

Modeling the MAPK pathway: Derive CCM model

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Introduction

Structural Causal Models(SCMs) is a widely-used causal modeling framework. However, it is not powerful enough in representing dynamical systems at equilibrium. To solve this problem, Blom and Mooij [1] proposed a generalize SCM, called Causal Constraints Model(CCM), which could capture the causal semantics in dynamical systems comparing with SCMs. This project focus on a biological process, Mitogrn-activated protein kinase (MAPK) pathway, and try to derive CCM on this pathway.

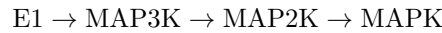
Derive Causal Constraints Models

Brief Introduction of CCM

To model the effects of all perfect interventions, SCM could only give several structural equations, which leaves no freedom to model the dependence on initial conditions, while CCM is specified by causal constraints, which are the relations between variables that constrain the solutions of the model under explicitly specified intervention targets.

Mitogrn-activated protein kinase (MAPK) pathway

In this project, our group mainly focus on an important biological process, MAPK pathway. It is a cascade of three proteins, a MAPK (aka Erk), a MAPK kinase (MAP2K, aka Mek), and a MAPK kinase kinase (MAP3K, aka Raf), represented with a causal diagram.



The boiochemical reactions

A protein molecule is in an active state if it has one or more attached phosphoryl groups. Each arrow in the equation above combines the reactions of the phosphorylation (i.e., activation) and dephosphorylation (i.e., desactivation). For example, $E1 \rightarrow \text{MAP3K}$ combines two reactions



From the above reaction, we can figure out the ordinary differential equations \mathbb{M} .

$$\mathbb{M} = \begin{cases} \frac{dK3}{dt} = & v_{K3}^{\text{act}} E1 (T_{K3} - K3(t)) - v_{K3}^{\text{inh}} K3(t) \\ \frac{dK2}{dt} = & \frac{(v_{K2}^{\text{act}})^2}{v_{K2}^{\text{inh}}} K3(t)^2 (T_{K2} - K2(t)) - v_{K2}^{\text{act}} K3(t) K2(t) - v_{K2}^{\text{inh}} K2(t) \\ \frac{dK1}{dt} = & \frac{(v_{K1}^{\text{act}})^2}{v_{K1}^{\text{inh}}} K2(t)^2 (T_{K1} - K1(t)) - v_{K1}^{\text{act}} K2(t) K1(t) - v_{K1}^{\text{inh}} K1(t) \end{cases}$$

Find Equilibrium

Before constructing the Causal Constraints Model (CCM), we need first figure out the Structural Causal Model (SCM) from the ordinary differential equations \mathbb{M} . In order to achieve that, first, find the steady state of K_1, K_2, K_3 .

First of all, to find the steady state of K_3 , we let $\frac{dK_3}{dt} = 0$, then we have:

$$\begin{aligned}\frac{dK_3}{dt} &= 0 \\ v_{K_3}^{act} E_1 (T_{K_3} - K_3(t)) - v_{K_3}^{inh} K_3(t) &= 0 \\ v_{K_3}^{act} E_1 T_{K_3} - v_{K_3}^{act} E_1 K_3(t) &= v_{K_3}^{inh} K_3(t) \\ \frac{v_{K_3}^{act}}{v_{K_3}^{inh}} E_1 T_{K_3} - \frac{v_{K_3}^{act}}{v_{K_3}^{inh}} E_1 K_3(t) &= K_3(t) \\ K_3(t) &= \frac{\frac{v_{K_3}^{act}}{v_{K_3}^{inh}} E_1 T_{K_3}}{1 + \frac{v_{K_3}^{act}}{v_{K_3}^{inh}} E_1}\end{aligned}$$

Let $g_1(u) = \frac{u}{1+u}$ so we have $K_3^* = T_{K_3} g_1(\omega_{K_3} E_1)$ where $\omega_{K_3} = \frac{v_{K_3}^{act}}{v_{K_3}^{inh}}$.

