

NATIONAL TAIWAN UNIVERSITY,
GRADUATE INSTITUTE OF BIOMEDICAL ENGINEERING AND BIOINFORMATICS

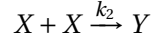
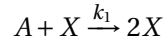
BEBI5009:
Mathematical Modeling of System Biology
Homework 3

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1 4.8.6 Global dynamics from local stability analysis.

a) Consider the chemical reaction network with mass-action kinetics:



i) Write a differential equation model describing the concentrations of X and Y

$$\begin{aligned}\frac{d[X]}{dt} &= -k_1[A][X] + 2k_1[A][X] - k_2[X]^2 = k_1[A][X] - k_2[X]^2 \\ \frac{d[Y]}{dt} &= k_2[X]^2 - k_3[Y]\end{aligned}$$

ii) Verify that the system has two steady states.

At steady-state,

$$\begin{aligned}\frac{d[X]_{ss}}{dt} = 0 &\Rightarrow k_1[A]_{ss}[X]_{ss} - k_2[X]_{ss}^2 = 0 \Rightarrow [X]_{ss}(k_1[A]_{ss} - k_2[X]_{ss}) = 0 \\ &\Rightarrow [X]_{ss} = 0 \quad \text{or} \quad [X]_{ss} = \frac{k_1}{k_2}[A]\end{aligned}$$

At the same time,

$$\frac{d[Y]_{ss}}{dt} = 0 \Rightarrow k_2[X]_{ss}^2 - k_3[Y]_{ss} = 0 \Rightarrow [Y]_{ss} = \frac{k_2}{k_3}[X]_{ss}^2$$

Then, we can conclude that there are two steady-state:

$$([X]_{ss}, [Y]_{ss}) = (0, 0) \quad \text{or} \quad \left(\frac{k_1}{k_2}[A], \frac{k_1^2}{k_2 k_3}[A]^2\right)$$

iii) Determine the system Jacobian at the steady states and characterize the local behavior of the system near these points

Let us denote the concentrations of species [X] as x and [Y] as y . Also, Let us denote the equations derived in (a) by $\frac{d[X]}{dt} = f(x, y)$ and $\frac{d[Y]}{dt} = g(x, y)$. Then,

$$J = \begin{bmatrix} \frac{\partial f(x,y)}{\partial x} & \frac{\partial f(x,y)}{\partial y} \\ \frac{\partial g(x,y)}{\partial x} & \frac{\partial g(x,y)}{\partial y} \end{bmatrix} = \begin{bmatrix} k_1[A] - 2k_2x & 0 \\ 2k_2x & -k_3 \end{bmatrix}$$

When the system is at steady-state that $([X]_{ss}, [Y]_{ss}) = (0, 0)$:

$$J = \begin{bmatrix} k_1[A] & 0 \\ 0 & -k_3 \end{bmatrix}$$

Eigen values of this Jacobian matrix are $k_1[A]$ and $-k_3$. Since one of is postive and the other is negative, the steady-state here is a saddle point. When the system is at steady-state that $([X]_{ss}, [Y]_{ss}) = (\frac{k_1}{k_2}[A], \frac{k_1^2}{k_2 k_3}[A]^2)$:

$$J = \begin{bmatrix} -k_1[A] & 0 \\ 2k_1[A] & -k_3 \end{bmatrix}$$

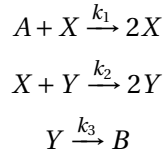
Eigen values of this Jacobian matrix are $-k_1[A]$ and $-k_3$. Since both eigen value of Jacobian matrix are negative, the steady-state here is a stable node.

- iv) By referring to the network, provide an intuitive description of the system behaviour starting from any initial condition for which $[X] = 0$.

Since the system is without X, the only reaction will happen in this system is $Y \rightarrow B$. The reaction will consume Y until that there is no Y any more.

