

Research Notes on Self-Coupled Spiking Neural Networks

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today

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1 The self-coupled SNN Model

Problem Statement

Given:

- A Linear Dynamical System $\frac{dx}{dt} = Ax(t) + Bc(t)$, $x \in \mathbf{R}^d$
- A Decoder Matrix $D \in \mathbf{R}^d \times N$ specifying the preferred directions of N neurons in d -dimensional space,

synthesize a spiking neural network that implements the linear dynamical system.

Features

1. **Long-Term Network Accuracy** The Deneve network assumes $\hat{x} = x$. We show this assumption produces estimation error between the network and its target system that increases with time. By avoiding this assumption, the self-coupled network remains accurate over time.
2. **Tuning Curve Rotation** To most efficiently use neurons, we use orthogonal coding directions via SVD. The dynamics matrix A is diagonalized by an orthonormal basis \mathcal{U} in d -dimensional space, while the decoder matrix D is chosen such that \mathcal{U} gives its left singular vectors. This choice of coding directions eliminates the need for coupling. ?
At least two neurons per dimension (2d in total) are required since ~~voltage thresholds~~ ^{spikes} are strictly positive. N -neuron ensembles can thus represent systems with $\frac{N}{2}$ dimensions or less.
3. **Post-synaptic Spike Dropping** At each synapse, neurotransmitter release due to an action potential is probabilistic. We incorporate probabilistic spike transmission by thinning at every synaptic connection. The pre-synaptic neuron's membrane potential is still deterministically reset by an action potential.
4. **Dimensionless Time** We describe both the network and target system in dimensionless time. Time is normalized by the synapses' time constant, τ_s . This dimensionless representation ensures consistent numerical simulation independent of simulation timestep. Furthermore, τ_s is implicitly specified as 1, reducing the model's parameters by one.

2 Basic Model

2.1 Derivation

1. Let τ_s be the synaptic time constant of each synapse in the network. Define dimensionless time as:

$$\xi \triangleq \frac{t}{\tau_s}.$$

We now assume our Linear Dynamical System is expressed in dimensionless time, i.e

$$\frac{dx}{d\xi} = Ax(\xi) + Bc(\xi). \quad (2.1)$$

To describe the neuron dynamics in dimensionless time, let $o(\xi) \in \mathbf{R}^N$ be the spike trains of N neurons composing the network with components

$$o_j(\xi) = \sum_{k=1}^{n_j \text{ spikes}} \delta(\xi - \xi_j^k),$$

where ξ_j^k is the time at which neuron j makes its k^{th} spike. Define the network's estimate of the state variable as

$$\hat{x}(\xi) \triangleq Dr(\xi), \quad (2.2)$$

where $D \in \mathbf{R}^{d \times N}$ and

$$\frac{dr}{d\xi} = -r + o(\xi). \quad (2.3)$$

When the probability of synaptic transmission is 1, component r_j is the total received post-synaptic current (PSC) from neuron j by the network estimator. Define the network error as

$$e(\xi) \triangleq x(\xi) - \hat{x}(\xi). \quad (2.4)$$

2. From equations (2.3) and (2.2), we have

$$D\dot{r} + Dr = Do$$

$$\implies \dot{\hat{x}} + \hat{x} = Do,$$

where the dot denotes derivative w.r.t dimensionless time ξ .

Subtract $\dot{\hat{x}}$ from \dot{x} to get \dot{e} :

$$\begin{aligned} \dot{e} &= \dot{x} - \dot{\hat{x}} \\ &= (Ax + Bc) - (Do - \hat{x}) \\ &= A(e + \hat{x}) + Bc - Do + \hat{x} \\ &= Ae + (A + I)\hat{x} + Bc - Do \\ &= Ae + (A + I)(Dr) + Bc - Do \\ \implies D^T \dot{e} &= D^T Ae + D^T(A + I)(Dr) + D^T Bc - D^T Do. \end{aligned}$$

The quantity $D^T e$ defines the membrane voltage of the predictive coding framework (PCF), a precursor to this model:

$$v_{pcf} \triangleq D^T e.$$

Note that the definition implies $e = D^{T\dagger} v_{pcf}$. The voltage dynamics are thus

$$\dot{v}_{pcf} = D^T A D^{T\dagger} v_{pcf} + D^T (A + I) (Dr) + D^T Bc - D^T Do, \quad (2.5)$$

where $D^{T\dagger}$ is the left pseudo-inverse of $D^T \in \mathbf{R}^{N \times d}$. The PCF thus defines a mapping between two vector spaces: the d-dimensional state space of the target system, and the N-dimensional voltage space of the spiking neural network. This mapping is visualized in figure (1).

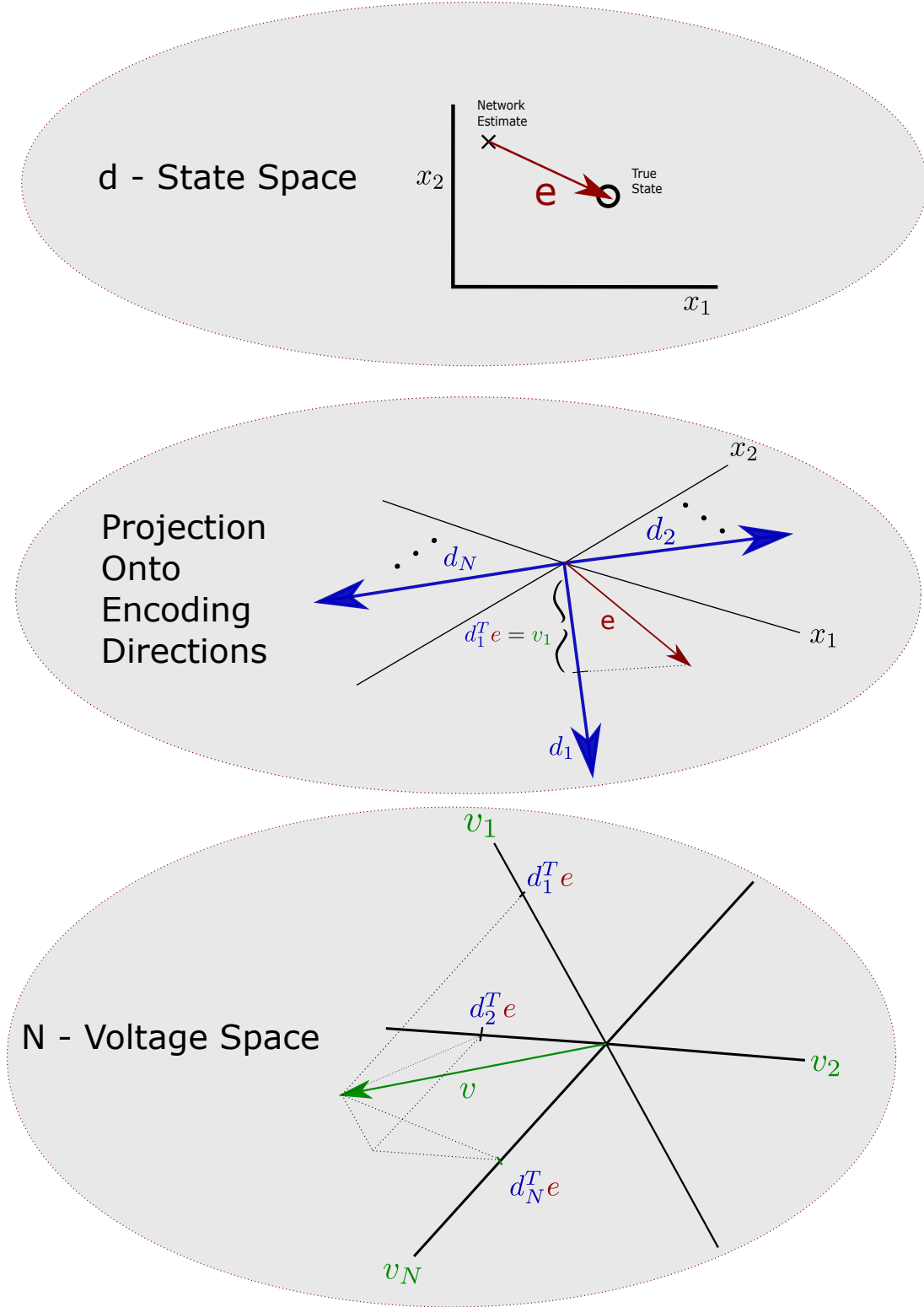


Figure 1: Mapping Between State and Voltage Spaces: **Top:** The estimation error e is computed by comparing the decoded network estimate to the true state of the target dynamical system. **Middle:** The e is projected onto the encoding directions of the neurons composing the network. The projection of error onto encoding direction j gives the membrane voltage of neuron j , $v_j = d_j^T e$. **Bottom:** The voltages form a N -dimensional vector contained in voltage space. 5

3. The self-coupled network is derived via a change of bases. Assuming both D and A are full rank, diagonalize each to a common left basis:

$$A = \mathcal{U} \Lambda \mathcal{U}^T = \sum_{j=1}^d \Lambda_j \mathcal{U}_j \mathcal{U}_j^T,$$

$$D = \mathcal{U} \begin{bmatrix} S & 0 \end{bmatrix} V^T = \sum_{j=1}^d S_j \mathcal{U}_j V_j^T,$$

$$D^T = V \begin{bmatrix} S \\ 0 \end{bmatrix} \mathcal{U}^T = \sum_{j=1}^d S_j V_j \mathcal{U}_j^T,$$

$$D^T D = V \begin{bmatrix} S \\ 0 \end{bmatrix} \begin{bmatrix} S & 0 \end{bmatrix} V^T = \sum_{j=1}^d S_j^2 V_j V_j^T,$$

with $\mathcal{U} \in \mathbf{R}^{d \times d}$ and $V \in \mathbf{R}^{N \times N}$, and $S \in \mathbf{R}^{d \times d}$.

In the original basis, the state is x . In the rotated basis we denote this quantity as y . It is the projection of x onto the d -dimensional \mathcal{U} basis:

$$y \triangleq \mathcal{U}^T x \tag{2.6}$$

The rotated target dynamics are thus

$$\begin{aligned} \dot{y} &= \mathcal{U}^T \dot{x} \\ &= \Lambda y(\xi) + \mathcal{U}^T B c(\xi) \\ &= \Lambda y(\xi) + \mathcal{U}^T B \mathcal{U} \mathcal{U}^T c(\xi) \\ &= \Lambda y(\xi) + \beta \tilde{c}(\xi) \end{aligned} \tag{2.7}$$

where

$$\beta \triangleq \mathcal{U}^T B \mathcal{U},$$

and

$$\tilde{c} \triangleq \mathcal{U}^T c,$$

give the projections of B and c respectively. The network estimate in the rotated basis is

$$\hat{y} \triangleq \mathcal{U}^T \hat{x}.$$

From equation (2.2),

$$\begin{aligned}
\hat{y} &= \mathcal{U}^T \hat{x} \\
&= \mathcal{U}^T D r \\
&= [S \quad 0] V^T r \\
&= [S \quad 0] \rho \\
\implies \dot{\hat{y}} &= [S \quad 0] V^T \dot{r} \\
&= [S \quad 0] (-V^T r + V^T o).
\end{aligned}$$

Note that $V^T r$ and $V^T o$ are projections of the N-neuron network's post-synaptic current and spike train respectively onto the rotated basis, denoted by

$$\rho \triangleq V^T r, \quad (2.8)$$

$$\tilde{o} \triangleq V^T o. \quad (2.9)$$

The preceding equality also gives \hat{y} in terms of ρ :

$$\hat{y} = [S \quad 0] \rho. \quad (2.10)$$

With these definitions, the last equality above also implies

$$\dot{\rho} = -\rho + \tilde{o}. \quad (2.11)$$

To finish describing the basic network quantities in terms of the rotated basis, let ϵ be the error in the rotated basis:

$$\begin{aligned}
\epsilon &\triangleq y - \hat{y} \\
&= \mathcal{U}^T e.
\end{aligned} \quad (2.12)$$

4. Repeat the derivation of equation (2.5) but with y , \hat{y} , and ϵ :

$$\begin{aligned}
\dot{\epsilon} &= \dot{y} - \dot{\hat{y}} \\
&= \Lambda y + \beta c - [S \ 0] (-\rho + \tilde{o}) \\
&= \Lambda (\epsilon + [S \ 0] \rho) + \beta \tilde{c} - [S \ 0] (-\rho + \tilde{o}) \\
&= \Lambda \epsilon + (\Lambda + I) [S \ 0] \rho + \beta \tilde{c} - [S \ 0] \tilde{o} \\
\Rightarrow \begin{bmatrix} S \\ 0 \end{bmatrix} \dot{\epsilon} &= \begin{bmatrix} S \\ 0 \end{bmatrix} \Lambda \epsilon + \begin{bmatrix} S \\ 0 \end{bmatrix} (\Lambda + I) [S \ 0] \rho + \begin{bmatrix} S \\ 0 \end{bmatrix} \beta \tilde{c} - \begin{bmatrix} S \\ 0 \end{bmatrix} [S \ 0] \tilde{o}.
\end{aligned}$$

The last equality gives a system of N equations of which only d are nontrivial. A comparison with equation (2.5) suggests the N -dimensional rotated membrane potential v is best defined as:

$$v \triangleq \begin{bmatrix} S \\ 0 \end{bmatrix} \epsilon \in \mathbf{R}^N. \quad (2.13)$$

it is invertible from v to ϵ .
This mapping is not invertible unless we only consider the first d components and neglect the remaining, trivial components. Abusing notation, we write

no abuse required $\epsilon = S^{-1}v$, $E = [S^{-1} \ 0]v$
giving an N vector whose first d elements are (well defined) and the remaining components of v are assumed to be zero. Using a similar abuse for the ρ and \tilde{o} terms, we arrive at the system of d equations describing the nontrivial network voltage dynamics:

$$\begin{aligned}
\dot{v} &= S \Lambda S^{-1} v + S (\Lambda + I) S \rho + S \beta \tilde{c} - S^2 \tilde{o} \\
\Rightarrow \dot{v} &= \Lambda v + S (\Lambda + I) S \rho + S \beta \tilde{c} - S^2 \tilde{o} \quad (2.14)
\end{aligned}$$

We can also write all N dimensions explicitly to respect the dimensionality of v and ρ :

$$\dot{v} = \begin{bmatrix} \Lambda & 0 \\ 0 & 0 \end{bmatrix} v + \begin{bmatrix} S (\Lambda + I_d) S & 0 \\ 0 & 0 \end{bmatrix} \rho + \begin{bmatrix} S \\ 0 \end{bmatrix} \beta \tilde{c} - \begin{bmatrix} S^2 & 0 \\ 0 & 0 \end{bmatrix} \tilde{o}. \quad 2.14$$

To summarize conceptually, there are 4 vector spaces in total: the error space which tracks the dynamical system and the network estimate, the voltage space which tracks the membrane potentials, and the transformed counterparts of each in the $\mathcal{U} - V$ bases. Figure (2) shows the relationships derived between these subspaces.

