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Guiding Synchrony through Random Networks

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(Received 9 August 2011; revised manuscript received 18 June 2012; published 13 December 2012)

Sparse random networks contain structures that can be considered as dilated feed-forward networks. Modeling of cortical circuits has shown that feed-forward structures, if strongly pronounced compared to the embedding random network, enable reliable signal transmission by propagating localized (subnetwork) synchrony. This assumed prominence, however, is not experimentally observed in local cortical circuits. Here, we show that nonlinear dendritic interactions, as discovered in recent single-neuron experiments, naturally enable guided synchrony propagation already in random recurrent neural networks that exhibit mildly enhanced, biologically plausible substructures.

DOI: 10.1103/PhysRevX.2.041016

Subject Areas: Biological Physics, Complex Systems

I. INTRODUCTION

Cortical neural networks generate a ground state of highly irregular spiking activity whose dynamics is sensitive to small perturbations such as missing or additional spikes [1–4]. A robust, reliable transmission of information in the presence of such perturbations and noise is nonetheless assumed to be essential for neural computation. It has been hypothesized that this reliable transmission might be achieved by propagation of pulses of synchronous spikes along feed-forward chains [5]. In current models, functionally relevant chains require a dense connectivity between the neuronal layers of the network [6] or strongly enhanced synapses and specifically modified response properties of neurons within the chain [7]. Such highly distinguished large-scale structures are not observed experimentally, however.

Can less-structured networks also guide synchrony? Recently, single-neuron experiments have revealed a mechanism that nonlinearly promotes synchronous inputs. On synchronous dendritic stimulation, neurons are capable of generating fast dendritic spikes. In the soma, these spikes induce rapid, strong depolarizations [8] that are nonlinearly enhanced compared to depolarizations expected from linear summation of single inputs. If the dendritic spike induces an action potential in the soma, the potential occurs at a fixed time after the stimulation, with submillisecond precision. Other experiments have found slow dendritic spikes that are comparably insensitive to input synchrony [9]. These slow dendritic spikes endow single neurons with computational capabilities comparable to multilayered feed-forward networks of simple-rate neurons [10]. Furthermore, they provide a possible mechanism underlying neural bursting and its propagation, which have been shown to enhance reliability and temporal precision of signal propagation [11,12]. The impact on collective circuit dynamics of fast dendritic spikes that induce nonadditive coupling has not been systematically investigated in a general setting so far.

In this article, we show that and describe how fast dendritic nonlinearities may support guided-synchrony propagation in neural circuits. First, we develop an analytical approach to describe such propagation in linearly and nonlinearly coupled networks. In particular, we derive expressions for the critical connectivity above which propagation occurs and for the size of the propagating pulse. We quantify how dendritic nonlinearities compensate for dense anatomical connections and thereby promote propagation of synchrony. Finally, using large-scale simulations of more detailed recurrent network models, we show that feed-forward networks that occur naturally as part of random circuits enable persistent guided synchrony propagation due to dendritic nonlinearities.

II. MODELS AND METHODS

A. Analytically tractable model

Model with linear summation of inputs. As a basis model, we consider networks of conventional leaky integrate-and-fire neurons that interact by sending and receiving spikes via directed connections. The membrane potential \( V_l(t) \) of a neuron \( l \) satisfies

\[
\dot{V}_l(t) = -\gamma_l V_l(t) + I_l(t),
\]

where \( \gamma_l \) is the inverse membrane time constant and \( I_l(t) \) is the total input current at time \( t \). In addition to inputs from the network, the neurons receive excitatory and inhibitory random inputs that emulate an embedding network, i.e.,
The subthreshold dynamics of the membrane potential converges to its asymptotic value $V^{\text{rest}}_l := V_l^0 / \gamma_l$. When the neuron’s membrane potential reaches or exceeds its threshold $\Theta_l$, its membrane potential is reset to $V^{\text{rest}}_l$ and a spike is emitted, which arrives at the postsynaptic neuron $j$ after a delay time $\tau_{jl}$. For a refractory period $t^R_{jl}$ after the reset, all incoming spikes to neuron $l$ are ignored, and the membrane potential is kept at $V^{\text{rest}}_l$.

We model the fast rise of the membrane potential on the arrival of a presynaptic spike by an instantaneous jump, such that the contributions of the arriving external spikes to the total input current are given by

$$I^\text{ext,ex}_l(t) = \sum_{k \in K} \epsilon^{\text{ext,ex}}_{lk} \delta(t - t^\text{ext,ex}_{lk}),$$

$$I^\text{ext,ln}_l(t) = \sum_{k \in K} \epsilon^{\text{ext,ln}}_{lk} \delta(t - t^\text{ext,ln}_{lk}),$$

where $t^\text{ext,ex}_{lk}$ ($t^\text{ext,ln}_{lk}$) are the arrival times of the $k$th excitatory (inhibitory) external spike at neuron $l$, $\epsilon^{\text{ext,ex}} > 0$ or $\epsilon^{\text{ext,ln}} < 0$ are the strengths of single external spikes, and $\delta(\cdot)$ is the Dirac $\delta$ distribution. Analogously, the contribution of spikes received from neurons of the network is given by

$$I^\text{net}_l(t) = \sum_j \sum_k \epsilon_{lj} \delta(t - t^j_{jk} - \tau_{lj}),$$

where $\epsilon_{lj}$ is the coupling strength from neuron $j$ to $l$ and $t^j_{jk}$ is the $k$th spike time of neuron $j$.