- v) Sketch a phase portrait for the system that is consistent with your conclusions from (iii) and (iv).

b) Repeat for the system



In this case, you'll find that the non-zero steady-state is a center: it is surrounded by concentric periodic trajectories.

i)

$$\begin{aligned} \frac{d[X]}{dt} &= -k_1[A][X] + 2k_1[A][X] - k_2[X][Y] = k_1[A][X] - k_2[X][Y] \\ \frac{d[Y]}{dt} &= -k_2[X][Y] + 2k_2[X][Y] - k_3[Y] = k_2[X][Y] - k_3[Y] \end{aligned}$$

ii) At steady-state,

$$\begin{aligned} \frac{d[X]_{ss}}{dt} = 0 &\Rightarrow k_1[A]_{ss}[X]_{ss} - k_2[X]_{ss}[Y]_{ss} = 0 \Rightarrow [X]_{ss}(k_1[A]_{ss} - k_2[Y]_{ss}) = 0 \\ \Rightarrow [X]_{ss} = 0 &\text{ or } [Y]_{ss} = \frac{k_1}{k_2}[A] \\ \frac{d[Y]_{ss}}{dt} = 0 &\Rightarrow k_2[X]_{ss}[Y]_{ss} - k_3[Y]_{ss} = 0 \Rightarrow [Y]_{ss}(k_2[X]_{ss} - k_3) = 0 \\ \Rightarrow [Y]_{ss} = 0 &\text{ or } [X]_{ss} = \frac{k_3}{k_2} \end{aligned}$$

Then, we can conclude that there are two steady-state:

$$([X]_{ss}, [Y]_{ss}) = (0, 0) \text{ or } \left(\frac{k_3}{k_2}, \frac{k_1}{k_2}[A]\right)$$

- iii) Let us denote the concentrations of species [X] as x and [Y] as y . Also, Let us denote the equations derived in (a) by $\frac{d[X]}{dt} = f(x, y)$ and $\frac{d[Y]}{dt} = g(x, y)$. Then,

$$J = \begin{bmatrix} \frac{\partial f(x,y)}{\partial x} & \frac{\partial f(x,y)}{\partial y} \\ \frac{\partial g(x,y)}{\partial x} & \frac{\partial g(x,y)}{\partial y} \end{bmatrix} = \begin{bmatrix} k_1[A] - k_2y & -k_2x \\ k_2y & k_2x - k_3 \end{bmatrix}$$

When the system is at steady-state that $([X]_{ss}, [Y]_{ss}) = (0, 0)$:

$$J = \begin{bmatrix} k_1[A] & 0 \\ 0 & -k_3 \end{bmatrix}$$

Eigen values of this Jacobian matrix are $k_1[A]$ and $-k_3$. Since one of is positive and the other is negative, the steady-state here is a saddle point. When the system is at steady-state that $([X]_{ss}, [Y]_{ss}) = (\frac{k_3}{k_2}, \frac{k_1}{k_2}[A])$:

$$J = \begin{bmatrix} 0 & -k_3 \\ k_1[A] & 0 \end{bmatrix}$$

Eigen values of this Jacobian matrix are roots of $\lambda^2 = -k_1 k_3 [A]$. Since both eigen value of Jacobian matrix don't have zero real parts, the steady-state here is a center, which has periodic trajectories.

- iv) The situation is the same as counterpart of part (a). Since the system is without X, the only reaction will happen in this system is $Y \rightarrow B$. The reaction will consume Y until that there is no Y any more.
- v)

2 4.8.8 Linearization.

Consider the simple reaction system $\rightarrow S \rightarrow$, where the reaction rates are

$$\text{production: } V_0 \quad \text{consumption: } \frac{V_{max}[S]}{K_M + [S]}$$

- a) Write the differential equation that describes the dynamics in $s = [S]$. Find the steady state. Next, approximate the original system by linearizing the dynamics around the steady state. This approximation takes the form of a linear differential equation in the new variable $x(t) = s(t) - s^{ss}$.

