Chemical reaction networks

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Algebraic Biology

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

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

Classification of reactions:

- $A \longrightarrow P$: "uni-molecular"
- $A + B \longrightarrow P$: "bi-molecular"
- $\blacksquare A + B + C \longrightarrow P: \quad \text{``tri-molecular''}$

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 [R].

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 $A \xrightarrow{k} P: \qquad \frac{d[P]}{dt} = k[A]$ $A \xrightarrow{k} P + Q: \qquad \frac{d[P]}{dt} = k[A]$ $A + B \xrightarrow{k} P: \qquad \frac{d[P]}{dt} = k[A][B]$ $2A \xrightarrow{k} P: \qquad \frac{d[P]}{dt} = k[A]^{2}$ $A + B \xrightarrow{k_{1}}{k_{2}} P: \qquad \frac{d[P]}{dt} = k_{1}[A][B] - k_{2}[P]$

An example

Consider the following chemical reaction network:

$$A + B \stackrel{k_1}{\underset{k_2}{\longleftrightarrow}} C, \qquad A \stackrel{k_3}{\longrightarrow} 2B$$

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

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

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:

$$A+B \stackrel{k_1}{\underset{k_2}{\longleftarrow}} C, \qquad A \stackrel{k_3}{\longrightarrow} 2B,$$

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 equilibria?
- Does the system have a *stable* positive equillbrium?
- Does the system have an *unstable* positive equilbrium?
- 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

$$E + S \underset{k_2}{\overset{k_1}{\longleftrightarrow}} ES \overset{k_3}{\longrightarrow} E + P$$

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

- 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]:

$$[ES] = \frac{E_0[S]}{\frac{k_2 + k_3}{k_1} + [S]}.$$

Alas, we can write

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

Michaelis-Menten equation

Recall the following chemical reaction:

$$E + S \xrightarrow[k_2]{k_1} ES \xrightarrow{k_3} E + P$$

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.

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

Remarks

- The "reaction rate", f([S]), is a strictly increasing function of [S].
- $\coprod_{[S] \to \infty} f([S]) = V_{\max}, \quad \text{(biologically, the maximum reaction rate)}$

$$\bullet f(K_m) = \frac{1}{2}V_{\max}.$$

• The reaction rate f([S]) is proportional to E_0 .

Michaelis-Menten equation

Recall the following chemical reaction:

$$E + S \underset{k_1}{\overset{k_2}{\longleftrightarrow}} ES \overset{k_3}{\longrightarrow} E + P$$

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

$$rac{d[S]}{dt} = -rac{d[P]}{dt} = -rac{V_{ ext{max}}[S]}{K_m + [S]} \, .$$

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:

$$E + nS \stackrel{k_1}{\underset{k_2}{\longleftrightarrow}} ES_n \stackrel{k_3}{\longrightarrow} E + P$$

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

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

Multi-molecule binding

Hill equation

Given the chemical reaction

$$E + nS \underset{k_1}{\overset{k_2}{\longleftrightarrow}} ES_n \overset{k_3}{\longrightarrow} E + P$$

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

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

This is called the Hill equation with Hill coefficient n.

Remarks

- The "reaction rate", f([S]), is a strictly increasing function of [S].
- $\blacksquare \lim_{[S] \to \infty} f([S]) = V_{\max}, \quad \text{(biologically, the maximum reaction rate)}$

$$\bullet f(K_m^{1/n}) = \frac{1}{2}V_{\max}.$$

- The reaction rate f([S]) is proportional to E_0 .
- 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:

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

Suppose two substrates compete for the same enzyme:

$$E + S \stackrel{p_1}{\underset{p_2}{\longrightarrow}} ES \stackrel{p_3}{\longrightarrow} P + E$$
$$E + T \stackrel{q_1}{\underset{q_2}{\longrightarrow}} ET \stackrel{q_3}{\longrightarrow} Q + E$$
$$E_0 = [E] + [ES] + [ET].$$

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:

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

But what if the binding is sequential?

$$E + S \stackrel{k_1}{\underset{k_2}{\leftrightarrow}} ES + S \stackrel{k_3}{\underset{k_4}{\leftrightarrow}} ES_2 \stackrel{k_5}{\longrightarrow} P + E.$$

This is really two separate reactions:

$$E + S \stackrel{k_1}{\underset{k_2}{\leftarrow}} ES, \qquad ES + S \stackrel{k_3}{\underset{k_4}{\leftarrow}} ES_2 \stackrel{k_5}{\longrightarrow} P + E.$$

Question

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

For more infomation

Lecture Notes in Mathematics

Karin Gatermann

Computer Algebra Methods for Equivariant Dynamical Systems

1728

Applied Mathematical Sciences

Martin Feinberg

Foundations of Chemical Reaction Network Theory



M. Macauley (Clemson)

Springer

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Books

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