**Model with nonlinear summation of inputs.** In the above model without nonlinear dendrites, the strengths of synchronous inputs are summed up linearly [cf. Eq. (5)]. We incorporate nonlinear dendrites by modulating this sum for excitatory inputs by a nonlinear function $\sigma$ that can be directly read off from experimental results [8]: $\sigma$ equals the identity for small excitatory input, increases steeply when the input exceeds a threshold $\Theta_b$, and saturates for larger inputs. We define the dendritic modulation function as

$$\sigma(\epsilon) = \begin{cases} 
\epsilon & \text{for } \epsilon \leq \Theta_b \\
_k & \text{otherwise}
\end{cases}$$

For simplicity, we consider only exactly simultaneous spikes as synchronous. Accordingly, conduction delays are chosen homogeneously, $\tau_{lj} = \tau$, so that synchronous presynaptic spiking can be amplified. In this scenario, the detection of synchronous events is straightforward. However, systems with heterogeneous delays and a finite dendritic integration window exhibit qualitatively the same phenomena [13]. The contribution of spikes received from the network is then given by

$$I^\text{net}_l(t) = \sum_j \left( \sigma\left( \sum_{k \in M_{\text{ex}}(t')} \epsilon_{lj} \right) + \sum_{k \in M_{\text{in}}(t')} \epsilon_{lj} \right) \delta(t - t' - \tau),$$

where $t'$ are all firing times in the network. The sets $M_{\text{ex}}(t')$ and $M_{\text{in}}(t')$ denote the sets of indices of neurons sending an excitatory or inhibitory spike at time $t'$, respectively. Networks with linear dendrites can be described by setting $\sigma(\epsilon) = \epsilon$.

**B. Biologically more detailed model**

**Conductance-based model.** In the last part of this article, we employ a biologically more detailed neuron model to highlight the generality of our findings on propagation enhancement. The neuron model is a conductance-based, leaky integrate-and-fire neuron that is augmented by terms introducing the impact of dendritic spikes (see also [14]). The subthreshold dynamics of the membrane potential $V_l$ of neuron $l$ obeys the differential equation

$$C^m_l \frac{dV_l(t)}{dt} = g^I_l [V_l^{\text{rest}} - V_l(t)] + g^E_l [E^{\text{ex}} - V_l(t)]$$

$$+ g^G_l [E^{\text{in}} - V_l(t)] + I^\text{DS}_l(t) + I^0_l.$$ (8)

Here, $C^m_l$ is the membrane capacity, $g^I_l$ is the resting conductance, $V^{\text{rest}}_l$ is the resting membrane potential, $E^{\text{ex}}$ and $E^{\text{in}}$ are the reversal potentials, and $g^E_l(t)$ and $g^I_l(t)$ are the conductances of excitatory and inhibitory synaptic populations, respectively. $I^\text{DS}_l(t)$ models the current pulses caused by dendritic spikes, and $I^0_l$ is a constant current gathering slow external and internal currents. The time course of single synaptic conductances contributing to $g^I_l(t)$ and $g^E_l(t)$ is given by the difference between two exponential functions (e.g., [15]). Whenever the membrane potential reaches the spike threshold $\Theta_l$, the neuron sends a spike to its postsynaptic neurons, is reset to $V^{\text{rest}}_l$, and becomes refractory for a period $t^R_{jl}$.

To account for dendritic spike generation, we consider the sum $g_{l,t\Delta t}$ of excitatory input strengths (characterized by the coupling strengths) arriving at an excitatory neuron $l$ within the time window $\Delta t$ for nonlinear dendritic interactions,

$$g_{l,t\Delta t}(t) = \sum_j \sum_k g_{lj}^{\text{max}} \chi_{[t, t - \Delta t]}(t^j_{jk} + \tau),$$

where $\chi_{[t, t - \Delta t]}$ is the characteristic function of the interval $[t, t - \Delta t]$, $t^j_{jk}$ is the $k$th firing time of excitatory neuron $j$, and $\tau$ denotes the synaptic delay. We denote the peak conductance (coupling strength) for a connection from
neuron $j$ to neuron $l$ by $g_{jl}^{\max}$. If $g_{j;\Delta t}$ exceeds a threshold $g_{\theta}$, a dendritic spike is initiated and the dendrite becomes refractory for a time window $\tau_{\text{DS},\text{ref}}$. The effect of the dendritic spike is incorporated into the model by the current pulse that reaches the soma at a time $\tau_{\text{DS}}$ thereafter. This current pulse is modeled as the sum of three exponential functions,

$$I_{i;\text{S}}(t) = c(g_{j;\Delta t})[-A e^{-t/\tau_{\text{DS},1}} + B e^{-t/\tau_{\text{DS},2}} - Ce^{-t/\tau_{\text{DS},3}}],$$

(10)

with prefactors $A > 0$, $B > 0$, $C > 0$, decay time constants $\tau_{\text{DS},1}$, $\tau_{\text{DS},2}$, $\tau_{\text{DS},3}$, and a dimensionless correction factor $c(g_{j;\Delta t})$, where $g_{j;\Delta t}$ is the summed excitatory input at the initiation time of the dendritic spike as given by Eq. (9). The factor $c(g_{j;\Delta t})$ modulates the pulse strength, ensuring that the peak of the excitatory postsynaptic potential (pEPSP) reaches the experimentally observed region of saturation. At very high excitatory inputs, the conventionally generated depolarization exceeds the level of saturation, and the pEPSP increases [cf. Fig. 1(a)].

Detection probability. In the last part of this article, we investigate recurrent networks where a feed-forward subnetwork consisting of a certain number of layers (groups) is created by modifying strengths of existing synaptic connections of the network. To decide whether propagation of synchrony in recurrent networks is successful, we consider the signal-to-noise ratio (SNR): We pick $\omega$ neurons, randomly selected from the network, to be a first group. After initiation of synchronous activity in this group, we count the number of spikes from neurons of the $i$th group, $S_i$, within a time window $[t_{i;\exp} - \tau_{i/2}, t_{i;\exp} + \tau_{i/2}]$. (For details on how the $i$th group is defined, see Sec. III C on recurrent networks.) Here, $t_{i;\exp}$ is the expected time for the synchronous pulse to reach layer $i$, and $\tau_{i}$ is the expected width of the synchronous pulse. We consider all spikes within the time window of size $\tau_{i}$ centered at $t_{i;\exp}$ as part of the synchronous pulse. We assume that $I_{i;\exp} = I_{i;\exp} + (i - 1)\Delta t_{\exp}$, where $\Delta t_{\exp}$ is chosen after simulation such that $\sum S_i$ becomes maximal:

$$S_i = \sum_{j \in \text{Gr}(i)} \sum_{k} \chi_{[t_{i;\exp} - (\frac{\tau_{i}}{2}), t_{i;\exp} + (\frac{\tau_{i}}{2})]}(t_{j;\exp})$$

(11)

Here, Gr($i$) are the indices of neurons of group $i$, $t_{j;\exp}$ is the $k$th firing time of neuron $j$, and $\chi$ denotes the characteristic function, as before.