5. The spike trains o are chosen minimize the network estimation error

$$\mathcal{L}(\xi) = \|x(\xi + d\xi) - \hat{x}(\xi + d\xi)\|^2. \quad (2.15)$$

The network greedily minimizes $\mathcal{L}(\xi)$ an instant $d\xi$ ahead in time. If no spike occurs at time ξ , then the objective is given above. If neuron j spikes, the estimate $\hat{x} \leftarrow \hat{x} + d_j$, where d_j is column j of D . The objective is now

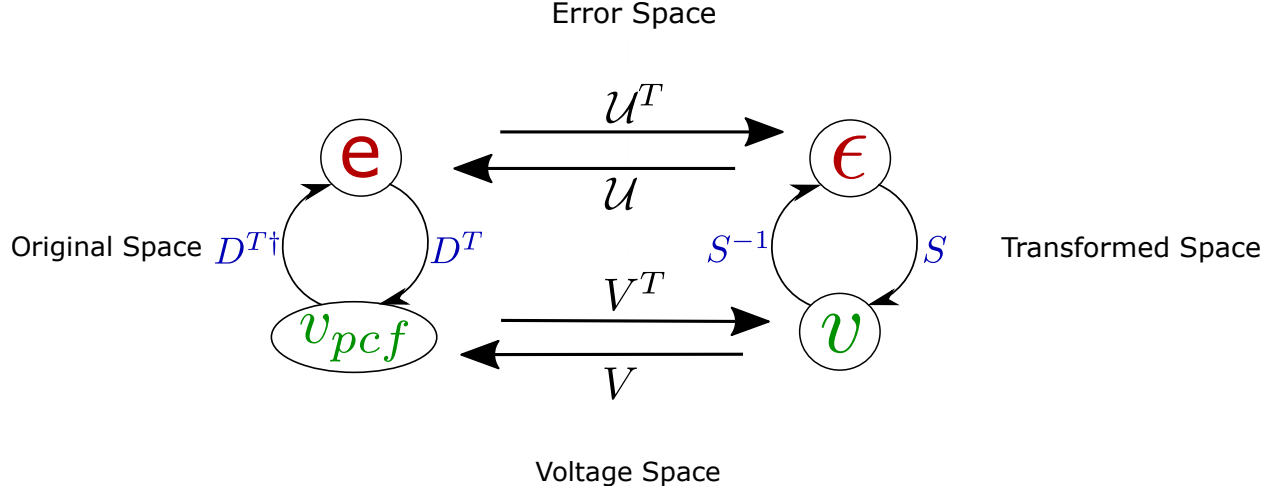


Figure 2: Depiction of the relationship between original and transformed spaces and their respective error and voltage spaces. An arrow represents left multiplication by the given matrix. The zeros in the full $N \times N$ matrices mapping between v and ϵ are omitted for clarity.

$$\begin{aligned}
\mathcal{L}_{sp}(\xi) &= \|x - (\hat{x} + d_j)\|^2 \\
&= x^T x - 2x^T \hat{x} - 2x^T d_j + \hat{x}^T \hat{x} + 2\hat{x}^T d_j + d_j^T d_j \\
&= x^T x - 2x^T \hat{x} + \hat{x}^T \hat{x} - 2d_j^T (x - \hat{x}) + d_j^T d_j \\
&= \|x - \hat{x}\|^2 - 2d_j^T (x - \hat{x}) + d_j^T d_j \\
&= \mathcal{L}_{ns}(\xi) - 2d_j^T (x - \hat{x}) + d_j^T d_j,
\end{aligned}$$

where $\mathcal{L}_{ns}(\xi)$ is the objective if no spike occurs. Spiking occurs when the objective decreases or

$$\begin{aligned}
&\mathcal{L}_{sp} < \mathcal{L}_{ns} \\
&\implies -2d_j^T (x - \hat{x}) + d_j^T d_j < 0 \\
&\implies d_j^T (x - \hat{x}) > \frac{\|d_j\|^2}{2}.
\end{aligned}$$

Since $d_j^T (x - \hat{x}) = d_j^T e$ is already defined as membrane voltage, the right hand side gives neuron j 's spike threshold voltage v_{th} ,

$$v_{th}^{pcf} = \frac{1}{2} \begin{bmatrix} d_1^T d_1 \\ \vdots \\ d_N^T d_N \end{bmatrix}.$$

For the rotated network, note \mathcal{U}^T is an orthonormal matrix by definition. Thus it is norm-preserving:

$$\begin{aligned}\mathcal{L}_{sp}(\xi) &= \|x - \hat{x}\|^2 \\ &= \|\mathcal{U}^T(x - \hat{x})\|^2 \\ &= \|y - \hat{y}\|^2.\end{aligned}$$

If we define the rotated network objective as

$$\tilde{L}(\xi) \triangleq \|y(\xi + d\xi) - \hat{y}(\xi + d\xi)\|^2,$$

it is equal to the original network objective when no spike occurs. However, a spike alters the readout by $\hat{y} \leftarrow \hat{y} + S_l$, where S_l is the l^{th} column of $[S \ 0]$. With the same approach as above, the objective when neuron l spikes is

$$\begin{aligned}\tilde{L}_{sp} &= \tilde{L}_{ns} + 2S_l^T \epsilon + S_l^T S_l \\ \implies v_l &> \frac{\|S_l\|^2}{2}.\end{aligned}$$

This leads to voltage thresholds

$$v_{th} = \frac{1}{2} \begin{bmatrix} S_1^T S_1 \\ \vdots \\ S_d^T S_d \\ 0 \\ \vdots \\ 0 \end{bmatrix}.$$

The voltage thresholds are strictly positive, such that neuron j will only spike if

$$v_l = S_l^T \epsilon > v_{th} > 0.$$

In spiking, neuron l encodes the quantity S_l into the network estimate as a correction to error ϵ . The strictly positive voltage ~~implies it is impossible for neuron l to encode in the quantity $(-S_l)$ since~~ *adds to the error* *correct antiparallel error* *addition*

$$S_l^T(\epsilon) = S_l^T(-S_l) = -\|S_l\|^2 < 0 < v_{th}.$$

To illustrate consider the space of errors $\epsilon \in \mathbf{R}^d$ which satisfy the voltage threshold of neuron l , i.e

$$\epsilon_{sp} = \{\epsilon \in \mathbf{R}^d \mid S_l^T \epsilon > v_{th}\}.$$

In \mathbf{R}^2 , ϵ_{sp} is the half-plane formed by the line normal to S_l minus the circle $\|\epsilon\|^2 > v_{th}$ as in figure (3). ~~This excludes $-S_l$.~~ *exceeds v_{th} .* *\propto* The optimization thus tells us that neuron l spikes when the projection $S_l^T \epsilon$ is nonnegative.

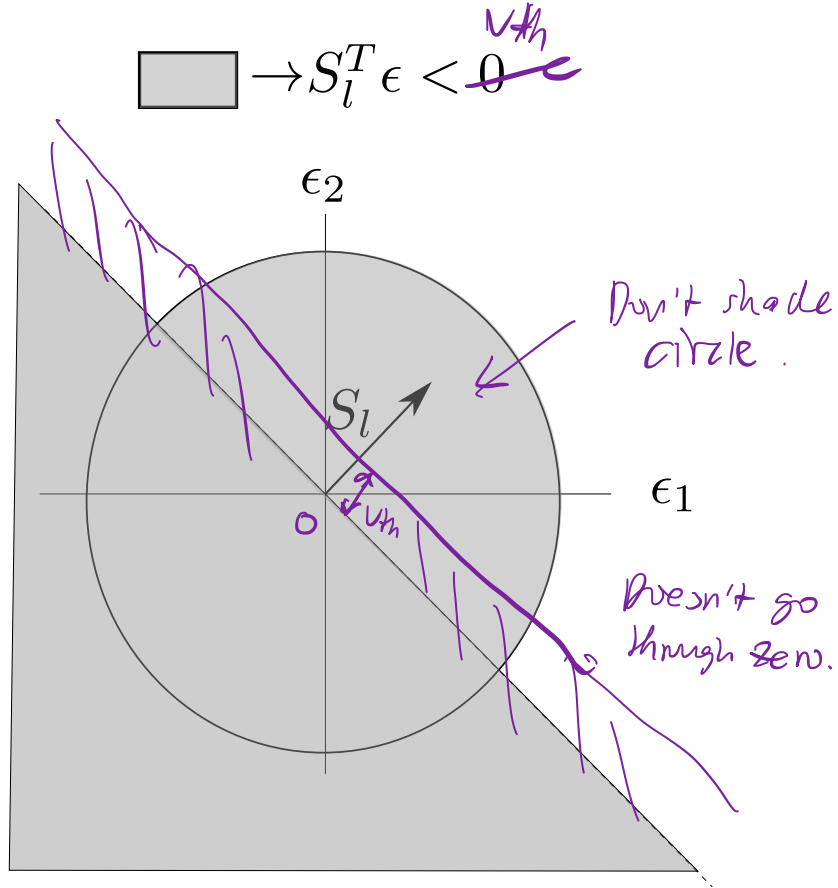


Figure 3: Sketch of the Encoding space ϵ_{sp} of neuron l with direction $S_l \in \mathbf{R}^2$. The radius of the circle is $v_{th} = \frac{\|S_l\|^2}{2}$. (Vector S_l not drawn to scale).

6. Equations (2.14) and (2.11) describe how we implement a network with d neurons that produces an accurate estimate \hat{x} of the given target system. As written, the network can only encode vectors y with strictly nonnegative elements. To see why we return to the optimization procedure performed by the network.

The network optimizes the objective

$$\mathcal{L} = ||y - \hat{y}||,$$

by choosing spike times \tilde{o} . The spikes are integrated into a post-synaptic feedback ρ . This feedback vector is scaled by S to generate the estimate \hat{y} that minimizes the objective at time ξ . In other words, the network performs the optimization

$$\min_{\rho \in \mathbf{R}^{d+}} ||y - S\rho||^2,$$

where \mathbf{R}^{d+} denotes the real d -vectors with nonnegative components. The components must be non-negative because the spikes \tilde{o} have nonnegative area when integrated, and will not decay below zero. In using a greedy approach with only one spike at a given time step, the network more specifically performs

$$\min_{x \in \mathbf{Z}^{d+} : \sum_j Z_j = 1} ||y - S(\rho + x)||^2.$$

I.e, it must choose one neuron to spike with unit area 1, which adds precisely one column of S to the estimate \hat{y} .

It is easier to analyze the former optimization of $\rho \in \mathbf{R}^{d+}$, so we do so here. Because $\{x \in \mathbf{Z}^{d+} : \sum_j Z_j = 1\} \subset \mathbf{R}^{d+}$, it is always the case that

$$\min_{\rho \in \mathbf{R}^{d+}} ||y - S\rho||^2 \leq \min_{x \in \mathbf{Z}^{d+} : \sum_j Z_j = 1} ||y - S(\rho + x)||^2,$$

i.e optimizing over arbitrary $\rho \in \mathbf{R}^{d+}$ will always give just as low or lower objectives than under the single-greedy spike optimization.

We're interested in the range of vectors representable by the network. That is the set

$$X^* = \{x \in \mathbf{R}^{d+} : Sx = y\}.$$

Over this set, the objective function is 0, i.e.

$$X^* = \{x \in \mathbf{R}^{d+} : \mathcal{L} = ||y - Sx||^2 = 0\}.$$

Let $x \in X^*$, and consider its negative $-x$. It follows that

$$\begin{aligned} x &= S\rho \\ \implies -x &= -S\rho \\ &= S(-\rho). \end{aligned}$$

However if $\rho \in \mathbf{R}^{d+}$, then it is impossible for $-\rho \in \mathbf{R}^{d+}$ to also be true. Thus $-\rho \notin X^*$ so that $\mathcal{L} > 0$. We conclude that for any vector \hat{y} , the network can represent with $\mathcal{L} = 0$, there exists a negative

vector that the network cannot represent with $\mathcal{L} = 0$. This is obviously undesirable as it restricts the set of vectors the network can reconstruct within a given error tolerance. In \mathbf{R}^2 for example, network representation where $\mathcal{L} = 0$ is restricted to the first quadrant. This restriction applies equally to the greedy single-spike optimization.

This issue is unique to the self-coupled network and does not occur in the original PCF network even though its spikes must also have positive unit area. The difference arises when we take the SVD of the decoder matrix.

$$D = U \begin{bmatrix} S & 0 \end{bmatrix} V^T.$$

The SVD decomposes D into orthonormal bases U and V which are mapped to one another by singular values S , as in figure (4). By rotating into the $U - V$ bases, we preemptively perform the first and last mappings, leaving only multiplication by a diagonal matrix. This eliminates ~~redundant neurons~~ ^{linearly dependent} encoding vectors, keeping only orthonormal ^{ones} encoding vectors. However, the SVD is agnostic to the nonnegativity restriction of its N -dimensional input vectors. A vector with opposite sign (antiparallel) to one of the orthonormal bases is a linear combination of that bases, and is considered within the range of the matrix. For example, suppose $x = D = U \begin{bmatrix} S & 0 \end{bmatrix} V^T y$. For an orthonormal basis, $-x$ is obtainable by $U \begin{bmatrix} S & 0 \end{bmatrix} V^T (-y)$. However the constraint that spikes have positive unit area prevents a vector $p = -y$ from being reachable by the network.

what does this mean? SVD applies to matrix and V is a $N \times N$ basis.
Mathematically $D = U \begin{bmatrix} S & 0 \end{bmatrix} V^T$

You are mixing bases here.
The V_i s are still orthonormal in \mathbf{R}^N
It's the D_i s that are not orthonormal in \mathbf{R}^d .

Yeah, but Denève does this even though it's also constrained to spikes with positive area.

$$D : \mathbf{R}^N \rightarrow \mathbf{R}^d = \mathcal{U} \begin{bmatrix} S & 0 \end{bmatrix} V^T$$

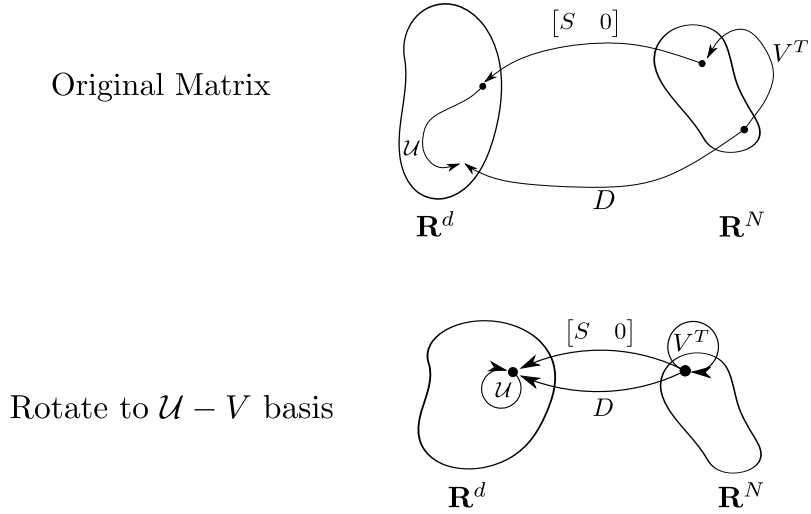


Figure 4: Visualizing D as a sequence of linear maps between subspaces. **Top:** The matrix $D \in \mathbf{R}^{d \times N}$ is decomposed via SVD into a sequence of 3 linear maps (matrices). The rightmost matrix $V^T \in \mathbf{R}^{N \times N}$ projects a vector x to give coefficients for the expansion in the basis V . The center matrix $\begin{bmatrix} S & 0 \end{bmatrix} \in \mathbf{R}^{d \times N}$ maps vectors from the V basis to a vector in \mathbf{R}^d by scaling and truncation. The leftmost matrix $\mathcal{U} \in \mathbf{R}^{d \times d}$ gives the resultant vector $Dx \in \mathbf{R}^d$ by using the scaled vector $\begin{bmatrix} S & 0 \end{bmatrix} V^T$ as coefficients for a basis expansion in \mathcal{U} . **Bottom:** We rotate the basis for vectors in \mathbf{R}^N and \mathbf{R}^d to the \mathcal{U} and V bases respectively. This negates the need of D to preemptively project and afterward rotate a vector, leaving only scaling by a diagonal matrix. The mapping D performs on a vector y simplifies to multiplication by a diagonal matrix S of y 's first d components.

