

## Goal

(*italicized words identify optimization goals*)

Wish to find  $W$  which has the *most* regular behavior over the *largest* contiguous subset of initial concentrations  $C_0$ , given gaussian disturbances of  $W$ . The contiguity may not be required, but seems to make sense (modelling a region which the cell can reliably keep the ambient concentrations within). We could take into account non-contiguous subsets, but that is not likely to be useful (it would be weird to have a motif which has similar behavior over different non-contiguous regions of ambient concentration).

## Formulation

We define the norm over a matrix.

$$W - W' = W'', \text{ where } W''_{ij} = W_{ij} - W'_{ij} \quad \forall i, j \quad (1)$$

$$|W| : \mathbb{R}_{n \times n} \rightarrow \mathbb{R} = \sqrt{\sum_{i,j \in [1,n]} W_{ij}} \quad (2)$$

Given fixed parameters  $W \in \mathbb{R}_{n \times n}$ ,  $\sigma_W \in \mathbb{R}$ , define a Gaussian distribution of weights.

$$D(W'|W, \sigma_W) = (2\sigma_W^2\pi)^{-2} e^{-\frac{(|W' - W|)^2}{2\sigma_W^2}} \quad (3)$$

This says “mutations modify the weights of the matrix  $W'$  a Euclidean distance from the original weight matrix  $W$  with probability according to the gaussian distribution with parameter  $\sigma_W$ .” This may not be realistic, but it captures the idea that if a mutation occurs, it is likely to modify multiple weights at once if it modifies any at all.

We define the “behavior” of a network  $(C_0, W)$  as the concentration vector  $C(T)$  at some given time  $T$ . This assumes linear differential equations modelling activation and inhibition rates with no external input or output.

$$C_{C_0, W}(t) = C_0 + \int_0^t W \cdot C(t) dt \quad (4)$$

This may not be appropriate; some networks may have *very* reliable periodic behavior, where the period simply changes slightly in response to disturbance, which would lead to vastly different  $C(T)$ . However, if we restrict ourselves to networks which reach some sort of equilibrium (which need not be defined or used at this time), we can use just  $C(T)$ . We define networks that reach equilibrium over some region “centered” at  $C_0$  as networks which have similar  $C(T)$  for some large  $T$  over  $C'_0$  normally distributed with mean  $C_0$ .  $T > 0$

and must be “large enough to allow the system to get to equilibrium.” We can accomplish this by simply choosing a timescale and picking networks which achieve equilibrium.

## Variability

Variability  $V$  of a network is defined using variation  $v$  with an initial concentration  $(C_0, W)$  over some gaussian distribution of  $C'_0$  is defined.

$$v(C_0, W', W) = \int_{C'_0 \in \mathbb{R}^n} D(C'_0|C_0, \sigma_C) \cdot |C_{C'_0, W'}(T) - C_{C_0, W}(T)| \quad (5)$$

$$\begin{aligned} V(C_0, W) &= \int_{W' \in \mathbb{R}^{n \times n}} D(W'|W, \sigma_W) \cdot v(C_0, W', W) \\ &= \int_{W', C'_0} D(W'|W, \sigma_W) D(C'_0|C_0, \sigma_C) \cdot |C_{C'_0, W'}(T) - C_{C_0, W}(T)| \end{aligned} \quad (6)$$

Finding stable networks  $X$ , then, is defined as an optimization problem. The most robust networks (the ones with the minimum variability) are (hopefully) likely to produce interesting network motifs.

$$X^* = \min_W \int_{C_0 \in \mathbb{R}^n} V(C_0, W) \quad (7)$$

Simulation of  $(C_0, W)$  seems possible by discretizing choices of  $C'_0$  and  $W'$  normally distributed about  $C_0$  and  $W$ .