To determine the noise level of group $i$, we measure the probability $P_{i;\Delta t_{\exp}, \tau_{i}}(k)$ of finding $k$ spikes from neurons of group $i$ within time windows $\tau_{i}$ over a control time interval during which no synchronous activity is initiated. The noise level $N_i$ of group $i$ is the minimal value satisfying

$$\sum_{k=0}^{N_i} P_{i;\Delta t_{\exp}, \tau_{i}}(k) \leq a,$$

(12)

with a constant $a \approx 1$.

Finally, we denote the propagation of synchrony up to the $i$th layer as successful if the SNR is larger than $b$,

$$\text{SNR}_i := \min_{j=1,...,\omega} \left( \frac{S_j}{N_j} \right) > b,$$

(13)

where $b \geq 1$. This means, in particular, that we can distinguish the background (spontaneous) activity from the
III. RESULTS

A. Feed-forward chains with linear coupling

How can diluted feed-forward networks (FFNs) propagate synchrony? FFNs consist of a sequence of layers, each composed of $\omega$ excitatory neurons; they forward connections to neurons in the subsequent layer randomly present with probability $p$. Present connections have strength $e$. Synchronous spiking activity is initiated by exciting neurons of the first layer to spike simultaneously. In the second layer, the synchronous pulse arriving from the first layer excites a certain subgroup of neurons to spike simultaneously which in turn generates a synchronous input to layer three, etc.

To understand the collective dynamics analytically, we consider networks of leaky integrate-and-fire neurons in the limit of fast synaptic currents (cf. Sec. II). In the absence of synchronous activity, each neuron of the FFN receives a large number of inputs from an emulated external network and only very few inputs from the previous layer, such that its dynamics is practically identical to the ground state of balanced networks. If the connections within the FFN are weak and/or the connection probability is low, the spontaneous spiking activity is influenced only weakly by spiking activity of the FFN. Therefore, we assume that the ground-state activity is exclusively governed by the external inputs, effectively setting couplings within the chain to $\epsilon_{ij} = 0$. The external input is balanced, i.e., the mean input is subthreshold, and spontaneous spiking is caused by fluctuations in the input. The network’s neurons thus spike in an asynchronous and irregular manner [1,2] and the stationary distribution of membrane potentials $P_V(V)$ can be calculated analytically in diffusion approximation [2,16].

$$p_f(x) := \int_{\Theta - x}^{\Theta} P_V(V) dV$$

(14)

is the probability of finding a neuron’s membrane potential in the interval $[\Theta - x, \Theta]$. We model the fast rise of the membrane potential on the arrival of (possibly nonlinear enhanced) presynaptic spikes by an instantaneous jump in the membrane potential (cf. Sec. II); thus, $p_f[\sigma(h\epsilon)]$ specifies the spiking probability of a single neuron, after receiving $h$ input spikes of strength $\epsilon$ from the preceding layer.

To assess the propagation of synchrony, we consider the average number of neurons that are activated in each layer in response to the initial synchronous pulse (cf. also [17]). When $g_i$ neurons spike synchronously in layer $i$,

$$p^p(g_i) := \sum_{k=0}^{g_i} \binom{g_i}{k} h^k (1 - p)^{g_i - k} p^p[\sigma(h\epsilon)]$$

(15)

is the probability of spiking of a particular neuron in layer $i + 1$, where the number of simultaneous inputs $h$ is binomially distributed, $h \sim B(g_i, p)$. Thus, for layers of size $\omega$, the average number of neurons spiking in layer $i + 1$ is

$$\langle g_{i+1} \rangle = \omega p^p(g_i).$$

(16)

Substituting the average group size $\langle g_i \rangle$ for the actual size $g_i$ yields the interpolated map $\langle g_{i+1} \rangle = \omega p^p(\langle g_i \rangle)$, whose fixed points qualitatively determine the propagation of synchronous activity (cf. Fig. 2).

The trivial, absorbing fixed point $G_0 = 0$, defining a state of extinguished activity, always exists. For sufficiently small $p$, $\epsilon$, and $\omega$, this is the only fixed point. With increasing connectivity and layer size, a pair of fixed
points ($G_1$, unstable, and $G_2$, stable) appears via a tangent bifurcation. Initial pulses in the basin of $G_2$ (i.e., those larger than $G_1$) typically initiate stable propagation of synchrony with group sizes around $G_2$. For given layer size $\omega$ and connection strength $\epsilon$, the critical connectivity $p^*$ for which $G_1 = G_2$ marks the minimal connectivity that supports stable propagation of synchrony.

To elaborate the influence of nonlinear dendritic interactions, we derive the critical connectivity for FFNs. The mechanisms underlying propagation of synchrony are different for networks with and without nonlinear dendritic interactions and thus require different analytical approaches to derive $p^*$. We first consider feed-forward chains with conventional, linear coupling, i.e., $\sigma(x) = x$. To obtain $p^*$, we first expand $p_f(x)$ into a Taylor series up to first order around the mean of the binomial distribution specifying the average number $pg_i$ of active neurons in each layer, such that Eq. (16) simplifies to

$$
\langle g_{i+1} \rangle = \omega \sum_{h=0}^{g_i} \binom{g_i}{h} p^h (1-p)^{g_i-h} p_f(h\epsilon)
$$

(17)

$$
= \omega \sum_{h=0}^{g_i} \binom{g_i}{h} p^h (1-p)^{g_i-h}
$$

(18)

$$
\times [p_f(g_ip\epsilon) + p'_f(g_ip\epsilon)(h\epsilon - g_ip\epsilon)]
$$

(19)

$$
= \omega p_f(g_ip\epsilon).
$$

(20)