For the network decoder to fully span the state space of interest, we must add anti-parallel encoding directions. One solution is to form a separate network of d neurons whose encoding directions are the antiparallel set $-\mathcal{U}$. That is, we form an identical network except the decode matrix is

$$-D = -\mathcal{U} \begin{bmatrix} S & 0 \end{bmatrix} V^T.$$

We then add the output of the two networks to recover the encoded state. PCF uses a similar approach to ensure consistency with Dale's law, which states that a neuron cannot both excite and inhibit other neurons.

No PCF does not comply with Dale's Law; a neuron can excite and inhibit at the same time.

We divide the error into its positive and negative components and encode each in a separate network. Let

$$\epsilon^+ = \epsilon \geq 0,$$

be the nonnegative component of ϵ . Note that the original estimate, e may contain negative or positive components, but the projection ϵ^+ does not. Similarly define the negative error by

$$\epsilon^- = -\epsilon < 0.$$

Note

$$\epsilon^- \neq -\epsilon^+.$$

Rather,

$$\epsilon = \epsilon^+ - \epsilon^-.$$

The relationship between ϵ^+ and ϵ^- resembles two orthogonal subspaces. The preceding relation is analogous to a direct sum.

Let

$$v^+ = S^T \epsilon^+,$$

be the voltage induced by projecting ϵ^+ onto the orthonormal bases given by $D = \mathcal{U} \begin{bmatrix} S & 0 \end{bmatrix} V^T$, and

$$v^- = S^T \epsilon^-$$

be the respective projection onto the antiparallel orthonormal bases, $D = \mathcal{U} \begin{bmatrix} -S & 0 \end{bmatrix} V^T$.

Note that

$$v^- \neq -v^+.$$

Rather,

$$v = v^+ - v^-,$$

where v is the voltage of an idealized neuron capable of positive and negative area spikes. Note that both v_j^+ and v_j^- are bounded by thresholds $v_{th} = \frac{\|S_j\|^2}{2}$, so that the idealized neuron is always within the voltage range $v \in [-v_{th}, v_{th}]$. This is equivalent to asserting the error along each encoding direction S_j is contained within the polytope $S_j^T \epsilon \leq \frac{\|S_j\|^2}{2}$.

Let ρ^+ and $\tilde{\rho}^+$ be the slow synaptic feedback and spike trains of the positive neurons, with ρ^- and $\tilde{\rho}^-$ defined similarly for the negative neurons. Finally split $\tilde{c} = \tilde{c}^+ - \tilde{c}^-$ into positive and negative components as with ϵ .

We now have two d -dimensional systems of equations.

$$\dot{v}^+ = \begin{bmatrix} \Lambda & 0 \\ 0 & 0 \end{bmatrix} v^+ + \begin{bmatrix} S(\Lambda + I_d) S & 0 \\ 0 & 0 \end{bmatrix} \rho^+ + \begin{bmatrix} S \\ 0 \end{bmatrix} \beta \tilde{c}^+ - \begin{bmatrix} S^2 & 0 \\ 0 & 0 \end{bmatrix} \tilde{o}^+,$$

$$\dot{v}^- = \begin{bmatrix} \Lambda & 0 \\ 0 & 0 \end{bmatrix} v^- + \begin{bmatrix} S(\Lambda + I_d) S & 0 \\ 0 & 0 \end{bmatrix} \rho^- + \begin{bmatrix} S \\ 0 \end{bmatrix} \beta \tilde{c}^- - \begin{bmatrix} S^2 & 0 \\ 0 & 0 \end{bmatrix} \tilde{o}^-.$$

These equations each produce estimates

$$\hat{y}^+ = S\rho^+,$$

$$\hat{y}^- = S\rho^-,$$

which give the network estimate

$$\hat{y} = \hat{y}^+ - \hat{y}^-.$$

Writing the above as a single network, assume $N = 2d$ so we need not fill with zeros:

$$\begin{bmatrix} \dot{v}^+ \\ \dot{v}^- \end{bmatrix} = \begin{bmatrix} \Lambda & 0 \\ 0 & \Lambda \end{bmatrix} \begin{bmatrix} v^+ \\ v^- \end{bmatrix} + \begin{bmatrix} S(\Lambda + I_d) S & 0 \\ 0 & S(\Lambda + I_d) S \end{bmatrix} \begin{bmatrix} \rho^+ \\ \rho^- \end{bmatrix} + \begin{bmatrix} S & 0 \\ 0 & S \end{bmatrix} \beta \begin{bmatrix} \tilde{c}^+ \\ \tilde{c}^- \end{bmatrix} - \begin{bmatrix} S^2 & 0 \\ 0 & S^2 \end{bmatrix} \begin{bmatrix} \tilde{o}^+ \\ \tilde{o}^- \end{bmatrix}.$$

We simplify this by writing

$$\dot{v} = \Lambda v + S(\Lambda + I_{2d}) S \rho + S \beta \tilde{c} - S^2 \tilde{o},$$

where we have made the following substitutions:

This may work, but it's for the next chapter. At the moment, we are analyzing the notated equivalent of Denere + G.J. That network is described by

$$\dot{v} = \frac{1}{2} \begin{bmatrix} S \\ -S \end{bmatrix} \Lambda [S^+ - S^-] v + \frac{1}{2} \begin{bmatrix} S \\ -S \end{bmatrix} (\Lambda + I) [S - S] r + \frac{1}{\sqrt{2}} \begin{bmatrix} S \\ -S \end{bmatrix} \beta \tilde{c} - \frac{1}{2} \begin{bmatrix} S \\ -S \end{bmatrix} [S - S] \tilde{o}$$

At the moment, our goal is to obtain theoretical predictions for this to compare with those for Denere and Denere + Gup-Junctions.

$$v \leftarrow \begin{bmatrix} v^+ \\ v^- \end{bmatrix} \in \mathbf{R}^{2d},$$

$$\Lambda \leftarrow \begin{bmatrix} \Lambda & 0 \\ 0 & \Lambda \end{bmatrix} \in \mathbf{R}^{2d \times 2d},$$

$$S \leftarrow \begin{bmatrix} S & 0 \\ 0 & S \end{bmatrix} \in \mathbf{R}^{2d \times 2d},$$

$$\rho \leftarrow \begin{bmatrix} \rho^+ \\ \rho^- \end{bmatrix} \in \mathbf{R}^{2d},$$

$$\tilde{o} \leftarrow \begin{bmatrix} \tilde{o}^+ \\ \tilde{o}^- \end{bmatrix} \in \mathbf{R}^{2d},$$

$$\beta \leftarrow \begin{bmatrix} \beta & 0 \\ 0 & \beta \end{bmatrix} \in \mathbf{R}^{2d \times 2d},$$

$$\tilde{c} \leftarrow \begin{bmatrix} \tilde{c}^+ \\ \tilde{c}^- \end{bmatrix} \in \mathbf{R}^{2d},$$

$$v_{th} \leftarrow \begin{bmatrix} v_{th} \\ v_{th} \end{bmatrix} \in \mathbf{R}^{2d}.$$

To decode from the network to the d -dimensional estimate, we multiply by $\begin{bmatrix} \mathcal{U} & -\mathcal{U} \end{bmatrix} \in \mathbf{R}^{d \times 2d}$, i.e

$$\hat{y} = \hat{y}^+ - \hat{y}^- = \begin{bmatrix} \mathcal{U} & -\mathcal{U} \end{bmatrix} \rho.$$

This approach suggests a balance between excitatory and inhibitory neurons when coding a signal that inhabits the full d -dimensional state space. To illustrate, consider a 2 neuron network as in figure (5). A sinusoidal input drives the neurons which encode v^+ and v^- respectively. A readout neuron performs leaky integration of the spike trains from the two driven neurons. We observe equal levels of excitatory and inhibitory input from the two neurons, suggesting a tight balance.

Note that the fast coupling matrix preceding \tilde{o} is diagonal. This implies that when a neuron j spikes, its antiparallel neuron is unchanged. In the PCF, a neuron spike resets its threshold to $-v_{th}$, but likewise sets a neuron antiparallel to it to v_{th} . Next, the antiparallel neuron spikes and likewise resets the neuron. This cycle repeats itself causing the network estimate to oscillate uncontrollably in a catastrophic network failure termed "ping-ponging". The PCF addresses this through regularization terms applied to the network objective \mathcal{L} and added noise, (Boerlin 2013) both of which are unnecessary here.

2.2 Simulation of Basic Equations

Here we simulate the above equations (2.14) and (2.11) with the $N = 2d$ neurons. The parameters are

$$\begin{aligned}
A &= - \begin{bmatrix} 1 & 0 \\ 0 & 1 \end{bmatrix} = \mathcal{U} \Lambda \mathcal{U}^T, \\
B &= \begin{bmatrix} 1 & 0 \\ 0 & 1 \end{bmatrix}, \\
c(\xi) &= \begin{bmatrix} \cos(\frac{\pi}{4}\xi) \\ \sin(\frac{\pi}{4}\xi) \end{bmatrix} \\
D &= \mathcal{U} \begin{bmatrix} S & 0 \end{bmatrix} V^T = \mathcal{U} \begin{bmatrix} .1 I_d & 0 \end{bmatrix} I_N, \\
d\xi &= 10^{-6}, \\
N &= 4, \\
x(0) &= \begin{bmatrix} \frac{1}{2} & \frac{1}{2} \end{bmatrix}.
\end{aligned} \tag{2.16}$$

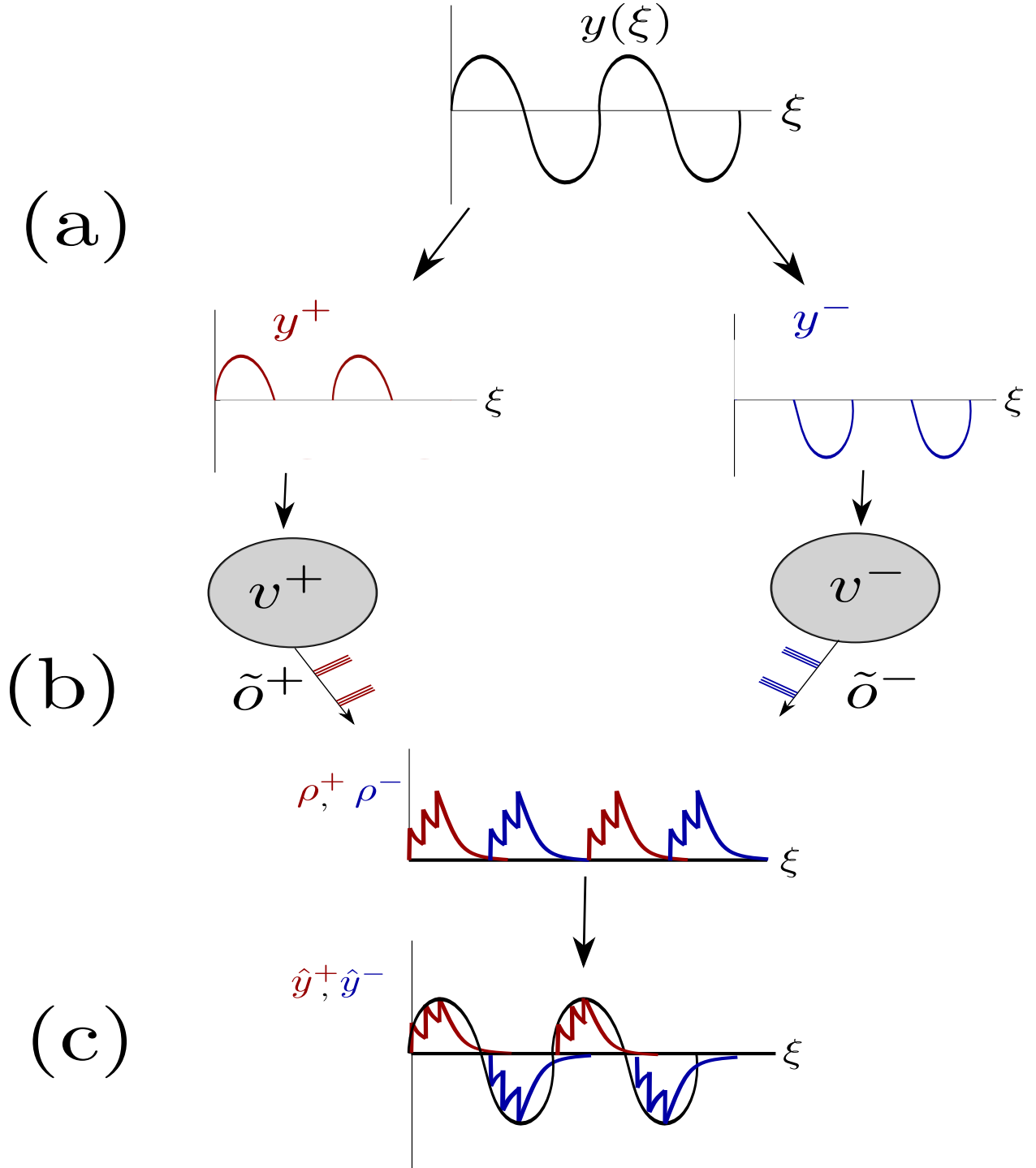


Figure 5: Balance Between Excitatory and Inhibitory Activity for Oscillatory Input. *(a)*: A sinusoidal input y is divided into its positive (excitatory) and negative (inhibitory) components. *(b)*: Each neuron encodes its respective input by spiking to produce a nonnegative filtered spike train ρ . *(c)*:

The network estimate is the sum of activity from excitatory and inhibitory filtered spike trains. For oscillatory input, both neurons must spike in equal amounts so that the amplitude of oscillation remains bounded.

