

Topological Modelling of Biological Interactions for Robustness

1 Goals

1. To define measures of similarity and robustness of network topologies so that small changes in the input do not affect the output.
2. To create a method of finding networks which produce the desired chemical concentrations with a high degree of robustness against changes in input.

2 Description as Dynamical System

Biological chemical networks can be described in terms of dynamical systems with inputs. At this point we make the assumption that inputs to the system D (degradation, external input) are linear, as well as repression or induction of one chemical on another W . We can then describe a system $S = \langle s_0, D, W \rangle$ such that $s_0 \in \mathbb{R}^n, D \in \mathbb{R}^n, W \in \mathbb{M}^n$, where $\mathbb{M}^n = \mathbb{R}^{n \times n}$. Let s_i be the i^{th} component of state vector s . Σ refers to the set of all such systems S .

$$S = \langle s_0, D, W \rangle \quad (1)$$

$$\frac{ds_i}{dt}(t) = D_i s_i(t) + \sum_{j=1}^n W_{ji} \cdot s_j(t) \quad (2)$$

3 Similarity of a Dynamical System

We define similarity of a dynamical system of this type to quantify the “closeness” of one network to another. We allow “shifting” and “stretching” of the attractor of the system to form a bijection from one system to a transformed system which is as close to the other system as possible. In the below equations, stretching is performed by the factors ϕ_1, ϕ_2 , and shifting is performed by the offsets t_{p_1}, t_{p_2} .

M_g is called the “generalized” similarity between two systems, and M_l is the “linear” form of that generalized similarity. In the equation for M_g , P is the “penalty” for shifting and stretching, and d is an arbitrary metric to determine closeness of two state vectors $s_1, s_2 \in \mathbb{R}^n$. $s(t)$ is the value of the state vector s at time t .

$$M : \Sigma \times \Sigma \rightarrow \mathbb{R} \quad (3)$$

$$s'_1 = s_1(\phi_1 \cdot (t + t_{p_1})) \quad (4)$$

$$s'_2 = s_2(\phi_2 \cdot (t + t_{p_2})) \quad (5)$$

$$M_g(S_1, S_2) = \min_{\substack{\phi_1, \phi_2 \geq 1 \\ t_{p_1}, t_{p_2} \geq 0}} P \left(\int_0^\infty d(s'_1(t), s'_2(t)) dt, \phi_1, \phi_2, t_{p_1}, t_{p_2} \right) \quad (6)$$

$$P_l(r, \phi_1, \phi_2, t_{p_1}, t_{p_2}) = r\phi_1\phi_2 + t_{p_1} + t_{p_2} \quad (7)$$

$$d_l(s_1, s_2) = |s'_1 - s'_2| \quad (8)$$

$$M_l(S_1, S_2) = \min_{\substack{\phi_1, \phi_2 \geq 1 \\ t_{p_1}, t_{p_2} \geq 0}} \left(\int_0^\infty |s'_1(t) - s'_2(t)| dt \right) \phi_1\phi_2 + t_{p_1} + t_{p_2} \quad (9)$$

4 Robustness of a Dynamical System

“Robustness” is meant to quantify the tendency of the system to remain similar (the “similarity” defined in 3) given small changes in input. There is a simple formulation $R_s(s)$ used to motivate a more general formulation $R_g(s)$, and finally a linear formulation $R_l(s)$, which is what is used in 5.

The simple formulation quantifies the condition that all “nearby” systems have “similar” behavior. This is extremely similar to the definition of continuity of a function at a point in a topological space. Nearby systems to a state are those within the ball centered at that state, for a given radius $\epsilon > 0$, using some metric m . m is *not* the similarity metric discussed in 3. An *example* of m is given in (11). s_{i_0} refers to the initial state vector s_0 for the system S_i .

$$R : \Sigma \rightarrow \mathbb{R} \quad (10)$$

$$m(S_1, S_2) = |s_{1_0} - s_{2_0}| \quad (11)$$

$$R_s(S) = \arg\inf_{\delta > 0} M(S, S') < \delta \forall S' \in B_m(s, \epsilon) \quad (12)$$

The general formulation of robustness of a system is defined in terms of three functions f , g , and h . f represents how “important” the initial states near some state s are, g represents how “important” the similarity of each nearby system is, and h combines the two measures.

$$f : \Sigma \times \Sigma \rightarrow \mathbb{R} \quad (13)$$

$$g : \Sigma \times \Sigma \rightarrow \mathbb{R} \quad (14)$$

$$h : \mathbb{R} \times \mathbb{R} \rightarrow \mathbb{R} \quad (15)$$

$$R_g(S) = \int_{S' \in \Sigma} h(f(S, S'), g(S, S')) \quad (16)$$

The general formulation of robustness can be used to model the simple formulation, given $\epsilon > 0$.

$$f(S, S') = \begin{cases} 1 & |s_0 - s'_0| \\ 0 & \text{else} \end{cases} \quad (17)$$

$$M_{sup}(S) = \sup_{\sigma \text{ s.t. } f(S, \sigma)=1} M(S, \sigma) \quad (18)$$

$$g(S, S') = \begin{cases} 1 & M(S, S') < M_{sup}(S) \\ 0 & \text{else} \end{cases} \quad (19)$$

$$h(a, b) = a \cdot b \quad (20)$$

Or, more simply:

$$M_{sup}(S) = \sup_{\sigma \text{ s.t. } |s_0 - \sigma| < \epsilon} M(S, \sigma) \quad (21)$$

$$f(S, S') = 1 \quad (22)$$

$$(23)$$

The linear formulation of robustness is a member of the class of functions defined by the general formulation, and is used in 5.

$$f(S, S') = |s_0 - s'_0|^{-1} \quad (24)$$

$$g = M_l \quad (25)$$

$$R_l(S) = \int_{S' \in \Sigma} \frac{M_l(S, S')}{|s_0 - s'_0|} \quad (26)$$

5 Finding Maximally Robust Approximately Isomorphic Systems

Given (1), (2), (9), and (26), we can attempt to solve the problem posed in the introduction. We start with these linear formulations because we believe they will be easier to solve in closed form. We wish to find a system S^* which is similar to some system S and which is maximally robust.

Given $\epsilon > 0$. Let $S' = \langle s'_0, D, W \rangle$ for some $s'_0 \in \mathbb{R}^n$.

$$\Sigma_{similar} = \{S' \in \Sigma \mid M_l(S, S') < \epsilon\} \quad (27)$$

$$S^* = \underset{S' \in \Sigma_{similar}}{\operatorname{argsup}} R_l(S') \quad (28)$$

We can probably do this by taking the derivative of R_l , somehow.