It's intuitive to construct the system:

$$f(s) = \frac{ds}{dt} = V_0 - \frac{V_{max}s}{K_M + s}$$

and we can find steady state.

$$\frac{ds^{ss}}{dt} = V_0 - \frac{V_{max}s^{ss}}{K_M + s^{ss}} = 0 \Rightarrow s^{ss} = \frac{K_M V_0}{V_{max} - V_0}$$

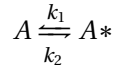
Then, we can approximate original system by linearizing the dynamics around the steady state.

$$\begin{aligned}
 f(s) &\approx f(s^{ss}) + \frac{df}{ds} \big|_{s=s^{ss}} (s - s^{ss}) \\
 &= 0 + \left(-\frac{V_{max}(K_M + s^{ss}) - V_{max}s^{ss}}{(K_M + s^{ss})^2} \right) (s - s^{ss}) \\
 &= -\frac{V_{max}K_M}{(K_M + s^{ss})^2} (s - s^{ss}) \\
 \Rightarrow f(x) &= -\frac{V_{max}K_M}{(K_M + s^{ss})^2} x(t) = -\frac{V_{max}K_M}{\left(K_M + \frac{K_M V_0}{V_{max} - V_0}\right)^2} x(t) = -\frac{V_{max}}{K_M \left(1 + \frac{V_0}{V_{max} - V_0}\right)^2} x(t)
 \end{aligned}$$

- b) Take parameter values $V_0 = 2$, $V_{max} = 3$, and $K_M = 1$ and run simulations of the nonlinear and linearized systems starting at initial conditions $[S] = 2.1$, $[S] = 3$, and $[S] = 12$. Comment on the discrepancy between the linear approximation and the original nonlinear model.

3 4.8.13 Sensitivity analysis: reversible reaction.

Consider the reversible reaction with mass-action rate constants as shown.



Let T be the total concentration of A and A*.

- a) Solve for the steady-state concentration of A* and verify that an increase in k_1 leads to an increase in $[A^*]^{ss}$.

First, we can write down these relationships:

$$\begin{aligned}
 \frac{d[A]}{dt} &= -k_1[A] + k_2[A^*] \\
 \frac{d[A^*]}{dt} &= k_1[A] - k_2[A^*] \\
 T &= [A] + [A^*]
 \end{aligned}$$

Then, we can utilize differential equation of one of species and total concentration of A and A*.

$$\begin{aligned}
 \frac{d[A]^{ss}}{dt} &= -k_1[A]^{ss} + k_2[A^*]^{ss} = -k_1[A]_{ss} + k_2(T - [A]^{ss}) = -(k_1 + k_2)[A]^{ss} + k_2T = 0 \\
 \Rightarrow [A]^{ss} &= \frac{k_2T}{k_1 + k_2} \\
 \Rightarrow [A^*]^{ss} &= T - [A]^{ss} = \frac{k_1T}{k_1 + k_2}
 \end{aligned}$$

To verify the statement of the question, we can simply take derivative of $[A^*]^{ss}$.

$$\frac{d[A^*]^{ss}}{dk_1} = \frac{d\left(\frac{k_1 T}{k_1 + k_2}\right)}{dk_1} = \frac{T(k_1 + k_2) - k_1 T}{(k_1 + k_2)^2} = \frac{k_2 T}{(k_1 + k_2)^2} > 0$$

where the positive slope verifies that the increasement in k_1 will lead to an increase in $[A^*]^{ss}$

- b) Use parametric sensitivity analysis to determine whether the steady state concentration of A^* is more sensitive to a 1 % increase in T or a 1 % increase in k_1 . Does the answer depend on the values of the parameters?

First, we can extend the result of (a).

$$\frac{\frac{d[A^*]^{ss}}{[A]^{ss}}}{\frac{dk_1}{k_1}} = \frac{d[A^*]^{ss}}{dk_1} \frac{k_1}{[A^*]^{ss}} = \frac{k_2 T}{(k_1 + k_2)^2} \frac{k_1}{\frac{k_1 T}{k_1 + k_2}} = \frac{k_2}{k_1 + k_2}$$

Then, we can also derive the relative sensitivity of T .

$$\frac{\frac{d[A^*]^{ss}}{[A]^{ss}}}{\frac{dT}{T}} = \frac{d[A^*]^{ss}}{dT} \frac{T}{[A^*]^{ss}} = \frac{k_1}{k_1 + k_2} \frac{T}{\frac{k_1 T}{k_1 + k_2}} = 1$$

Finally, we can conclude that the steady state concentration of A^* is more sensitive to a 1 % increase in T than a 1 % increase in k_1 , because $\frac{k_2}{k_1 + k_2} < 1$ and it's independent to the values of the parameters.