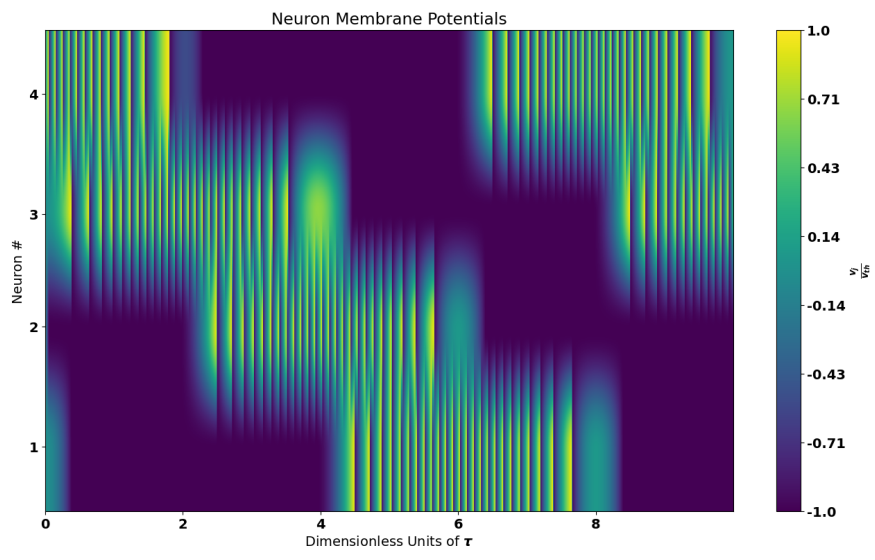
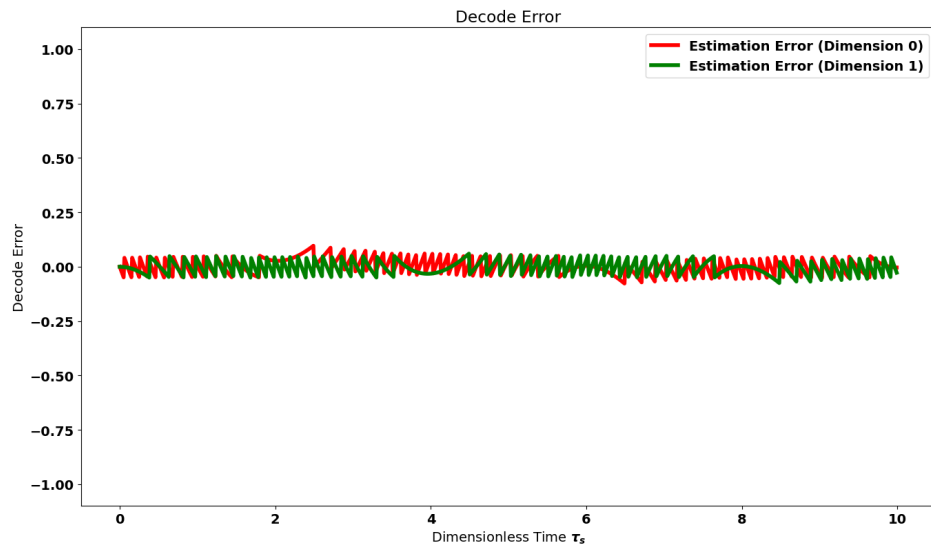
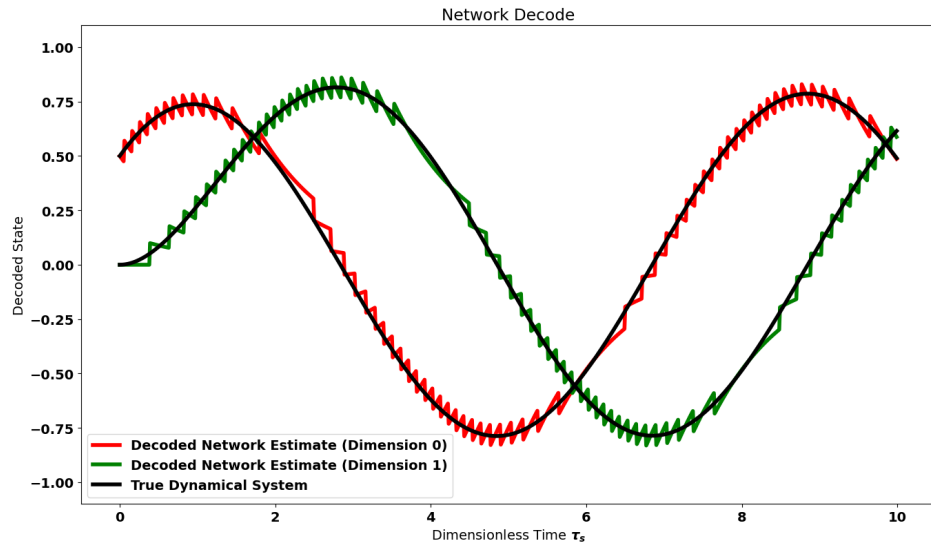


Figure 6: Simulation of equations (2.14) and (2.11) with parameters listed in equation (2.16). **Top:** The decoded network estimate plotted alongside the target dynamical system. **Middle:** The estimation error along each state-space dimension. **Bottom:** The membrane potentials of the 4 neurons during the same time period.

For the numerical implementation, the matrix exponential was used to integrate the continuous terms over a simulation time step. Continuous terms include all equation terms excepting the delta functions ω handled separately. After integrating over a timestep, any neuron above threshold was manually reset (action of fast inhibition). If multiple neurons are above threshold, the system is integrated backwards in time until only one neuron is above threshold before spiking. The matrix exponential was computed using a Padé approximation via the Python package Scipy: `scipy.linalg.expm()`.

3 Analysis: RMSE vs Spike Rate for Constant Driving Force

We analyse the network described by equations (2.14) and (2.11) for the case of a constant (in time) driving force $c(\xi) = k$. First we derive explicit expressions for the network estimate, then we compute the resulting RMSE for various driving strengths k .

1. Let

$$A = - \begin{bmatrix} 1 & 0 \\ 0 & 1 \end{bmatrix} = \Lambda,$$

$$B = \begin{bmatrix} 1 & 0 \\ 0 & 1 \end{bmatrix} = I,$$

$$D = \mathcal{U} \begin{bmatrix} S & 0 \end{bmatrix} V^T = \mathcal{U} \begin{bmatrix} I_d & 0 \end{bmatrix} I_N,$$

$$c(\xi) = k \in \mathbf{R}^d, \text{ expressed in the } \mathcal{U} - V \text{ basis,}$$

$$d\xi = 10^{-4},$$

$$N = 4,$$

$$x(0) = \begin{bmatrix} \frac{1}{2} & 0 \end{bmatrix}.$$

The system simplifies to one neuron by noting the following: The voltage of the $2d$ non trivial neurons is

$$v = S\epsilon.$$

Voltage j spikes i.f.f. $v_j = \frac{\|S_j\|^2}{2}$. Let 1 index the neuron whose encoding direction S_j is closest in angle to k , i.e

$$S_1^T e \geq S_j^T e \quad \forall j$$

$$\implies v_1 \geq v_k \quad \forall j.$$

It follows from equation (2.14) that v_1 will reach its threshold first. Neuron 1 spikes and its voltage is reset, and sequence repeats. Let us integrate until the first spike, avoiding the discontinuity from \tilde{o} . The nontrivial dynamics simplify to

$$\dot{v}_1(\xi) = \Lambda_1 v_1(\xi) + (\Lambda_1 + 1) \|S_1\|^2 \rho_1(\xi) + S_1^T k, \quad (3.1)$$

$$\dot{\rho}_1(\xi) = -\rho_1(\xi).$$

With initial conditions $v_1(0) = v_1^0$ and $\rho_1(0) = \rho_1^0$, the system of equations has the general solution

$$\begin{aligned} \rho_1(\xi) &= \rho_1^0 e^{-\xi}, \\ v_1(\xi) &= e^{\Lambda_1 \xi} \left(\frac{S_1^T k}{\Lambda_1} + \|S_1\|^2 \rho_1^0 + v_1^0 \right) - e^{-\xi} \|S_1\|^2 \rho_1^0 - \frac{S_1^T k}{\Lambda_1}. \end{aligned} \quad (3.2)$$

Between spikes, neuron 1's voltage is independent from other neurons j . The voltage v_1 only depends on its own history and feedback from its own spike train, ρ_1 . Replace index 1 we see this is true for all neurons so that all neurons are *self-coupled*, hence the name.

A spike occurs when $v(\xi_{spike}) = \frac{\|S_1\|^2}{2}$, or

$$\frac{\|S_1\|^2}{2} = e^{\Lambda_1 \xi_{spike}} \left(\frac{S_1^T k}{\Lambda_1} + \|S_1\|^2 \rho_1^0 + v_1^0 \right) - e^{-\xi_{spike}} \|S_1\|^2 \rho_1^0 - \frac{S_1^T k}{\Lambda_1}.$$

This equation is transcendental in that a closed form expression for ξ_{spike} does not exist. However, under certain initial conditions we can obtain a solution. Let $r_1^0 = v_1^0 = -v_{th} = \frac{\|S\|^2}{2}$. Moreover, for a sufficiently small angle between S_1 and k , $S_1^T k \simeq \|S_1\| \|k\|$. Under these conditions,

$$\xi_{spike} = \frac{1}{\Lambda} \ln \left(\frac{1 + \frac{\Lambda_1 \|S_1\|}{2 \|k\|}}{1 - \frac{\Lambda_1 \|S_1\|}{2 \|k\|}} \right)$$

The preceding expression is the amount of time required for neuron 1 to spike starting from its membrane reset potential. This expression is exact for the case of no slow-synaptic feedback $\rho_1^0 = 0$. The intrinsic firing rate of the neuron is the inverse:

$$\phi(s, k) \triangleq \Lambda_1 \ln \left(\frac{1 + \frac{\Lambda_1 S_1}{2 k}}{1 - \frac{\Lambda_1 S_1}{2 k}} \right)^{-1}, \quad (3.3)$$

where the vertical bars for the norms are omitted for clarity.

The network will encode the constant driving force by spiking at a fixed rate determined by equation (3.3). Figure (7) shows a plot of equation (3.3) along with numerically computed spike rates for a simulated network driven with constant drive strength ratio $\frac{\|k\|}{\|s\|}$. Similar to membrane voltage, the resulting slow feedback and readout dynamics are reduced to one neuron periodically spiking:

$$\begin{aligned} \dot{\rho}_1 &= -\rho_1 + \tilde{o}_1 \\ \implies \dot{\hat{x}} &= -S_1 \rho_1 + S_1 \tilde{o}_1 \\ &= -\hat{x} + S_1 \tilde{o}_1, \end{aligned}$$

where $S_1 \in \mathbf{R}^d$.

2. The spike train \tilde{o}_1 is a periodic sequence of impulses spaced in time by $\frac{1}{\phi}$. If the first spike occurs at ξ_1^1 , then $\tilde{o}_1(\xi) = \sum_{l=0}^{\infty} \delta \left(\xi - \xi_1^1 - \frac{l}{\phi} \right)$. The network estimate therefore has dynamics

$$\dot{\hat{x}} = -\hat{x} + S_1 \sum_{l=0}^{\infty} \delta \left(\xi - \frac{l}{\phi} \right). \quad (3.4)$$

The target dynamical system is

$$\begin{aligned} \dot{x} &= -x + k \\ x(0) &= \begin{bmatrix} \frac{1}{2} & 0 \end{bmatrix}, \end{aligned}$$

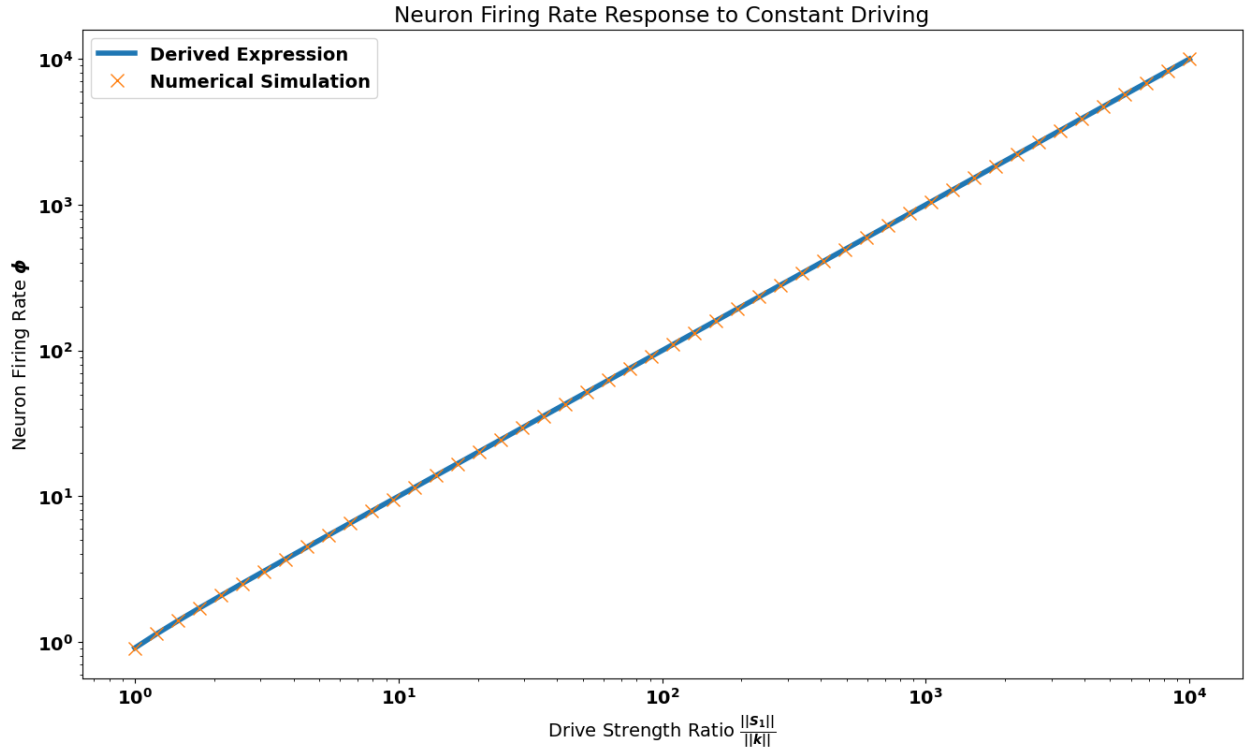


Figure 7: A log-log plot of equation (3.3) alongside the rates measured from numerical simulations. The simulation parameters are described at the beginning of this section. The rate was measured as the number of spike resets divided by the duration of the simulation.

which has a stable fixed point at

$$x = k. \quad (3.5)$$

Equation (3.4) implies that the network estimate \hat{x} will decay until the first spike ξ_1^1 occurs:

$$\hat{x}(\xi) = x(0)e^{-\xi}, \quad 0 \leq \xi < \frac{1}{\phi}.$$

At this instant, the vector S_1 is added to the network estimate.

$$\hat{x}\left(\frac{1}{\phi}\right) = x(0)e^{-\frac{1}{\phi}} + S_1.$$

Decay again occurs until the next spike

$$\begin{aligned} \hat{x}(\xi) &= \hat{x}\left(\frac{1}{\phi}\right)e^{-(\xi - \frac{1}{\phi})}, \\ &= \left(x(0)e^{-\frac{1}{\phi}} + S_1\right)e^{-(\xi - \frac{1}{\phi})}, \quad \frac{1}{\phi} \leq \xi < \frac{2}{\phi} \\ \implies \hat{x}\left(\frac{2}{\phi}\right) &= \left(x(0)e^{-\frac{1}{\phi}} + S_1\right)e^{-\left(\frac{1}{\phi}\right)} + S_1 \\ &= x(0)e^{-\frac{2}{\phi}} + S_1e^{-\frac{1}{\phi}} + S_1. \end{aligned}$$

