Criticality in neural networks paper title

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Abstract

It has been posited that biological neural networks, such as a brain, may naturally exist in critical states. We propose two mechanisms for signal transduction in two such networks as encoding strategies which are optimized by criticality. First, we examine compressive sensing in a 2-dimensional Ising model at or near its critical temperature. Secondly, we examine the dynamical synchronization capabilities of a random neural network model as it transitions into chaotic behavior. We propose that both techniques should be most successful at the critical state of either model.

1 Introduction

- 1. Biological neural networks have been posited to exist at or near critical states.
- A critical state is characterized by the divergence of statistical correlation lengths in the system.
- 3. We propose that criticality may play a role
- 4. We also examined a dynamical model which exhibits critical behavior near the transition to chaotic dynamics. The model incorporates a capacitive leak current for each element, as well as a coupling to "neighboring" elements through a saturating, nonlinear I/V relationship.
- 5. The ability of the model to dynamically reconstruct an input signal from measurements of a limited subset of network elements was tested near the transition to chaotic behavior and compared to the case away from the critical point.

2 Dynamical synchronization technique

The model equations are coupled to a set of data through a linear term as

$$\frac{dx_i}{dt} = f_i(x) + \sum_j g_{ij} (y_j - x_j) \tag{1}$$

where x(t) and y(t) are the model and the data trajectories, respectively, and g_{ij} is a set of positive coupling constants. In practice, only a subset of the couplings will be nonzero (corresponding to the components of the system which can be measured) and the matrix g_{ij} is diagonal, since in this scheme there is no obvious advantage to coupling different components of the system to each other¹.

¹In a further examination of the model, we hope to test a coupling scheme which couples components to each other in time-delayed coordinates. Because the components are coupled through the dynamics, the time-delayed trajectories can provide information about each other and enhance the scheme by coupling unmeasurable components to the data.

It may be possible for *all* the components of a model to synchronize to a data set if enough measurements are made, and if the coupling is large enough. What constitutes "enough" seems to come on a case-by-case basis. Roughly, what is required is for the linear terms to regularize the solution manifold by making the value of the largest Lyapunov exponent in the coupled system non-positive. However, the ratio of the *i*-th coupling term to the *i*-th term of the dynamics must remain small, so as to not wash out the physicality of the problem (in practice, this may also increase the speed of convergence to the actual solution). In other words, adjusting the coupling is a delicate balancing act for which there is no general procedure.

The success of the procedure, however, is not ultimately measured by the ability to synchronize, but by the ability of the model *make predictions* after the coupling is turned off, since in general we're only able to measure a few components of the system. In the case where we have access to all of the data states, however, we can additionally test the procedure by examining the normally unobservable states. This is called a *twin experiment* in the literature and is a useful technique for assessing the viability of the model to synchronize in a controlled setting.

3 Random neural network model

We now consider a neural network model, originally presented by [1], which is a dynamical system describing a network of N randomly connected resitive and capacitive elements. The equations of motion describing the system are

$$\frac{dV_i}{dt} = -V_i + \sum_{j=1}^{N} J_{ij}\phi(V_j)$$
(2)

where $\phi(V_i)$ is an I/V relationship which may be thought of as an "activity" proportional to the synaptic current between neurons i and j. One possible choice is a sigmoid function of V, i.e. $\phi_i = \tanh{(\alpha V_i)}$. This choice is both biologically motivated as acting to saturate synaptic activity as a function of membrane voltage, as well as mathematically to avoid highly unstable, runaway solutions to eq. (2). α acts as a control parameter on the turnaround rate of the synaptic activity around V=0; in some sense it controls the "degree of nonlinearity" in the system.

The matrix J_{ij} describes the connectivity in the network; in this particular model, J_{ij} is chosen to be a Gaussian random matrix with elements distributed according to the statistics

$$\langle J_{ij} \rangle = 0, \quad \langle J_{ij} J_{kl} \rangle = \frac{\tilde{J}^2}{N} \, \delta_{ik} \delta_{jl}$$
 (3)

with zero on the diagonals to eliminate self-coupling terms. This means synaptic connections are totally decorrelated and, in general, asymmetric. This also means that, on average, half of the connections are inherently inhibitory and half are excitatory. If J_{ij} were symmetric, (2) would describe a system that relaxes to a global minimum of an energy function; with asymmetric couplings the dynamics are that of a spin glass, which in general have nonrelaxational and possibly chaotic behavior.

A detailed mathematical treatment of this model in the limit of large N is given in [1]. The result is that under the replacement $\alpha \to \tilde{J}$ in the expression for ϕ , the model undergoes a "phase transition" when the control parameter \tilde{J} reaches a critical value of 1. This is manifest in the structure of the attracting solution manifold in the $\{V_i\}$ state space, which acquires at least one positive Lyapunov exponent; in other words a family of chaotic solutions to (2) appears.

The similarity of this transition to the phase transition in the 2D Ising model near T_{crit} is most apparent in the behavior of the quantity

$$\Delta(t) \equiv \langle V_i(t_0)V_i(t_0+t)\rangle \tag{4}$$

which is the autocorrelation of a single voltage as a function of time.

3.1 Numerical analysis

We examined the model for N=256 neurons with a single instantiation of J_{ij} . Ideally, we would like to scale up the simulation size to a larger $N\sim 1000$, and to gather statistics about an *ensemble* of

models parameterized by different instantiations of J_{ij} . However, limited by time, we now present said results as at least a preliminary examination of the model.

First, we performed a comparison of numerical results to the analytic results of [1]. For a single randomly drawn instantiation of J_{ij} , (2) was integrated forward in time using the LSODA solver in scipy.integrate. The initial conditions were drawn randomly from a ball of radius 1 centered about the origin $V_i = 0$. We found that near $\tilde{J} = 1$, limit cycle solutions appeared in a Hopf bifurcation from an attracting fixed point at the origin (which became unstable in the bifurcation).

To examine the behavior of $\Delta(t)$, it was assumed that the steady state of the system was ergodic so as to allow the replacement

$$\langle V_i(t_0)V_i(t+t_0)\rangle \to \int_{t_0}^{T-t} dt' \ V_i(t')V_i(t+t')$$

(up to an overall scale factor) where $T - t_0$ is the length of the recorded time series. In other words, the ensemble average was replaced with a time average. This was necessary because only a single instantiation of J_{ij} was used, and the LHS is an ensemble average over the distribution of J_{ij} .

Finding the critical value of \tilde{J} provides us with a place to examine the ability of the model to synchronize. The observer system looks like

$$\frac{dV_i}{dt} = -V_i + \sum_{j=1}^{N} J_{ij} \tanh(\tilde{J}V_j) + g_i(V_i^{(D)} - V_i)$$
(5)

where M elements of g_i are nonzero, corresponding to the M measured components of the system (corresponding to the M data time series $V_i^{(D)}$).

References

[1] H Sompolinsky, A Crisanti, and H. Sommers. Chaos in random neural networks. *Physical Review Letters*, 61(3):259–262, July 1988.