The linear approximation becomes exact in the limit of large layer sizes $\omega$ and small couplings $\epsilon$, where the product $\omega \epsilon$ is kept constant. We obtain an interpolated map from expression (20) by replacing $g_i$ by its mean value $\langle g_i \rangle$. At the fixed point $G := \langle g_{i+1} \rangle = \langle g_i \rangle$, the function

$$
F(G, p^+; \omega, \epsilon) := G - \omega p_f(p^+G\epsilon) = 0
$$

(21)

vanishes. Here, the values $G$ and $p^+$ are the average group size and the connection probability at the fixed point for given layer size $\omega$ and coupling strength $\epsilon$, respectively. Furthermore, $F$ has a double root at the bifurcation point, so the derivative

$$
\frac{\partial F(G, p^*, \omega, \epsilon)}{\partial G} = 1 - \omega p^* \epsilon p'_f(p^*G\epsilon) = 0
$$

(22)

also vanishes such that the derivative of $p_f$ at the bifurcation point is given by

$$
p'_f(p^*G\epsilon) = \frac{1}{\omega p^* \epsilon}.
$$

(23)

Combining the above equations, we express the derivatives of $F$ at the bifurcation point by

$$
\frac{\partial F(G, p^*, \omega, \epsilon)}{\partial p^*} = -\omega G \epsilon p'_f(p^*G\epsilon) = -\frac{G}{p^*},
$$

(24)

and

$$
\frac{\partial F(G, p^*, \omega, \epsilon)}{\partial \epsilon} = -\omega p^* \epsilon p'_f(p^*G\epsilon) = -\frac{G}{\epsilon}.
$$

(25)

Applying the implicit function theorem yields the set of derivatives of $p^*$,

$$
\frac{\partial p^*(G, \omega, \epsilon)}{\partial G} = -\left(\frac{\partial F(G, p^*, \omega, \epsilon)}{\partial p^*}\right)^{-1} \frac{\partial F(G, p^*, \omega, \epsilon)}{\partial G} = 0,
$$

(27)

and

$$
\frac{\partial p^*(G, \omega, \epsilon)}{\partial \omega} = -\left(\frac{\partial F(G, p^*, \omega, \epsilon)}{\partial p^*}\right)^{-1} \frac{\partial F(G, p^*, \omega, \epsilon)}{\partial \omega} = -\frac{p^*(G, \omega, \epsilon)}{\omega},
$$

(28)

and

$$
\frac{\partial p^*(G, \omega, \epsilon)}{\partial \epsilon} = -\left(\frac{\partial F(G, p^*, \omega, \epsilon)}{\partial p^*}\right)^{-1} \frac{\partial F(G, p^*, \omega, \epsilon)}{\partial \epsilon} = -\frac{p^*(G, \omega, \epsilon)}{\epsilon},
$$

(29)

which are solved by

$$
p^*_f := p^* = \frac{1}{\lambda \epsilon \omega},
$$

(30)

where $\lambda$ is a constant independent of $\omega$ and $\epsilon$. We note that we did not make explicit assumptions on the distribution of membrane potentials $P_i(V)$, which is determined by the setup of the external network, i.e., the external input current $I^0$, the coupling strengths $\epsilon^{\text{ext,ex}}$ and $\epsilon^{\text{ext,in}}$, as well as the firing rates $\nu^{\text{ext,ex}}$ and $\nu^{\text{ext,in}}$. With a different, lengthier approach based on a second-order expansion of $p_f$, one can derive an analytical estimate of $G$ [13]. Figure 3(a) displays this analytical approximation for $p^*_f$ and its agreement with numerical simulations. For connectivity larger than $p^*_f$, there is stable propagation of synchrony even in networks with linear dendritic interactions, and the size of the propagating pulse fluctuates around the stable fixed point $G_s$ of Eq. (16). (For very large connectivity, pathological high-frequency spiking activity can emerge due to spontaneous chain activation.)

**B. Feed-forward chains with nonlinear coupling**

We now consider networks incorporating nonlinear dendritic interactions and show that the connectivity and number of active neurons required for propagation of synchrony are smaller. In such networks, the mechanism underlying propagation of synchrony is different, because it is supported predominantly by nonlinearly enhanced inputs. As a consequence, the maximal input is bounded by $\kappa$, leading to a saturation in the return map (16)
[cf. Fig. 2(c)]. The saturation enables propagating pulses of a size substantially smaller than $\omega$, in contrast to linearly coupled networks. The discontinuity in the modulation function $\sigma$ induces a discontinuity in $p_f(\sigma(x))$, which prevents our previous analytical method. We thus determine the critical connectivity by a self-consistency approach. When a synchronous pulse arrives at a specific layer, the summed excitatory input strength $x$ is either smaller or larger than the dendritic threshold $\Theta_b$. For sufficiently small $\Theta_b$, the spiking probability of a neuron due to a subthreshold input is much smaller than that due to a suprathreshold input, i.e., $p_f(\Theta_b) \ll p_f(\kappa)$. Thus, only a small fraction of neurons receives an input smaller than $\Theta_b$ and is elicited to spike. We approximate $p_f(x) = 0$ for $x \leq \Theta_b$. When there is persistent propagation of synchrony, $p_\gamma$ (which denotes the fraction of neurons that receive sufficiently strong input to reach the dendritic threshold) is constant throughout the layers. The total spiking probability of a single neuron on the arrival of the synchronous pulse is then given by the product $p_\gamma p_f(\kappa)$. The probability

$$p^g(\delta) = \binom{\omega}{g} [p_\gamma p_f(\kappa)]^g [1-p_\gamma p_f(\kappa)]^{\omega-g}$$

for $g$ neurons to spike synchronously follows a binomial distribution. By combining the total spiking probability and the topological connection probability $p$, we compute the probability

$$P(k) = \sum_{g=k}^{\omega} \binom{\omega}{g} p^g(1-p)^{\omega-g}$$

$$= \binom{\omega}{k} [p_\gamma p_f(\kappa)]^k [1-p_\gamma p_f(\kappa)]^{\omega-k}$$

that a neuron of the subsequent layer receives exactly $k$ synchronous spikes. Thus, $k$ itself is binomially distributed, and we denote its mean value by $\delta$ and its standard deviation by $\sigma_\delta$. Using a Gaussian approximation of the binomial distribution yields the self-consistent equation

$$p_\gamma = \sum_{k=\lceil \Theta_b/\epsilon \rceil}^{\omega} P(k)$$

$$= \int_{\Theta_b/\epsilon}^{\omega} \frac{1}{\sqrt{2\pi}\sigma_\delta} \exp \left[ -\frac{1}{2} \left( \frac{k-\delta}{\sigma_\delta} \right)^2 \right] dk$$