The third spike more clearly shows the recursive behavior

$$\begin{aligned} \hat{x}\left(\frac{3}{\phi}\right) &= \left[x(0)e^{-\frac{2}{\phi}} + S_1e^{-\frac{1}{\phi}} + S_1\right]e^{-\frac{1}{\phi}} + S_1 \\ &= x(0)e^{-\frac{3}{\phi}} + S_1e^{-\frac{2}{\phi}} + S_1e^{-\frac{1}{\phi}} + S_1 \end{aligned}$$

Let us consider the n^{th} spike sufficiently far from $\xi = 0$ such that the transient term $x(0)e^{-\frac{n}{\phi}}$ can be neglected. This leads to the expression

$$\begin{aligned} \hat{x}\left(\frac{n}{\phi}\right) &= \sum_{l=0}^{n-1} S_1 e^{-\frac{l}{\phi}} \\ &= S_1 \frac{1 - e^{-\frac{n}{\phi}}}{1 - e^{-\frac{1}{\phi}}}. \end{aligned}$$

For sufficiently large n , this converges to

$$\hat{x}(\xi_1^n) = \frac{S_1}{1 - e^{-\frac{1}{\phi}}}. \quad (3.6)$$

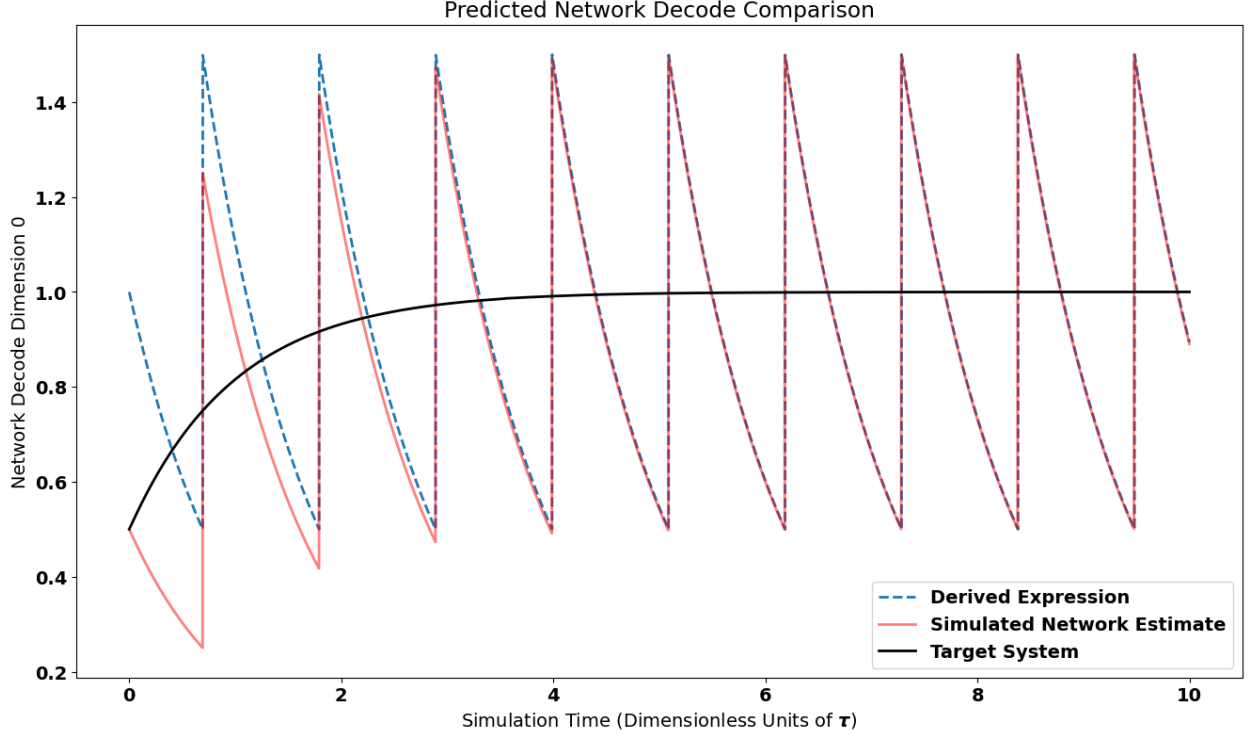


Figure 8: Comparison of the derived long-term network estimate equation (3.8) to numerical simulation. Parameters are the same as the previous figure, with $\frac{\|S_1\|}{\|k\|} = 1$.

3. The preceding argument states that after a transient interval, the network estimate at any spike time ξ_1^n is given by equation (3.6). As shown in figure (8), this convergence occurs after roughly 5 spikes under the given parameters.

We know from equation (3.4) that the estimate will decay exponentially from this value over an interval $\frac{1}{\phi}$ until a spike returns it to the initial value. Thus the network estimate between two consecutive spikes is given by

$$\hat{x}(\xi) = \frac{S_1}{1 - e^{-\frac{1}{\phi}}} e^{-(\xi - \xi_1^n)}, \quad 0 \leq \xi - \xi_1^n < \frac{1}{\phi}.$$

Combine this expression with equation (3.6), we have an explicit expression for the long-term behavior of the network estimate given by

$$\hat{x}(\xi) = \frac{S_1}{1 - e^{-\frac{1}{\phi}}} e^{-(\xi - \xi_1^1) \bmod \frac{1}{\phi}}, \quad (3.7)$$

where $x \bmod y$ denotes the fractional remainder of x after division by y .

Writing this equation in terms of provided network parameters, we use (3.3), to first obtain:

$$e^{-\frac{1}{\phi}} = \left(\frac{2k - \Lambda_1 S_1}{2k + \Lambda_1 S_1} \right)^{1/\Lambda_1},$$

which then gives

$$\hat{x}(\xi) = \frac{S_1}{\left(\frac{2k-\Lambda_1 S_1}{2k+\Lambda_1 S_1}\right)^{1/\Lambda_1}} e^{-(\xi-\xi_1^1) \bmod \left(\frac{2k+\Lambda_1 S_1}{2k-\Lambda_1 S_1}\right)^{1/\Lambda_1}}. \quad (3.8)$$

4. Assume the true system dynamics have settled to their fixed point $x = k$. From equation (3.8) the network estimate \hat{x} and therefore error $e = x - \hat{x}$ is a periodic function of ξ with period $\frac{1}{\phi}$. The RMSE over any integer number of spike periods is easily calculated from the RMSE over a single spike period. We compute the per-spike RMSE of the error signal e by

$$RMSE_{spike} \triangleq \sqrt{\phi \int_0^{\frac{1}{\phi}} \|e(\tau)\|^2 d\tau}. \quad (3.9)$$

The integrand $\|e(\tau)\|^2$ simplifies to

$$\begin{aligned} e^T e &= (x - \hat{x})^T (x - \hat{x}) \\ &= x^T x - 2x^T \hat{x} + \hat{x}^T \hat{x} \\ &= \|k\|^2 - 2S_1^T k \frac{e^{-\tau}}{1 - e^{-\frac{1}{\phi}}} + \|S_1\|^2 \left(\frac{e^{-\tau}}{1 - e^{-\frac{1}{\phi}}} \right)^2 \\ &= \|k\|^2 - 2\|S_1\| \|k\| \frac{e^{-\tau}}{1 - e^{-\frac{1}{\phi}}} + \|S_1\|^2 \left(\frac{e^{-\tau}}{1 - e^{-\frac{1}{\phi}}} \right)^2. \end{aligned}$$

Note that

$$\int_0^{\frac{1}{\phi}} e^{-\tau} d\tau = 1 - e^{-\frac{1}{\phi}},$$

while

$$\begin{aligned} \int_0^{\frac{1}{\phi}} (e^{-\tau})^2 d\tau &= \frac{1 - e^{-\frac{2}{\phi}}}{2} \\ &= \frac{1}{2} \left(1 - e^{-\frac{1}{\phi}} \right) \left(1 + e^{-\frac{1}{\phi}} \right). \end{aligned}$$

Therefore the integral is

$$\phi \int_0^{\frac{1}{\phi}} \|e(\tau)\|^2 d\tau = \|k\|^2 - 2\phi \|S_1\| \|k\| + \phi \frac{\|S_1\|^2}{2} \frac{1 + e^{-\frac{1}{\phi}}}{1 - e^{-\frac{1}{\phi}}}.$$

The per-spike RMSE of the network estimate is thus

$$RMSE_{spike}(k, S_1, \phi) = \sqrt{\|k\|^2 - 2\phi \|S_1\| \|k\| + \phi \frac{\|S_1\|^2}{2} \frac{1 + e^{-\frac{1}{\phi}}}{1 - e^{-\frac{1}{\phi}}}}. \quad (3.10)$$

To write the RMSE explicitly as a function of given parameters S_1, k, Λ_1 , we substitute our earlier expression for $e^{-\frac{1}{\phi}}$ and use equation (3.3) to obtain

$$RMSE_{spike}(k, S_1, \Lambda_1) = \sqrt{\|k\|^2 - 2 \frac{\Lambda_1 \|S_1\| \|k\|}{\ln\left(\frac{2\|k\| + \Lambda_1 \|S_1\|}{2\|k\| - \Lambda_1 \|S_1\|}\right)} + \frac{\Lambda_1}{\ln\left(\frac{2\|k\| + \Lambda_1 \|S_1\|}{2\|k\| - \Lambda_1 \|S_1\|}\right)} \frac{\|S_1\|^2}{2} \left(\frac{1 + \left(\frac{2\|k\| - \Lambda_1 \|S_1\|}{2\|k\| + \Lambda_1 \|S_1\|}\right)^{1/\Lambda_1}}{1 - \left(\frac{2\|k\| - \Lambda_1 \|S_1\|}{2\|k\| + \Lambda_1 \|S_1\|}\right)^{1/\Lambda_1}} \right)} \quad (3.11)$$

5. For the case where $\Lambda_1 = -1$ and $S_1 = 1$, we can solve for the per-spike RMSE as a simple function of ϕ . First note that from equation (3.3),

$$\frac{\|k\|}{\|S_1\|}(\phi) = \frac{1}{2} \frac{1 + e^{-\frac{1}{\phi}}}{1 - e^{-\frac{1}{\phi}}}.$$

From the preceding expression equation (3.11) simplifies to

$$RMSE_{spike}(k, S_1, \phi) = \sqrt{\|k\|^2 - \phi \|S_1\| \|k\|}.$$

If we normalize by the driving force $\|k\|$, we can isolate the change in network accuracy due to the intrinsic neuron firing rate. Divide the preceding expression by $\|k\|$ to get

$$NRMSE_{spike}(\phi) = \sqrt{1 - 2\phi \frac{1 - e^{-\frac{1}{\phi}}}{1 + e^{-\frac{1}{\phi}}}}. \quad (3.12)$$

Equations (3.11) and (3.12) are plotted in figure (9). Note that the drive strength varies the amplitude of the target system's steady state. Thus we have derived the the network performance over its dynamic range of representable state space.

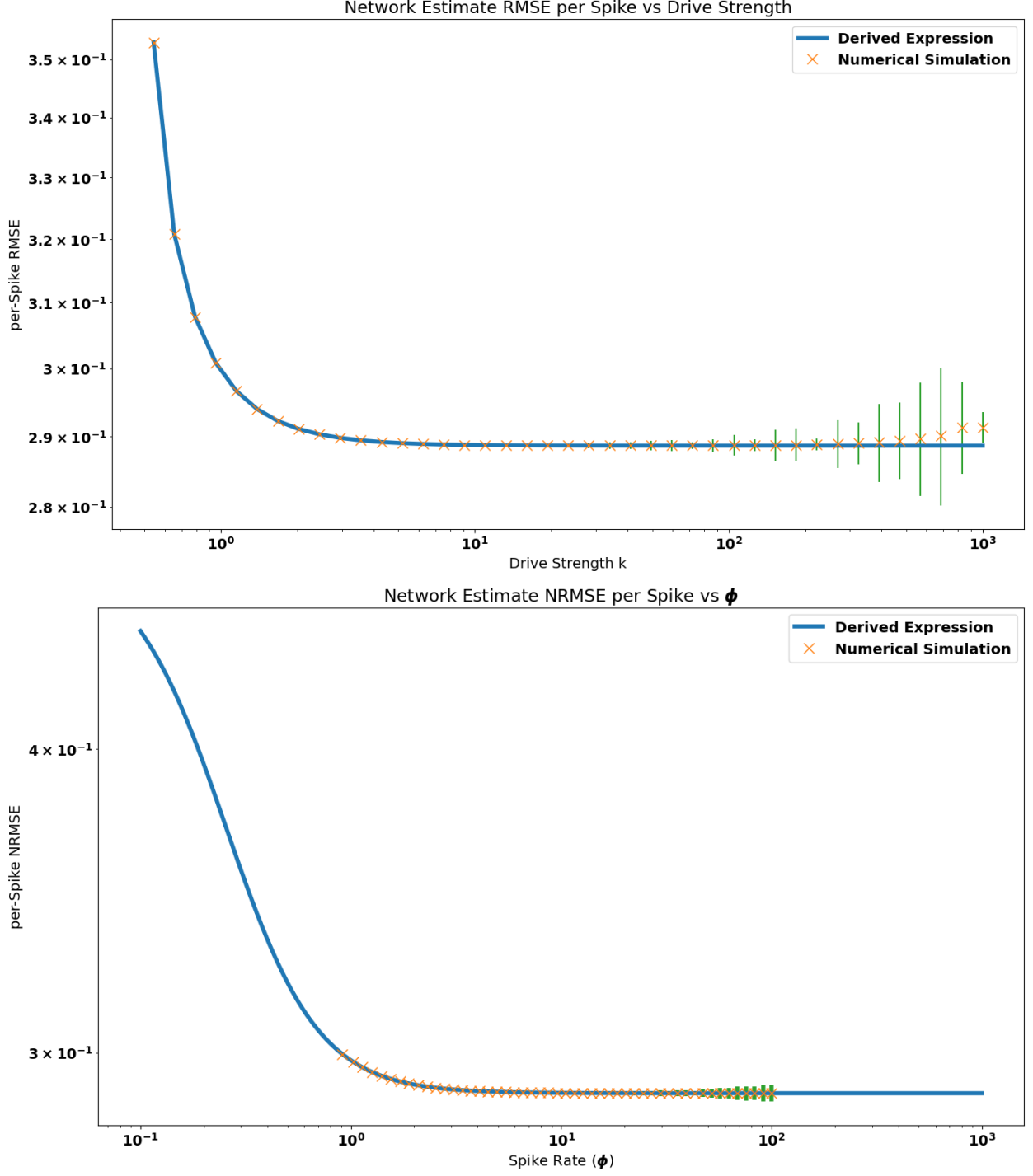


Figure 9: **Top:** A log-log plot of equation (3.11). **Bottom:** A log-log plot of equation (3.12). **Both:** Each simulated data point is the RMSE averaged over all inter-spike intervals in a simulation of length $T = 80\tau_s$ at a constant (in time) drive strength. Between simulations, the spike rates were varied by sweeping drive strength. Green vertical lines towards the larger values are ± 1 standard deviation. The spike rates $\hat{\phi}$ were computed numerically via dividing the number of spikes in a simulation by the simulation duration. The RMSE between two adjacent spikes was computed by numerical integration as a discrete sum: $RMSE = \sqrt{\hat{\phi} \sum_{\tau \text{ between spikes}} e(\xi)^T e(\xi) d\xi}$.