Similarly, we can find the steady state for K_2, K_1

$$\begin{aligned}\frac{dK_2}{dt} &= 0 \\ \frac{(v_{K_2}^{act})^2}{v_{K_2}^{inh}} K_3^2(t) T_{K_2} - \frac{(v_{K_2}^{act})^2}{(v_{K_2}^{inh})^2} K_3^2(t) K_2(t) - v_{K_2}^{inh} K_2(t) &= 0 \\ (\frac{v_{K_2}^{act}}{v_{K_2}^{inh}})^2 K_3^2(t) T_{K_2} - (\frac{v_{K_2}^{act}}{v_{K_2}^{inh}})^2 K_3^2(t) K_2(t) - \frac{v_{K_2}^{act}}{v_{K_2}^{inh}} K_3(t) K_2(t) - K_2(t) &= 0 \\ K_2(t) &= \frac{(\frac{v_{K_2}^{act}}{v_{K_2}^{inh}})^2 K_3^2(t) T_{K_2}}{1 + (\frac{v_{K_2}^{act}}{v_{K_2}^{inh}})^2 K_3^2(t) + \frac{v_{K_2}^{act}}{v_{K_2}^{inh}} K_3(t)}\end{aligned}$$

Here we can also define a function $g_2(u) = \frac{u^2}{1+u+u^2}$ so that we can rewrite the above equation as $K_2(t) = T_{K_3} g_2(\omega_{K_2} K_3)$ where ω_{K_2} represents $\frac{v_{K_2}^{act}}{v_{K_2}^{inh}}$

Find the equilibrium of K_1 is same as K_2 , we can easily get the equation of K_1 :

$$K_1(t) = \frac{(\frac{v_{K_1}^{act}}{v_{K_1}^{inh}})^2 K_2^2(t) T_{K_1}}{1 + (\frac{v_{K_1}^{act}}{v_{K_1}^{inh}})^2 K_2^2(t) + \frac{v_{K_1}^{act}}{v_{K_1}^{inh}} K_2(t)}$$

Then we rewrite this equation as $K_1(t) = T_{K_2} g_2(\omega_{K_1} K_2)$ where $\omega_{K_1} = \frac{v_{K_1}^{act}}{v_{K_1}^{inh}}$.

To conclude, the equilibrium of this reaction is:

$$\begin{cases} K_3^* = T_{K_3} g_1(\omega_{K_3} E_1), \\ K_2^* = T_{K_2} g_2(\omega_{K_2} K_3), \\ K_1^* = T_{K_1} g_2(\omega_{K_1} K_2) \end{cases}$$

Where K_3^*, K_2^*, K_1^* are the equilibrium values of $K_3(t), K_2(t), K_1(t)$, $g_1(u) = \frac{u}{1+u}$, $g_2(u) = \frac{u^2}{1+u+u^2}$ and $\omega_X = \frac{v_{act}^X}{v_{inh}^X}$

Proof of CCM

Our MAPK pathway fulfills basic enzymatic reaction model described in Alon's paper (Appendix).

They are both a model of the action of an enzyme X on its substrate S, to catalyze formation of product P. Enzyme X and substrate S bind with rate k_{on} to form a complex $[XS]$, which dissociates with rate k_{off} . [2]

Blom and Mooij's basic enzymatic reaction model (equation 2-6)[1] is equivalent to Alon's model (Appendix)[2]. Proof is below.

1) In Alon's model,



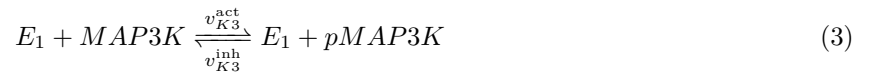
2) In Blom and Mooij's model,



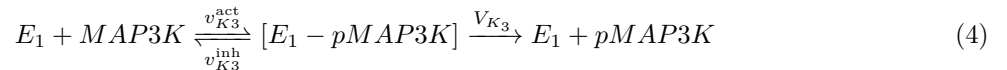
When $K_0 = 0, K_3 = 0$, two models are equivalent. So the CCM model with all constraints can applied to Alon's reaction class, including MAPK pathway.

There is no constraints for any one of three enzyme reaction in MAPK pathway. Proof is shown below.

Reaction $E1 \rightarrow MAP3K$ can be written as



It can also be rewritten as



Using conclusion from Blom and Mooij's paper, there will be two constraints.

$$\frac{d[E_1 - pMAP3K]}{dt} + \frac{d[E_1]}{dt} = E_{1_0} + [MAP3K]_0 \quad (5)$$

$$\frac{d[MAP3K]}{dt} - \frac{d[E_1]}{dt} = -V_{K_3} \frac{d[E_1 - pMAP3K]}{dt} \quad (6)$$

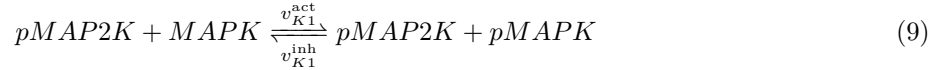
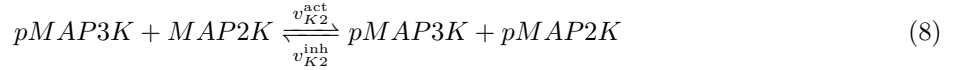
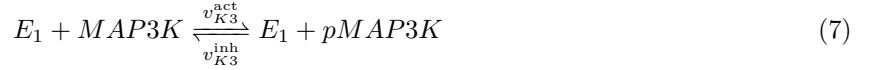
Obviously, two constraints are both related to intermediate product $[E_1 - pMAP3K]$, which is not existed in our degenerated model.