$$= \frac{1}{2} \left[ 1 - \text{Erf} \left( \frac{\Theta_b - \delta}{\sqrt{2}\sigma_\delta} \right) \right]$$

$$= \frac{1}{2} \left[ 1 + \text{Erf} \left( \frac{n}{\sqrt{2}} \right) \right]$$

where we defined

$$n := \frac{\delta - \Theta_b/\epsilon}{\sigma_\delta}$$

as the distance between the average number of inputs and the number needed to reach the onset of the nonlinearity, measured in units of $\sigma_\delta$. Solving definition (38) for $p$ that occurs as an argument of $\delta$ and $\sigma_\delta$ and using Eq. (37) yields the connection probability in terms of $n$,

$$p_{NL} = \frac{n^2 \epsilon + 2\Theta_b + \sqrt{n^2 \epsilon^2 + 4\Theta_b (\epsilon - \Theta_b)}}{p_f(\kappa) \epsilon(n^2 + \omega) [1 + \text{Erf} \left( \frac{n}{\sqrt{2}} \right) \sigma^2]}.$$
corresponding to the group sizes $G_1$ and $G_2$, $p^{\ast}_{NL}$ has one preimage, and any $p_{NL}$ below $p^{\ast}_{NL}$ has none [cf. Fig. 2(c)]. Thus, $p_{NL}(n)$ has one global minimum at $n = n^\ast$ where $\frac{dp_{NL}(n)}{dn} \big|_{n_n} = 0$, and the critical connectivity is $p_{NL}(n^\ast) = p^{\ast}_{NL}$.

The comparison of the results for linearly and nonlinearly coupled FFNs is particularly enlightening in the limit of large layer size ($\omega \gg 1$) and small coupling strengths ($\epsilon \ll \Theta - \Theta_{reset}$). We fix the maximal input to a neuron from the previous layer, $\epsilon \omega = \text{const}$, to preserve the network state and expand Eq. (40) in a power series around $\omega \to \infty$ and $\epsilon \to 0$. Considering the leading terms, we find

$$p_{NL} = 2 \frac{\Theta_b + n \sqrt{\Theta_b (\epsilon - \Theta_b) \omega}}{p_{f}(\kappa) \epsilon \omega [1 + \text{Erf}(\frac{n}{\sqrt{2}})]},$$

(41)

We note that propagation of synchrony mediated by dendritic spikes is enabled if a sufficiently large fraction of neurons of each layer receives a total input larger than or equal to $\Theta_b$, which implies in particular that $\Theta_b < \omega \epsilon$. Moreover, if the connectivity within the FFN is low, stable propagation even requires that $\Theta_b \ll \omega \epsilon$, and $p_{NL}$ further simplifies to

$$p_{NL} = 2 \frac{\Theta_b}{p_{f}(\kappa) \epsilon \omega} + n \sqrt{\Theta_b \omega} \frac{1}{1 + \text{Erf}(\frac{n}{\sqrt{2}})}.$$  

(42)

As described above, the critical connectivity is given by the minimum of $p_{NL}$ as a function of $n$, which is assumed at $n = n^\ast$. $\frac{dp_{NL}(n)}{dn} \big|_{n_n} = 0$ yields $n^\ast$ as an implicit function of $\frac{\Theta_b}{\epsilon}$,

$$\sqrt{\frac{\Theta_b}{\epsilon}} = \frac{\sqrt{\pi}}{\sqrt{2}} \exp \left( \frac{n^\ast}{2} \right) \left[ 1 + \text{Erf} \left( \frac{n^\ast}{\sqrt{2}} \right) \right] - n^\ast.$$  

(43)

For better readability, we define

$$\beta \left( \frac{\Theta_b}{\epsilon} \right) := \frac{1}{2} \left[ 1 + \text{Erf} \left( \frac{n^\ast}{\sqrt{2}} \right) \right] - n^\ast e^{-\left( n^\ast / 2 \right)^2 / 2 \pi},$$

(44)

where $n^\ast = n^\ast (\Theta_b / \epsilon)$ as given by Eq. (43). Combining Eqs. (42)–(44) enables to simplify the critical connectivity to

$$p^{\ast}_{NL} = \frac{\Theta_b}{p_{f}(\kappa) \epsilon \omega} \frac{1}{\beta (\Theta_b / \epsilon)},$$

(45)

which depends nonlinearly on the number of spikes needed to reach the dendritic threshold $\Theta_b / \epsilon$ through the function $1 / \beta(\cdot)$. One can show that $\beta (\Theta_b / \epsilon)$ increases with decreasing coupling strength $\epsilon$ from $\beta (\Theta_b / \epsilon) = 0.5$ for large $\epsilon$ and becomes maximal in the limit of small couplings, $\lim_{\epsilon \to 0} \beta (\Theta_b / \epsilon) = 1$. Figure 3(b) displays the results for $p^{\ast}_{NL}$ together with the results of numerical simulations. As in the linearly coupled network, the critical connectivity decays with layer size and coupling strength, but the dependence on $1 / \epsilon$ is nonlinear. The factor

$$c := \frac{p^{\ast}_{NL}}{p_{NL}} = \frac{p_{f}(\kappa)}{\lambda \Theta_b} \beta \left( \frac{\Theta_b}{\epsilon} \right),$$

(46)

by which the nonlinear dendritic interactions reduce the required network-connectivity, increases with decreasing threshold $\Theta_b$ and increasing enhancement $\kappa$. Figure 3(c) illustrates the numerically obtained reduction of connectivity: The critical connectivity $p^{\ast}_{NL}$ is smaller over the whole parameter range; the reduction is most effective for small $\epsilon$ and largely independent of $\omega$.

Nonlinear dendrites thus foster propagation of synchrony. We note that our model still overestimates the capability of linearly coupled networks to propagate synchrony: On synchronous input, linearly coupled groups of neurons generate synchronous output (if they generate output at all). This fact is a consequence of the infinitesimally short synaptic currents. In neurons with extended synaptic currents, the timing of the output strongly depends on the neurons’ state and input strength. In contrast, the timing of somatic action potentials elicited by dendritic spikes is largely independent of neuron state and input strength. Therefore we expect the effect of nonlinear dendrites to be even stronger in networks of biologically more detailed neurons, as considered in the next section.