4 Derivation: The Predictive Coding Framework and Gap-Junction Network

Here we derive the a form of the predictive coding framework (PCF) as defined in Boerlin & Deneve, 2013. We note an assumption in this model that we later show leads to errant behavior in the network estimate. The correction of this assumption produces an intermittent mode featuring direct membrane voltage coupling. We loosely term this a gap-junction network. We compare the network estimate of all three models (PCF, gap-junction, and self-coupled) for the case of a constant driving stimulus.

1. **The Predictive Coding Framework (PCF):** The PCF synthesizes a spiking neural network that implements a given dynamical system. It is briefly derived as follows:

Assume the following are given:

- A Linear Dynamical System $\dot{x}(\xi) = Ax(\xi) + Bc(\xi)$, $x \in \mathbf{R}^d$
- A Decoder Matrix $D \in \mathbf{R}^{d \times N}$ specifying The tuning curve of N neurons in d-dimensional space.

Let $o(t) \in \mathbf{R}^N$ describe the spike trains whose j^{th} component is given by

$$o_j(t) \triangleq \sum_{k=0}^{\infty} \delta(t - t_j^k),$$

where t_j^k is the time of the k^{th} spike of neuron j . Define the time-varying firing rate of the neurons by

$$\frac{dr}{dt}(t) \triangleq -\tau_s^{-1}r(t) + \tau_s^{-1}o(t),$$

where τ_s^{-1} is the decay rate of $r(t)$ given by the inverse synaptic time constant τ_s . For consistency across models, we transform the preceding two equations to dimensionless time via $\xi = \frac{t}{\tau_s} \implies \tau_s d\xi = dt$. This gives

$$o_j(\xi) \triangleq \sum_{k=0}^{\infty} \delta(\xi - \xi_j^k), \tag{4.1}$$

where ξ_j^k is the k^{th} spike of neuron j in dimensionless time, and

$$\frac{dr}{dt}(t) = -\tau_s^{-1}r(t) + \tau_s^{-1}o(t),$$

$$\implies \frac{dr}{\tau_s d\xi}(\xi) = -\tau_s^{-1}r(\xi) + \tau_s^{-1}o(\xi),$$

$$\implies \frac{dr}{d\xi}(\xi) = -r(\xi) + o(\xi).$$

Letting $\dot{}$ denote differentiation w.r.t. dimensionless time ξ , we arrive at

$$\dot{r}(\xi) \triangleq -r(\xi) + o(\xi). \tag{4.2}$$

The network estimate is defined as

$$\hat{x}(\xi) \triangleq Dr(\xi), \tag{4.3}$$

which gives rise to the network estimation error

$$e(\xi) \triangleq x(\xi) - \hat{x}(\xi). \quad (4.4)$$

The network chooses spike times ξ_j^k to greedily optimize the objective function

$$\mathcal{L}(\xi) = \|x(\xi + d\xi) - \hat{x}(\xi + d\xi)\|^2.$$

The PCF features regularized rate terms $r(\xi)$ for the sake of biological plausibility. At present we ignore these terms. They only increase the network estimation error e by sacrificing accuracy to minimize $r(\xi)$. Using an identical approach to the derivation of the self-coupled network in section (2), we arrive at

$$d_j^T (x - \hat{x}) = \frac{d_j^T d_j}{2}$$

where d_j is the j^{th} column of D . We define membrane voltage to get the spiking condition:

$$v_j \triangleq d_j^T (x - \hat{x}) \quad (4.5)$$

$$\implies d_j^T e = v_{th},$$

where $v^{th} = \frac{d_j^T d_j}{2}$.

Deriving the dynamics, the preceding equation defines voltage, which in matrix form is given by

$$\begin{aligned} V &= D^T (x - \hat{x}) \\ \implies \dot{V} &= D^T \dot{x} - D^T \dot{\hat{x}} \\ &= D^T (Ax + Bc) - D^T (D\dot{r}) \\ &= D^T Ax + D^T Bc - D^T D(-r + o). \end{aligned}$$

The PCF makes the assumption that when the network performs correctly, $x = \hat{x}$. We later quantify the estimation error introduced by this assumption and correct it to form the gap-junction model. For now make the assumed substitution $x = \hat{x} = Dr$.

$$\begin{aligned} \dot{V} &= D^T A(Dr) + D^T Bc + D^T Dr - D^T Do \\ &= D^T (A + I) Dr + D^T Bc - D^T Do. \end{aligned}$$

The model is finalized by the addition of a voltage leakage term to ensure stability, giving the final dynamics equation

$$\dot{V} = -v + D^T (A + I) Dr + D^T Bc - D^T Do. \quad (4.6)$$

Equation (4.6) scales the spike train o_j by $d_j^T d_j$. Thus the spiking behavior is described by

$$\begin{aligned}
v_{th} &= \frac{d_j^T d_j}{2} \\
&\text{if } v_j > v_j^{th}, \\
&\text{then } v_j' = v_j - d_j^T d_j \int \delta(\tau) d\tau, \\
&\text{and } r_j' = r_j + \int \delta(\tau) d\tau.
\end{aligned} \tag{4.7}$$

Equations (4.6) and (4.7) specify the PCF model we compare against. Figure (10) shows simulations of the PCF model with the following parameters:

$$\begin{aligned}
A &= - \begin{bmatrix} 1 & 0 \\ 0 & 1 \end{bmatrix}, \\
B &= \begin{bmatrix} 1 & 0 \\ 0 & 1 \end{bmatrix}, \\
c(\xi) &= 10 \begin{bmatrix} \cos(\frac{\pi}{2}\xi) \\ \sin(\frac{\pi}{2}\xi) \end{bmatrix} + 8 \\
D_{ij} &\sim \mathcal{N}(0, 1) \text{ Columns Normalized to Unit Length} \\
d\xi &= 10^{-5}, \\
N &= 32, \\
x(0) &= \begin{bmatrix} \frac{1}{2} & \frac{1}{2} \end{bmatrix}.
\end{aligned} \tag{4.8}$$

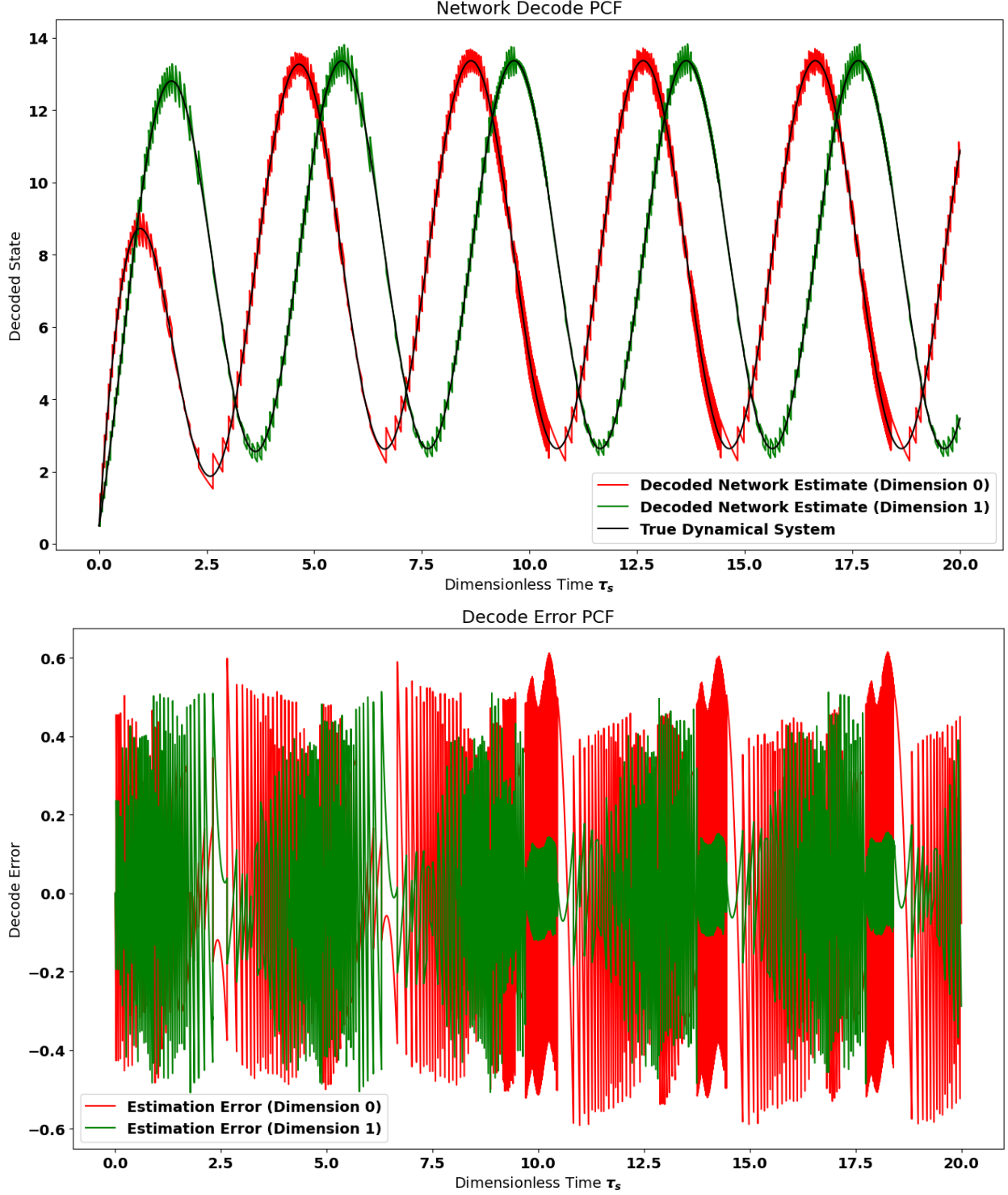


Figure 10: Simulation of PCF model given by equations (4.7) and (4.6). **Top:** Network estimate given by equation (4.3). **Bottom:** Estimation Error for PCF network from equation (4.4). The simulation parameters are given in equation (4.8). The numerical implementation is identical to that in section (2). A Padé approximation is used to compute a matrix exponential, then used to integrate the continuous terms of the differential equations. The spikes are handled separately at each time step by manually changing the values of neurons above threshold. For reasons of numerical stability, only one spike per time-step is allowed in the PCF model.

2. **The Gap-Junction Correction:** Here we correct the assumption that $\hat{x} = x$ made in the PCF model. We restart the previous derivation from this point and derive more a accurate form of equation (4.6) termed the gap-junction model. The derivation is identical as the PCF until we derive the voltage dynamics.

$$\dot{V} = D^T A x + D^T B c + D^T D r - D^T D o.$$

Instead of assuming $x = \hat{x}$, we apply the definition of voltage, equation (4.5) in matrix form.

$$\begin{aligned} v_j &= d_j^T e \\ \implies V &= D^T e \\ &= D^T (x - \hat{x}) \\ \implies x &= D^{T\dagger} V + \hat{x} \\ &= D^{T\dagger} V + D r, \end{aligned}$$

where $D^{T\dagger}$ is the left Moore-Penrose pseudo-inverse of D^T . Substitute this for x in \dot{V} above to get

$$\begin{aligned} \dot{V} &= D^T A (D^{T\dagger} V + D r) + D^T D r + D^T B c - D^T D o \\ \implies \dot{V} &= D^T A D^{T\dagger} V + D^T (A + I) D r + D^T B c - D^T D o. \end{aligned} \tag{4.9}$$

Equation (4.9) in conjunction with an identical spiking rule from PCF, equation (4.7) specifies the gap-junction model. It is simulated in figure (11). While the two simulations are similar, there are noticeable differences in their behavior e.g. $\tau_s \simeq 10, 13$.

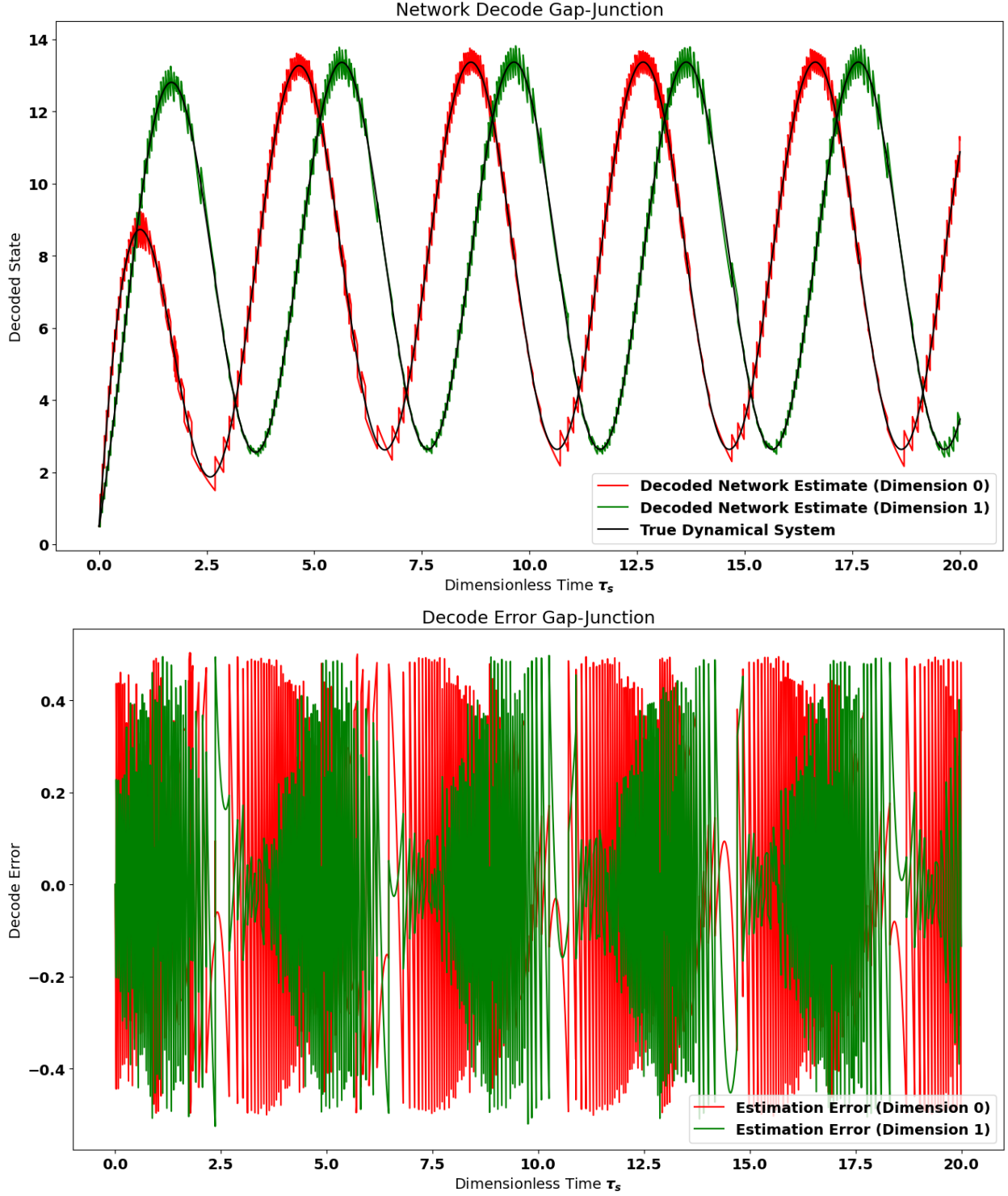


Figure 11: Simulation of the Gap-Junction model given by equations (4.7) and (4.9). **Top:** Network estimate given by equation (4.3). **Bottom:** Estimation Error for the Gap-Junction network from equation (4.4). The simulation parameters are the same as the previous figure. As with the PCF model, the network is only numerically stable if spikes are restricted to one per time step.

