\rightleftharpoons} ES$$
, $ES+S \stackrel{k_3}{\rightleftharpoons} ES_2 \stackrel{k_5}{\longrightarrow} P+E$.

Question

When does this system become roughly the same as the Hill equation with coefficient n=2?

For more infomation

Karin Gatermann Lecture Notes in Mathematics **Computer Algebra Methods for Equivariant Dynamical Systems** 1728 Springer



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