### C. Recurrent networks

The main findings generalize in two ways: to FFNs occurring in recurrent random networks and to biologically more detailed models. For such systems, we show that, in nonlinearly coupled networks, stable propagation naturally emerges, whereas it is difficult to achieve in linearly coupled networks. In contrast to isolated FFNs studied above, we now account for effects of the FFN on the surrounding network and its feedback. Further, we choose a more detailed neuron model (see Sec. II) to ensure that the main assumptions underlying the analytically tractable model are not crucial for stable propagation of synchrony. In particular, we show that systems with temporally extended postsynaptic responses and a temporally extended nonlinear dendritic interaction window exhibit qualitatively the same phenomena as found above.

We consider networks of randomly connected conductance-based leaky integrate-and-fire neurons [cf. Eqs. (8)–(10)]. The networks consist of $N^E$ excitatory and $N^I$ inhibitory neurons. A directed connection between two neurons is present with probability $p$. As for the isolated FFNs considered above, we construct the network such that the ground state in the absence of synchronous activity is characterized by balanced excitatory and inhibitory input, which results in an asynchronous irregular spiking activity. For simplicity, all neurons have the same parameters, e.g., $C_i^m = C^m$, $g_i^m = g^m$, etc.
First, we set up a model network in which the total excitation and inhibition to the neurons is balanced such that the spiking activity is asynchronous irregular. The external constant current $I^0$, together with the leak conductance $g^L$ and the resting potential $V^{\text{rest}}$, determines the asymptotic membrane potential in the absence of incoming spikes:

$$V^\infty = V^{\text{rest}} + \frac{I^0}{g^L}.$$  

(47)

Additionally, each neuron receives excitatory and inhibitory random Poissonian spike trains. The frequencies are denoted by $\nu^{\text{ext},\text{ex}}$ and $\nu^{\text{ext},\text{in}}$, and the ratio between them is chosen such that it equals the ratio of the number of excitatory and inhibitory neurons in the network:

$$\frac{\nu^{\text{ext},\text{ex}}}{\nu^{\text{ext},\text{in}}} = \frac{N^E}{N^I}. \quad (48)$$

This ensures that each neuron receives the same ratio of excitatory and inhibitory input from both the network and the external sources when neurons in the excitatory and inhibitory network populations spike on average with the same mean rate. All excitatory as well as all inhibitory connections have the same strength, i.e., $g_{lj}^{\text{max}} = g^{\text{ex}}$ for excitatory and $g_{lj}^{\text{max}} = g^{\text{in}}$ for inhibitory connections. The ratio of the peak postsynaptic potentials due to an inhibitory input and an excitatory input at the asymptotic membrane potential $V^\infty$ is approximately given by

$$g^\text{rat} := \frac{g^{\text{in}}[V^\infty - E^{\text{in}}]}{g^{\text{ex}}[V^\infty - E^{\text{ex}}]}.$$  

(49)

We set

$$g^\text{rat} = \frac{N^E}{N^I} = \frac{\nu^{\text{ext},\text{ex}}}{\nu^{\text{ext},\text{in}}}, \quad (50)$$

and

$$g^{\text{in}} = \frac{|V^\infty - E^{\text{ex}}|}{|V^\infty - E^{\text{ex}}|} \frac{N^E}{N^I} g^{\text{ex}}.$$  

(51)

to obtain balanced activity.

In contrast to the model considered in the first sections, now excitatory neurons have a nonzero time window $\Delta t$ for nonlinear dendritic modulation. When the strength of the excitatory input within $\Delta t$ exceeds a threshold, a current pulse is injected into the soma, modeling the effect of a dendritic spike. The neuron parameters for this phenomenological model are chosen according to experimental findings to reproduce quantitatively the time course of the membrane potential in response to a dendritic spike. [See Sec. II and Figs. 1(b) and 1(c).]

Considering a random network, we detect naturally occurring weak feed-forward structures suitable for signal transmission in the following way: We randomly choose a group of $x$ neurons to be the first layer. The second layer is composed of $x$ neurons out of those receiving the largest numbers of connections from the initial group. By repeating this selection process $l$ times, we identify a FFN consisting of $l$ layers. In each selection step, we exclude the $x$ neurons of the previous layer, but do not exclude neurons that are members of the layers preceding the previous one. The high-connectivity subnetwork selected from an existing random network as described above by construction has a slightly higher-than-average connection probability. Therefore, this structure is particularly well suited to enable propagation of synchrony. Alternatively, one can assign neurons randomly to the different layers and compensate for smaller connectivity by, e.g., larger layer sizes, according to Eq. (45).

The measurements start after an equilibration phase. (Initially, the network is at rest.) In the ground state, the network generates balanced irregular activity. Propagation of synchrony is initiated by exciting the neurons of the first layer to spike within a short time interval that is smaller than the time window of dendritic integration, $\Delta t$. This synchronous spiking leads to an increased input to the second layer after a delay time $\tau$. This input, in turn, may lead to highly synchronous spiking of a certain number of neurons of the second layer (possibly supported by dendritic spikes) and therewith to synchronous spiking after another delay time $\tau$ in the third layer, etc. Propagation of synchrony requires (i) that the total input of a layer to its successor within the FFN is sufficiently strong and (ii) that the input to the remaining network is sufficiently weak to avoid excitation of too many neurons to synchronous spiking. Requirement (ii) prevents pathological activity such as “synfire explosions” [6].

After initiating a propagation of synchrony by exciting the neurons of the first group to spike within a short time interval, we measure the probability of detecting a synchronous pulse in the subsequent groups [see Sec. II; cf. Figs. 4(a) and 4(b)]. Although the average connectivity within the identified FFN is significantly larger than the overall connectivity $p$, it is still small, and propagation of synchronous activity is very unlikely [upper insets of Figs. 4(a) and 4(b)]. We find that it is not sufficient to choose high-connectivity subnetworks as FFNs (as described above) to obtain a stable propagation of synchrony, but that the synapses within the FFN have to be strengthened. To study the transition to propagation, we gradually strengthen the synapses within the FFN. As suggested by the results on isolated chains, we observe a propagation of synchrony over more and more layers for moderate enhancements [Figs. 4(c) and 4(d)]. For very strong enhancements, the feedback from the network becomes important: The synaptic amplification leads to an increased spontaneous activity within the FFN, and this in turn results in an increased background activity. The overall increased spiking activity causes spontaneous synchronous pulses, and a separation of the induced synchronous signal from the
background activity is not possible anymore (the detection probability decreases; see Figs. 4(a) and 4(b) and lower insets).