5 Analysis: PCF and Gap-Junction Response to Constant Stimulus

We compare the network estimate of all three models (PCF, gap-junction, and self-coupled) for the case of a constant driving stimulus.

Let all 3 models have the same parameters as given by equation (4.8) with the exception that

$$c(\xi) = c = \begin{bmatrix} 1 \\ 0 \end{bmatrix},$$

and $x(0) = [\frac{1}{2} \ 0]$.

5.1 PCF Network Response to Constant Stimulus:

From equation (4.6), the PCF dynamics become

$$\begin{aligned} \dot{V}_{pcf} &= -V_{pcf} + D^T (-I + I) D^T r + D^T \begin{bmatrix} 1 \\ 0 \end{bmatrix} - D^T D o \\ &= -V_{pcf} + D^T \begin{bmatrix} 1 \\ 0 \end{bmatrix} - D^T D o. \end{aligned}$$

All voltages are initially 0. From equation (4.7) the thresholds are identically $\frac{1}{2}$. Until the first spike, neuron j 's voltage integrates the quantity $d_j^T \begin{bmatrix} 1 \\ 0 \end{bmatrix}$. Denote the neuron j whose tuning curve d_j is closest in angle to c by

$$j_{max} \triangleq \underset{i \in [1, \dots, N]}{\operatorname{argmax}} \ d_j^T c.$$

Neuron j_{max} will receive the highest driving force and will therefore reach its threshold before any other neuron. It will then be reset by 1 to $-\frac{1}{2}$. Each other neuron k will also be reset (decremented) by $d_k^T d_{j_{max}}$, proportional to their angle relative to both neuron j_{max} and the driving strength c . This sequence will repeat periodically so that only neuron j_{max} fires at a constant rate.

We write the PCF network as the one-dimensional equation

$$v_{pcf} = -v_{pcf} + d_{j_{max}}^T c - o_{j_{max}}.$$

This is a form of the leaky integrate-and-fire (LIF) model, with drive term $d_j^T c(\xi)$. The neuron is driven by inner product $d_{j_{max}}^T c$. Note from equation (4.7) that the threshold voltage varies with $\|d_{j_{max}}\|^2$. For clarity, we drop the subscripts j, j_{max} in the following equations. It is understood that we are referring to the solely spiking neuron j_{max} . With initial condition $v_{pcf}(0) = -\frac{\|d\|^2}{2}$, the neuron's trajectory is integrated as

$$v_{pcf}(\xi) = d^T c - e^{-\xi} \left(d^T c + \frac{\|d\|^2}{2} \right). \quad (5.1)$$

The neuron spikes when it reaches the threshold $v_{pcf} = \|d\|^2$. To compare with the self-coupled network, we note that the singular value associated with neuron j of the decoder matrix $S = \|d\|^2$.

From the preceding equation with voltage at threshold $\frac{\|d\|^2}{2}$,

$$\begin{aligned}\frac{\|d\|^2}{2} &= d^T c - e^{-\xi_{spike}} \left(d^T c + \frac{\|d\|^2}{2} \right) \\ \implies e^{-\xi_{spike}} &= \frac{d^T c - \frac{\|d\|^2}{2}}{d^T c + \frac{\|d\|^2}{2}} \\ \implies \xi_{spike} &= \ln \left(d^T c + \frac{\|d\|^2}{2} \right) - \ln \left(d^T c - \frac{\|d\|^2}{2} \right)\end{aligned}$$

This leads to a firing rate

$$\phi_{pcf}(d) = \frac{1}{\ln \left(d^T c + \frac{\|d\|^2}{2} \right) - \ln \left(d^T c - \frac{\|d\|^2}{2} \right)} \quad (5.2)$$

Deriving the network estimate, suppose it begins at $x(0)$. The trajectory until the first spike at time ξ_1 is

$$\hat{x}(\xi) = x(0)e^{-\xi}, \quad 0 \leq \xi < \xi_1.$$

The spike adds d to the readout followed by exponential decay:

$$\hat{x}(\xi) = (x(0)e^{-\xi_1} + d) e^{-(\xi - \xi_1)}, \quad 0 \leq \xi - \xi_1 < \frac{1}{\phi}.$$

Until the third spike the readout is

$$\hat{x}(\xi) = \left(x(0)e^{-\xi_1} e^{-\frac{1}{\phi}} + d e^{-\frac{1}{\phi}} + d \right) e^{-(\xi - \frac{1}{\phi} - \xi_1)}, \quad \frac{1}{\phi} \leq \xi - \xi_1 < \frac{2}{\phi}.$$

The recursive pattern is visible after the third spike

$$\hat{x}(\xi) = \left(x(0)e^{-\xi_1} e^{-\frac{2}{\phi}} + d e^{-\frac{2}{\phi}} + d e^{-\frac{1}{\phi}} + d \right) e^{-(\xi - \frac{2}{\phi} - \xi_1)}, \quad \frac{2}{\phi} \leq \xi - \xi_1 < \frac{3}{\phi}.$$

Consider the n^{th} term for n big enough so that the $x(0)$ term is approximately 0. The readout at spike time ξ_n is given by the sum

$$\hat{x}(\xi_n) = d \sum_{l=0}^{n-1} e^{-\frac{l}{\phi}}.$$

The series converges to

$$\hat{x}(\xi_n) = \frac{d}{1 - e^{-\frac{1}{\phi}}}.$$

Between the spikes the readout exponentially decays so that the network estimate is given by

$$\hat{x}_{pcf}(\xi) = \frac{d}{1 - e^{-\frac{1}{\phi}}} e^{-\left(\xi - \xi_1^1 \right) \bmod \frac{1}{\phi}}. \quad (5.3)$$

5.2 Gap-Junction Network Response to Constant Stimulus:

Here we derive the decoded estimate of a gap-junction network driven by a constant stimulus, $c(\xi) = \begin{bmatrix} 1 & 0 \end{bmatrix}$. All other parameters are identical to those in equation (4.8).

From the dynamics equation (4.9) gap-junction voltages are continuously coupled to one another via $D^T A D^{T\dagger}$. We thus need to solve the entire system between spikes rather than reducing it to a single dimension. Let $\tilde{\cdot}$ denote the Laplace transform of a variable. Assume neuron j has just spiked so that

$$V(0) = -\frac{1}{2} \begin{bmatrix} d_1^T d_j \\ \vdots \\ d_j^T d_j \\ \vdots \\ d_N^T d_j \end{bmatrix}.$$

Since $c(\xi) = \begin{bmatrix} 1 \\ 0 \end{bmatrix}$, $B = I$, and $o(\xi) = 0$ between spikes, we have

$$\dot{V} = D^T A D^{T\dagger} V + D^T (A + I) D r + d_1,$$

where d_1 is the first column of D . Apply the one-sided Laplace Transform to both sides and use the Laplace derivative property:

$$\begin{aligned} s\tilde{V} - V(0) &= D^T A D^{T\dagger} \tilde{V} + D^T (A + I) o D \tilde{r} + \mathcal{L}[d_1] \\ \implies (sI - D^T A D^{T\dagger}) \tilde{V} &= V(0) + D^T (A + I) D \tilde{r} + \tilde{d}_1 \\ \implies \tilde{V} &= (sI - D^T A D^{T\dagger})^{-1} [V(0) + D^T (A + I) D \tilde{r} + \tilde{d}_1] \\ &= (sI - D^T A D^{T\dagger})^{-1} V(0) + (sI - D^T A D^{T\dagger})^{-1} D^T (A + I) D \tilde{r} + \tilde{d}_1. \end{aligned}$$

Now apply the inverse Laplace transform. Note that by definition of matrix exponential,

$$\mathcal{L}^{-1} (sI - D^T A D^{T\dagger})^{-1} = e^{\xi D^T A D^{T\dagger}}.$$

Therefore,

$$\begin{aligned} V(\xi) &= e^{\xi D^T A D^{T\dagger}} V(0) + \mathcal{L}^{-1} \left[(sI - D^T A D^{T\dagger})^{-1} D^T B \tilde{c} \right] \\ &\quad + \mathcal{L}^{-1} \left[(sI - D^T A D^{T\dagger})^{-1} D^T (A + I) D \tilde{r} \right]. \quad (5.4) \end{aligned}$$

To simplify the second term of equation (5.4), use the convolution-product property of the Laplace transform to get

$$\mathcal{L}^{-1} \left[(sI - D^T A D^{T\dagger})^{-1} D^T B \tilde{c} \right] = \mathcal{L}^{-1} \left[(sI - D^T A D^{T\dagger})^{-1} \right] * \mathcal{L}^{-1} [D^T B \tilde{c}].$$

Note $c(\xi) = \begin{bmatrix} 1 \\ 0 \end{bmatrix}$, and $B = I$. Therefore,

$$\mathcal{L}^{-1} [D^T B \tilde{c}] = D^T B c = d_1,$$

where d_1 is the first column of D . The entire second term in $V(\xi)$ above becomes

$$\mathcal{L}^{-1} \left[(sI - D^T A D^{T\dagger})^{-1} D^T B \tilde{c} \right] = e^{\xi D^T A D^{T\dagger}} * d_1.$$

Evaluating the convolution, bring d_1 outside the integral, a linear operator:

$$e^{\xi D^T A D^{T\dagger}} * d_1(\xi) = \int_{\tau=-\infty}^{\infty} e^{(\xi-\tau) D^T A D^{T\dagger}} d\tau d_1.$$

The state $V(\xi)$ depends only on the past up to $V(0)$ so that $0 < \xi - \tau \leq \xi$:

$$e^{\xi D^T A D^{T\dagger}} * d_1(\xi) = \int_{\tau=0}^{\xi} e^{(\xi-\tau) D^T A D^{T\dagger}} d\tau d_1.$$

The integral of the matrix exponential $\int_{t=0}^T e^{tX} dt = X^{-1} (e^{Tx} - I)$. Thus,

$$\mathcal{L}^{-1} \left[(sI - D^T A D^{T\dagger})^{-1} D^T B \tilde{c} \right] = (D^T A D^{T\dagger})^{-1} \left(e^{\xi D^T A D^{T\dagger}} - I \right) d. \quad (5.5)$$

Note the notation $d = D^T B c$. Looking at the final term of equation (5.4), assume the network estimate is periodic with period $\frac{1}{\phi}$, where ϕ is the unknown spike rate. Between spikes, the dynamics of $r(\xi)$ are known from equation (4.2) solved as

$$r(\xi) = e^{-\xi I} r(0), \quad 0 < \xi \leq \frac{1}{\phi}.$$

Hence,

$$\begin{aligned} \mathcal{L}^{-1} \left[(sI - D^T A D^{T\dagger})^{-1} D^T (A + I) D \tilde{r} \right] &= \mathcal{L}^{-1} \left[(sI - D^T A D^{T\dagger})^{-1} \right] * \mathcal{L}^{-1} \left[D^T (A + I) D \tilde{r} \right] \\ &= e^{\xi D^T A D^{T\dagger}} * D^T (A + I) D e^{-\xi I} r(0) \\ &= e^{\xi D^T A D^{T\dagger}} * (D^T A D e^{-\xi I} r(0) + D^T D e^{-\xi I} r(0)) \\ &= e^{\xi D^T A D^{T\dagger}} * D^T A D e^{-\xi I} r(0) + e^{\xi D^T A D^{T\dagger}} * D^T D e^{-\xi I} r(0). \end{aligned}$$

The two convolutions are nearly identical so we solve the simpler of the two:

$$e^{\xi D^T A D^{T\dagger}} * D^T D e^{-\xi I} r(0) = \int_{\tau=0}^{\xi} e^{(\xi-\tau) D^T A D^{T\dagger}} D^T D e^{-\tau I} r(0) d\tau.$$

Note that $e^{-\tau I}$ simplifies as

$$\begin{aligned} e^{-\tau I} &= \sum_{k=0}^{\infty} \frac{(-\tau I)^k}{k!} \\ &= \left(\sum_{k=0}^{\infty} \frac{(-\tau^k)}{k!} \right) I \\ &= e^{-\tau I}. \end{aligned}$$

The scalar and identity matrix can both move to the beginning of the integral and reformed into a matrix:

$$\begin{aligned}
\int_{\tau=0}^{\xi} e^{(\xi-\tau)D^T A D^{T\dagger}} D^T D e^{-\tau I} r(0) d\tau &= \int_{\tau=0}^{\xi} e^{-\tau I} e^{(\xi-\tau)D^T A D^{T\dagger}} D^T D r(0) d\tau \\
&= e^{\xi D^T A D^{T\dagger}} \int_{\tau=0}^{\xi} e^{-\tau (I + D^T A D^{T\dagger})} d\tau D^T D r(0) \\
&= e^{\xi D^T A D^{T\dagger}} (I + D^T A D^{T\dagger})^{-1} \left(e^{\xi(I + D^T A D^{T\dagger})} - I \right) D^T D r(0).
\end{aligned}$$

From this expression it follows that

$$e^{\xi D^T A D^{T\dagger}} * D^T (A + I) D e^{-\xi I} r(0) = e^{\xi D^T A D^{T\dagger}} (I + D^T A D^{T\dagger})^{-1} \left(e^{\xi(I + D^T A D^{T\dagger})} - I \right) D^T (A + I) D r(0).$$

Hence,

$$\mathcal{L}^{-1} \left[(sI - D^T A D^{T\dagger})^{-1} D^T (A + I) D \tilde{r} \right] = e^{\xi D^T A D^{T\dagger}} (I + D^T A D^{T\dagger})^{-1} \left(e^{\xi(I + D^T A D^{T\dagger})} - I \right) D^T (A + I) D r(0). \quad (5.6)$$

Using equations (5.5) and (5.6), the voltage trajectory equation (5.4) becomes

$$\begin{aligned}
V(\xi) &= \\
&e^{\xi D^T A D^{T\dagger}} V(0) \\
&+ (D^T A D^{T\dagger})^{-1} \left(e^{\xi D^T A D^{T\dagger}} - I \right) d \\
&+ e^{\xi D^T A D^{T\dagger}} (I + D^T A D^{T\dagger})^{-1} \left(e^{\xi(I + D^T A D^{T\dagger})} - I \right) D^T (A + I) D r(0).
\end{aligned} \quad (5.7)$$