As a result, our $E_1 \rightarrow MAP3K$ has no constraint. Apparently, $MAP3K \rightarrow MAP2K$ and $MAP2K \rightarrow MAPK$ have no constraints either.

There is no constraints for combination of all three enzyme reactions in MAPK pathway. Proof is shown below.

Lemma: To claim a constraint, there must exists some algebra manipulation on time-derivatives that result in non-convergence in equilibrium under some conditions.

In our system,



$$\frac{d[E_1]}{dt} = 0 \quad (10)$$

$$\frac{d[MAP3K]}{dt} = -\frac{dK_3}{dt} = v_{K_3}^{act} E_1 (T_{K_3} - K_3(t)) + v_{K_3}^{inh} K_3(t) \quad (11)$$

$$\frac{d[pMAP3K]}{dt} = \frac{dK_3}{dt} \quad (12)$$

$$\frac{d[MAP2K]}{dt} = -\frac{dK_2}{dt} = \frac{(v_{K_2}^{act})^2}{v_{K_2}^{inh}} (K_3(t))^2 (T_{K_3} - K_2(t)) + v_{K_2}^{act} K_3(t) K_2(t) + v_{K_2}^{inh} K_2(t) \quad (13)$$

$$\frac{d[pMAP2K]}{dt} = \frac{dK_2}{dt} \quad (14)$$

$$\frac{d[MAPK]}{dt} = -\frac{dK_1}{dt} = \frac{(v_{K_1}^{act})^2}{v_{K_1}^{inh}} (K_2(t))^2 (T_{K_2} - K_1(t)) + v_{K_1}^{act} K_2(t) K_1(t) + v_{K_1}^{inh} K_1(t) \quad (15)$$

For all the nine time-derivatives above (eq.(7) to (15)), there is no combination that yields an outcome that is not dependent on parent variables. So there is no constraints on the system.

To conclude, our ODE derived CCM has no constraints on steady state for any intervention.

Table of intervention

Since there is no constrain in this case, we can do intervention on all conditions we want, following tables summarise all possible intervention and their results.

Intervention	K_1	K_2	K_3
none	$T_{k_1}g_2(\omega_{K_1}T_{k_2}g_2(\omega_{K_2}T_{k_3}g_1(\omega_{K_3}e_{10})))$	$T_{k_2}g_2(\omega_{K_2}T_{k_3}g_1(\omega_{K_3}e_{10}))$	$T_{k_3}g_1(\omega_{K_3}e_{10})$
$do(K_1 = k_1)$	k_1	$T_{k_2}g_2(\omega_{K_2}K_3)$	$T_{k_3}g_1(\omega_{K_3}E_1)$
$do(K_2 = k_2)$	$T_{k_1}g_2(\omega_{K_1}k_2)$	k_2	$T_{k_3}g_1(\omega_{K_3}E_1)$
$do(K_3 = k_3)$	$T_{k_1}g_2(\omega_{K_1}T_{k_2}g_2(\omega_{K_2}k_3))$	$T_{k_2}g_2(\omega_{K_2}k_3)$	k_3
$do(K_1 = k_1, K_2 = k_2)$	k_1	k_2	$T_{k_3}g_1(\omega_{K_3}E_1)$
$do(K_1 = k_1, K_3 = k_3)$	k_1	$T_{k_2}g_2(\omega_{K_2}k_3)$	k_3
$do(K_2 = k_2, K_3 = k_3)$	$T_{k_1}g_2(\omega_{K_1}k_2)$	k_2	k_3
$do(K_1 = k_1, K_2 = k_2, K_3 = k_3)$	k_1	k_2	k_3

References

- [1] T. Blom, J.M.Mooij (2019), *35th Annual Conference on Uncertainty in Artificial Intelligence*, Causal Constrain Models.
- [2] Uri Alon. Network Motif(book chapter 6 and appendix).