In agreement with previous studies (cf. [7]), we find that, in the linearly coupled networks considered, a synchronous pulse propagates only over a few layers, even in the optimal enhancement range [Fig. 4(a)]. In contrast, networks incorporating nonlinear dendrites support stable propagation of synchrony [Fig. 4(b)] in a substantial region of parameter space. In addition, the propagation is enabled for enhancements considerably smaller than the optimal enhancement for networks with linear dendrites.

**IV. DISCUSSION**

In conclusion, we have analyzed strongly diluted networks with linear and nonlinear dendritic interactions. We have demonstrated how nonlinear dendritic interactions may enhance and stabilize synchrony propagation in both isolated feed-forward chains and recurrent network structures. Moreover, our results show that such local nonlinear interactions support the separation of propagating synchrony and asynchronous background activity. Earlier works [6,7] did not take into account supralinear amplification of synchronous activity. One study [7] used existing connections in recurrent networks to create diluted chains assuming strongly enhanced synapses and at the same time partially decoupling the chain from the rest of the network; still, synchrony could propagate only over a few groups. In contrast, the results presented here indicate that a reliable propagation is achieved by only mildly adapted synapses and without specifically tuning or changing neuron properties or rewiring the network.

The recent study [18] incorporating nonlinear dendrites has shown that synchronous activity can propagate in purely random networks without modified connections. There are no specific propagation paths, but neurons are recruited in a quasirandom manner. Our results described here now indicate that specific feed-forward chains that naturally occur in random neural circuits are capable of persistently propagating synchronous signals if their synaptic strengths are increased. The strengths required in the presence of nonlinear interactions are common in biological neural circuits [19] and may well be generated by learning, e.g., through spike-timing-dependent plasticity.

Dendritic (coupling) nonlinearities therefore offer a viable mechanism for guiding synchrony through weakly structured random topologies.

Recently [12], feed-forward chains with slow dendritic (probably calcium) spikes have been simulated to check the possibility of the occurrence of specific spike patterns that are experimentally observed in the higher vocal center of song birds. Our theoretical work now yields analytic insights about the collective dynamics of circuits with fast dendritic (sodium) spikes. Fast dendritic spikes have been found in the hippocampus and in the neocortex and may thus be involved in hippocampal replay, memory formation, and other computational processes. Experimentally, the influence of fast dendritic spikes could be directly

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**FIG. 4.** Nonlinear dendritic interactions enable guided propagation of synchrony in random networks ($N = 10000$, $p = 0.03$, $\omega = 30$). (a),(b) Detection probability of induced synchronous activity in (a) conventional networks and (b) in networks with nonlinear dendrites. Insets of (a),(b) and Figs. (c),(d) Typical examples of network activity for different synaptic-enhancement factors. Red and blue indicate spikes of neurons within the FFN; green indicates spikes of neurons randomly sampled from the remaining neurons.
checked by selectively blocking dendritic sodium channels (see, e.g., [20], which indicates that the types of sodium channels in the dendrite and soma are different) and thereby distinguishing those effects that come from nonadditive coupling via fast dendritic spikes from those induced by other mechanisms. During the last decade, the number of neurons simultaneously accessible has multiplied from a few to the order of 10² neurons, with this rapid trend continuing. When recording the activity of a substantial fraction of neurons of a local circuit, synchrony propagation should be clearly detectable and analyzable. Our results suggest that synchrony propagation and thus spike patterns should be influenced if dendritic sodium channels and thus fast dendritic spikes are blocked. Specifically, in the hippocampus, the precision of (replayed) spike patterns decreases or the patterns vanish after blocking. Such experiments would thus provide a direct test of how nonadditive coupling is exploited for the collective dynamics of neural circuits. Once the connectome, i.e., the structural synaptic connectivity, of neural circuits becomes available in the future [21], the relative impact of synaptic, structural to dynamic features of single neurons on circuit dynamics may be well distinguishable.

The basic model of pulse-coupled units considered here is applicable to a range of systems in nature, not only to neural circuits but also, e.g., to earthquakes emerging from abruptly relaxing tectonic plates, and fireflies interacting by exchanging light flashes (e.g., [22]). We have now studied the impact of nonlinear input modulation on collective network dynamics and derived methods for their analysis that may also be useful in a non-neuronal setting. Interestingly, very recent results [23] have shown that fireflies are more prone to respond to synchronous flashes rather than to asynchronous ones, suggesting a direct application of our model.

ACKNOWLEDGMENTS

This work was supported by the BMBF (Grant No. 01GQ1005B), the DFG (Grant No. Ti 629/3-1), and the Swartz Foundation. Simulation results were partly obtained using the simulation software NEST [26].

APPENDIX

1. Parameters for Figs. 2 and 3

The single-neuron parameters and the coupling delay are \( \tau = \frac{1}{\gamma} = 14 \text{ ms} \), \( \Theta = 15 \text{ mV} \), \( V_{\text{reset}} = 0 \text{ mV} \), \( t_{\text{ref}} = 2 \text{ ms} \), and \( \tau = 10 \text{ ms} \) [15,24]. The external input is characterized by \( \tau_{\text{ext}}^{\text{ext}} = -e_{\text{ext}}^{\text{lin}} = 0.5 \text{ mV} \) [19,25], \( \tau_{\text{ext}}^{\text{ext}} = \tau_{\text{ext}}^{\text{lin}} = 3 \text{ kHz} \), and \( V^{\infty} = 5 \text{ mV} \). The parameters of the dendritic modulation function were chosen according to single-neuron measurements as \( \Theta = 4 \text{ mV} \) and \( \kappa = 11 \text{ mV} \) [8].