In the case $A = -I$, equation (5.7) simplifies considerably:

$$V(\xi) = e^{-\xi D^T D^{T\dagger}} V(0) + (D^T D^{T\dagger})^{-1} \left(I - e^{-\xi D^T D^{T\dagger}} \right) d. \quad (5.8)$$

Simplify the matrix $D^T D^{T\dagger}$ via its SVD:

$$\begin{aligned}
D^T &= V \begin{bmatrix} S \\ 0 \end{bmatrix} \mathcal{U}^T, \\
\implies D^{T\dagger} &= \mathcal{U} \begin{bmatrix} S & 0 \end{bmatrix} V^T \\
\implies D^T D^{T\dagger} &= V \begin{bmatrix} S \\ 0 \end{bmatrix} \mathcal{U}^T \mathcal{U} \begin{bmatrix} S & 0 \end{bmatrix} V^T \\
&= V \begin{bmatrix} S \\ 0 \end{bmatrix} \begin{bmatrix} S & 0 \end{bmatrix} V^T \\
&= V \begin{bmatrix} I_d & 0 \\ 0 & 0 \end{bmatrix} V^T \\
&= \begin{bmatrix} I_d & 0 \\ 0 & 0 \end{bmatrix} \in \mathbf{R}^{N \times N},
\end{aligned}$$

where I_d denotes the d -dimensional identity matrix. Equation (5.8) becomes

$$V(\xi) = e^{-\xi I_d} V(0) + \begin{bmatrix} I_d & 0 \\ 0 & 0 \end{bmatrix}^{-1} (I - e^{-\xi I_d}) d.$$

The matrix $\begin{bmatrix} I_d & 0 \\ 0 & 0 \end{bmatrix}$ is not invertible. Consider instead only the first d equations of the preceding system.

$$V_j(\xi) = e^{-\xi} V_j(0) + (1 - e^{-\xi}) d_j, \quad j = 1, \dots, d.$$

Note that $d_j = d^T c$, and consider neuron $j_{max} = j$ the first to reach the spike threshold. Recall its initial condition $v(0) = -\frac{\|d\|^2}{2}$ to arrive at

$$V(\xi) = -e^{-\xi} \frac{\|d\|^2}{2} + (1 - e^{-\xi}) d^T c$$

Compare with the corresponding PCF trajectory, equation (5.1). The preceding equation rearranges to

$$V(\xi) = d^T c - e^{-\xi} \left(d^T c + \frac{\|d\|^2}{2} \right),$$

which is identical to equation (5.1). Since only one neuron spikes, $r(\xi)$ and thus $\hat{x}(\xi)$ are identical for both PCF and gap-junction networks. The preceding, somewhat painful analysis shows that if the PCF and gap-junction models begin on their steady-state trajectories with the same initial conditions, their network estimates are identical in time. The statement is limited to the case of a constant driving stimulus. It does not, for example, show which network reaches the steady state trajectory first.

5.3 Per-spike RMSE of the PCF and Gap-Junction Networks for a Constant Stimulus

Equation (5.3) gives both gap-junction and PCF trajectories. We compute the per-spike RMSE by the integral

$$RMSE_{spike} \triangleq \sqrt{\phi \int_0^{\frac{1}{\phi}} e^T e(\tau) d\tau}.$$

The target dynamical system over this interval is $x(\xi) = \mathcal{U}_1$. Assuming the first spike is at $\xi_1 = 0$, we have

$$\begin{aligned}
e(\xi) &= x(\xi) - \hat{x}(\xi) \\
&= \mathcal{U}_1 - \frac{d}{1 - e^{-\frac{1}{\phi}}} e^{-\xi} \\
\Rightarrow e^T e &= \mathcal{U}_1^T \mathcal{U}_1 - 2\mathcal{U}_1^T \frac{d}{1 - e^{-\frac{1}{\phi}}} e^{-\xi} + \frac{d^T d}{\left(1 - e^{-\frac{1}{\phi}}\right)^2} e^{-2\xi} \\
&= 1 - 2 \frac{c^T d}{1 - e^{-\frac{1}{\phi}}} e^{-\xi} + \frac{d^T d}{\left(1 - e^{-\frac{1}{\phi}}\right)^2} e^{-2\xi}.
\end{aligned}$$

Integrate over a spike interval to arrive at

$$\begin{aligned}
\int_0^{\frac{1}{\phi}} e^T e(\tau) d\tau &= \frac{1}{\phi} - 2 \frac{c^T d}{1 - e^{-\frac{1}{\phi}}} \left(1 - e^{-\frac{1}{\phi}}\right) + \frac{d^T d}{\left(1 - e^{-\frac{1}{\phi}}\right)^2} \frac{1}{2} \left(1 - e^{-\frac{2}{\phi}}\right) \\
&= \frac{1}{\phi} - 2 c^T d + \frac{\|d\|}{2} \frac{1 - e^{-\frac{2}{\phi}}}{\left(1 - e^{-\frac{1}{\phi}}\right)^2}.
\end{aligned}$$

The per-spike RMSE of both PCF and Gap-Junction Networks is therefore

$$RMSE_{spike} = \sqrt{1 - 2\phi c^T d + \phi \frac{\|d\|^2}{2} \frac{1 - e^{-\frac{2}{\phi}}}{\left(1 - e^{-\frac{1}{\phi}}\right)^2}}.$$

To write the RMSE as a function of only firing rate ϕ , we invert equation (5.2) to obtain $d(\phi)$:

$$\frac{1}{\phi} = \ln \left(d^T c + \frac{\|d\|^2}{2} \right) - \ln \left(d^T c - \frac{\|d\|^2}{2} \right).$$

Note that $d^T c = d^T \begin{bmatrix} 1 \\ 0 \end{bmatrix} = d_0$. Because d is the most parallel vector to c , for large enough networks with uniformly distributed directions d , we have that $d = d_0 c$. This implies that $\|d\|^2 = d_0^2$. The preceding equation becomes

$$\begin{aligned}
\frac{1}{\phi} &= \ln \left(d_0 + \frac{d_0^2}{2} \right) - \ln \left(d_0 - \frac{d_0^2}{2} \right) \\
\Rightarrow e^{-\frac{1}{\phi}} &= \frac{d_0 - \frac{d_0^2}{2}}{d_0 + \frac{d_0^2}{2}} \\
&= \frac{1 - \frac{d_0}{2}}{1 + \frac{d_0}{2}} \\
\Rightarrow e^{-\frac{1}{\phi}} + e^{-\frac{1}{\phi}} \frac{d_0}{2} &= 1 - \frac{d_0}{2} \\
\Rightarrow d_0 \frac{1 + e^{-\frac{1}{\phi}}}{2} &= 1 - e^{-\frac{1}{\phi}} \\
\Rightarrow d_0(\phi) &= 2 \frac{1 - e^{-\frac{1}{\phi}}}{1 + e^{-\frac{1}{\phi}}} \\
&= 2 \tanh \frac{1}{2\phi}
\end{aligned}$$

Thus the per-spike RMSE simplifies to

$$RMSE_{spike} = \sqrt{1 - 4\phi \tanh \frac{1}{2\phi} + 2\phi \tanh^2 \frac{1}{2\phi} \frac{1 - e^{-\frac{2}{\phi}}}{\left(1 - e^{-\frac{1}{\phi}}\right)^2}}. \quad (5.9)$$

5.4 Comparison of Self-Coupled, Gap-Junction, and PCF Networks for a Constant Stimulus

We now compare all three models as they respond to a constant driving stimulus, while varying their firing rate.

Let the parameters be

$$A = - \begin{bmatrix} 1 & 0 \\ 0 & 1 \end{bmatrix},$$

$$B = \begin{bmatrix} 1 \\ 0 \end{bmatrix},$$

$$c(\xi) = \begin{bmatrix} 1 \\ 0 \end{bmatrix},$$

$$D = \mathcal{U} \begin{bmatrix} S & 0 \end{bmatrix} V^T = \mathcal{U} \begin{bmatrix} I_d & 0 \end{bmatrix} I_N,$$

$$d\xi = 10^{-4},$$

$$N = 8,$$

$$x(0) = \begin{bmatrix} \frac{1}{2} & 0 \end{bmatrix}.$$

We simulate the self-coupled, PCF, and gap-junction networks and compare their derived estimates given by equations (??) and (5.3) respectively. Figure (12) shows the network estimates of each model for the above parameters. We see that all trajectories are identical as predicted by our derived estimates, shown by the dotted line.

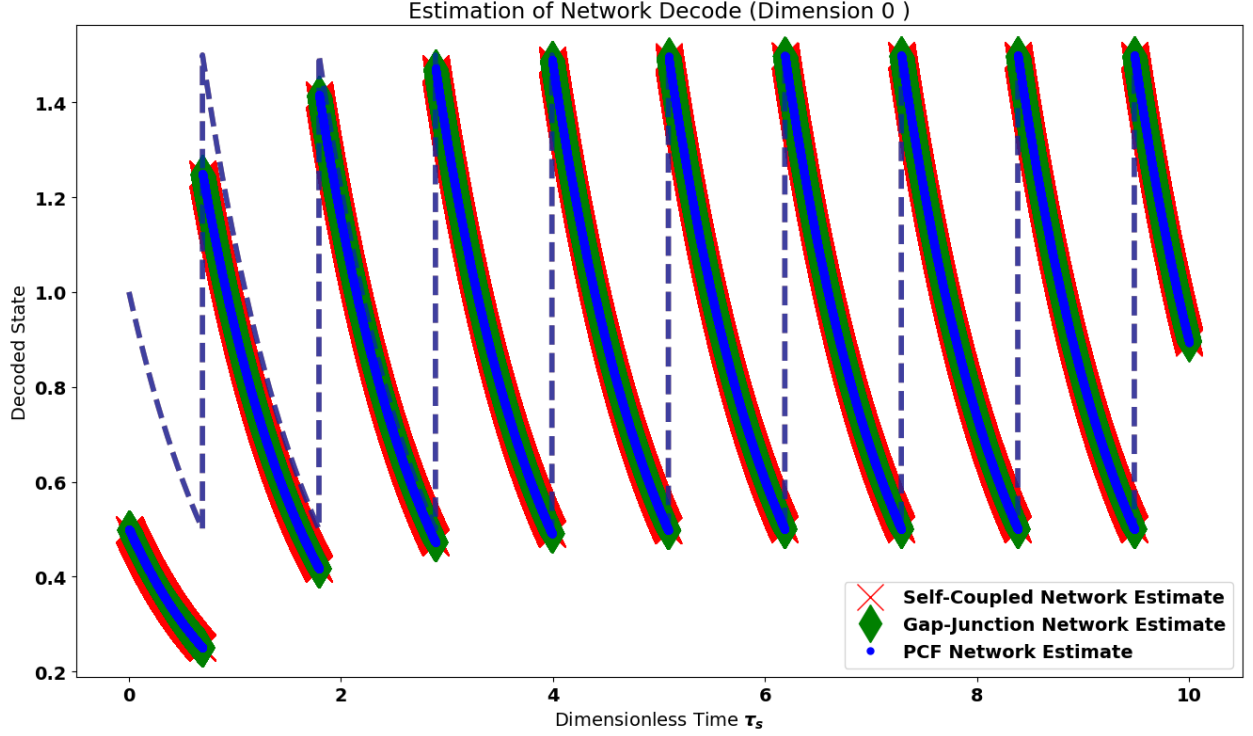


Figure 12: Simulation of self-coupled, gap-junction, and PCF networks. Network readouts for each are plotted. The dotted line is the derived expression(s) given by equations (??) and (5.3). Note that all three trajectories are identical.

Next we plot the per-spike RMSE of each model for the same parameters while varying the spike rate. Figure (13) shows the numerically measured per-spike RMSE for each model and their derived expressions, equations (??) and (5.9). As the firing rate approaches the simulation timestep, 10^4 , the curves deviate from the derived expression.

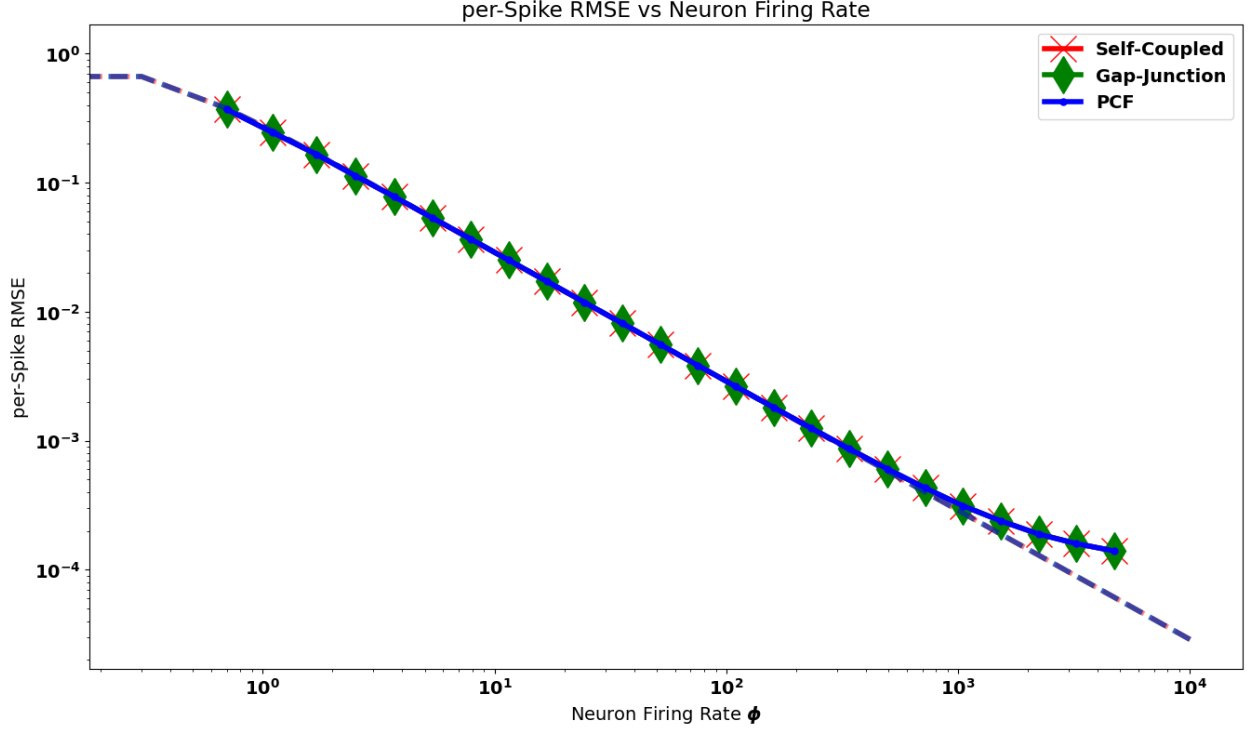


Figure 13: Simulated per-spike RMSE for self-coupled, gap-junction, and PCF networks. The dotted lines are the derived expression for each model given by given by equations (??) for the self-coupled and (5.9) for the gap-junction and PCF models respectively. Spike rates were estimated numerically by dividing the number of spikes by the simulation length. The RMSE was computed numerically by the discrete integral $RMSE = \sqrt{\hat{\phi} \sum_{\tau \text{ between spikes}} e(\xi)^T e(\xi) d\xi}$. All computations used the numerically estimated spike rate.