The maps and transition matrices presented in Fig. 2 are derived for \( \omega = 100 \) and \( \epsilon = 0.4 \text{ mV} \). To obtain the distribution of active neurons \( g_{i+1} \) in layer \( i + 1 \), we excite \( g_{i} \) neurons of the first layer to spike simultaneously and measure the number of active neurons in the following layer. For each value of \( g_{i} \), we calculate the distribution for \( m = 1000 \) different realizations of the FFN and initial conditions.

In Fig. 3, existing connections within the FFN have strengths \( e_{ij} = \epsilon \). We determine the critical connectivity for \( \epsilon \in [0.05 \text{ mV}, 0.6 \text{ mV}] \) and layer sizes \( \omega \in [50, 600] \) as follows: We construct a FFN consisting of 20 layers, with \( \omega \) neurons in each layer, and connect neurons of successive layers with probability \( p \in [0, 1] \). After an equilibration time \( t_{\text{eq}} \) (initially, the network is at rest), we initiate propagation of synchrony by exciting all neurons of the first layer to spike simultaneously. We then check whether the synchronous pulse propagates up to layer \( i \), i.e., whether there is synchronous activity in layer \( i \) at time \( t_{\text{eq}} = t_{\text{eq}} + (i - 1) \tau \). We consider the propagation for a certain setup specified by \( \epsilon, \omega, \) and \( p \) to be successful if a synchronous pulse propagates along the whole FFN in more than 50% of \( \omega = 31 \) realizations of the FFN with different initial conditions. We derive the critical connectivities \( p_{L}^{L} \) and \( p_{L}^{NL} \) up to a resolution of \( \Delta p = 5 \times 10^{-3} \) by repeatedly bisecting the interval [0,1] and testing the success of propagation.

2. Parameters for Fig. 4

For the network simulations, we employed the simulation software NEST [26]. The networks had a total number of \( N = 10000 \) neurons with \( N^E = 8000 \) and \( N^I = 2000 \). For simplicity, all neurons are considered identical, i.e., \( C_{ij} = C_{i}, g_{ij}^{L} = g_{i}^{L}, V_{\text{reset}}^{L} = V_{\text{reset}}^{i}, t_{\text{ref}}^{i} = t_{\text{ref}}, \Theta = \Theta, t_{\text{ref}} = t_{\text{ref}} \). The single-neuron parameters are \( C_{i} = 400 \text{ pF} \), \( V_{\text{reset}} = V_{\text{reset}} = -65 \text{ mV} \), \( g_{i}^{L} = 25 \text{ nS} \), \( \Theta = -50 \text{ mV} \), \( t_{\text{ref}} = 3 \text{ ms} \) [24,27], and \( \omega = 250 \text{ pA} \), and the frequencies of the external inputs are \( \nu_{\text{ext}}^{\text{lin}} = 2.4 \text{ kHz} \) and \( \nu_{\text{ext}}^{\text{lin}} = 0.6 \text{ kHz} \). The recurrent connectivity in cortical and hippocampal networks is sparse: Connection probabilities between 1% and 10%, depending on the distance and the region, have been estimated (e.g., [19,24,25]). For our simulations, we choose \( p = 0.3 \).

The time constants of the excitatory (AMPA) conductances are \( \tau_{\text{exc}} = 2.5 \text{ ms} \) and \( \tau_{\text{exc}} = 0.5 \text{ ms} \) [28]. For simplicity, we choose the same time constants for the inhibitory (GABA) conductances: \( \tau_{\text{exc}} = 2.5 \text{ ms} \) and \( \tau_{\text{exc}} = 0.5 \text{ ms} \). The reversal potentials are \( E_{\text{exc}} = 0 \text{ mV} \) and \( E_{\text{in}} = -75 \text{ mV} \) [15,24]. The strengths of experimentally observed pEPSPs due to single inputs range from small values like 0.1 mV to larger values like 2 mV [19,24,25]. For nonenhanced couplings, we set \( g_{\text{in}}^{\text{exc}} = 0.6 \text{ nS} \), which corresponds to a pEPSP of approximately 0.3 mV at rest. According to Eq. (51), the coupling strengths of the inhibitory synapses are \( g_{\text{in}}^{\text{in}} = -6.6 \text{ nS} \) to maintain balanced input. This configuration results in an
asynchronous irregular ground state with a spontaneous firing rate $\nu = 1.8$ Hz.

The parameters of the dendritic spike current are chosen according to single-neuron measurements in hippocampal cells: $\Delta t = 2$ ms [8], $g^{\Theta} = 8.65$ nS (corresponding to a pEPSP of about 3.8 mV at rest [8]), $\tau_{\text{DS}} = 2.7$ ms (such that $\tau + \tau_{\text{DS}} = 4.7$ ms and the peak of the depolarization is reached approximately 5 ms after presynaptic spiking), $A = 55$ nA, $B = 64$ nA, $C = 9$ nA, $\tau_{\text{DS},1} = 0.2$ ms, $\tau_{\text{DS},2} = 0.3$ ms, $\tau_{\text{DS},3} = 0.7$ ms, and $\rho_{\text{det,DS}} = 5.2$ ms. The correction factor, which modulates the strength of the dendritic spike, is found by fitting a linear correction function, $c(g) = \max(1.5 - g \times 0.053 \text{nS}^{-1}, 0)$, such that the experimentally observed region of saturation is obtained. The dynamics of the neuron model incorporating the mechanism for dendritic spike generation is illustrated in Fig. 1.

For calculating the SNR, we use an $a = 0.99$ and $b = 2$ and an expected width of the synchronous pulse $t^* = 10$ ms; the result is insensitive to changes in these parameters. The expected interval between successive synchronous active layers, $\Delta t^{\text{exp}}$, is chosen from the interval [2 ms, 7 ms] such that the signal, $\sum S_i$, is maximized (cf. Sec. II). The time interval for the estimation of the noise level is $\Delta t^{\text{obs}} = 15$ s. The detection probability shown in Figs. 4(a) and 4(b) is the fraction of successful propagations obtained from 10 different network realizations, where, for each network setup, propagation of synchrony was tested for 20 initial conditions.

All measurements start after an initial equilibrium phase of $t^0 = 4000$ ms.


