



Information encoding and reconstruction from the phase of action potentials

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Fundamental questions in neural coding are how neurons encode, transfer, and reconstruct information from the pattern of action potentials (APs) exchanged between different brain structures. We propose a general model of neural coding where neurons encode information by the phase of their APs relative to their subthreshold membrane oscillations. We demonstrate by means of simulations that AP phase retains the spatial and temporal content of the input under the assumption that the membrane potential oscillations are coherent across neurons and between structures and have a constant spatial phase gradient. The model explains many unresolved physiological observations and makes a number of concrete, testable predictions about the relationship between APs, local field potentials, and subthreshold membrane oscillations, and provides an estimate of the spatio-temporal precision of neuronal information processing.

Keywords: neural coding, action potential, local field potential, subthreshold membrane potential oscillations, gamma

INTRODUCTION

Whatever code neurons use for encoding, transferring, and decoding sensory information in the central nervous system (CNS) must be robust to a number of compromising factors. The integrity of the information transferred in massively parallel pathways of the brain is highly sensitive to distortions from different conduction delays (Rockland et al., 1997) and from intrinsic correlations due to the divergence and convergence of axonal projections (de la Rocha et al., 2007). Nevertheless, neuronal responses are increasingly specific to objects and progressively invariant with respect to incidental physical features, the times of occurrences, and the spatial locations of stimuli as we follow the activation from the sensory neurons through the primary sensory and ultimately to associational cortical areas. The high specificity of neuronal representations in higher cortical and associational areas relies on an unknown mechanism that enables the generation of action potentials (APs) in the same neuron consistently and independent of its neighbors (“sparse coding”) (Quiroga et al., 2005). This requires precise coincidences of membrane depolarization with the arrival of excitatory postsynaptic potentials (EPSPs) at the level of individual neurons (Abeles, 1982). I investigated through numerical simulations whether the phase of APs relative to the intrinsic subthreshold membrane potential oscillations (SMOs) could encode and retain information with high spatial and temporal selectivity. Our model relied on two basic assumptions: first, that the timing of AP initiation is dependent on the phase of the SMO (Llinas et al., 1991); and second, that SMO is nearly synchronized with a phase gradient across a local population of neurons (Benucci et al., 2007; Bringuier et al., 1999; Grinvald et al., 1994; Prechtl et al., 2000). To relate the second assumption to empirical data, we further assumed that the SMO correlates with local field potentials (LFP, Buzsáki et al., 2003; Lagier et al., 2004; Leung and Yim, 1986). The results derived from the model are consistent with empirical data, and they

provide a unified framework to answer a diverse set of daunting problems (see Table 1 in Supplementary Material).

The indirect evidence for the tight relationship between AP timing and SMO derives from observations of high correlation between extracellular APs and LFP. High coherency between APs and LFP oscillations is predominant at the gamma and theta frequency bands in the awake brain (Bragin et al., 1995; Chrobak and Buzsáki, 1998; Kamondi et al., 1998; Womelsdorf et al., 2007), and phase lock between APs and high frequency LFP occurs during slow wave sleep (Buzsáki et al., 1992; Chrobak and Buzsáki, 1996; Cisse et al., 2007). Gamma and theta oscillations also phase lock in the hippocampus and in the entorhinal cortex during active exploration of the environment and processing of sensory input (Chrobak and Buzsáki, 1998; Robbe et al., 2006). At the same time, APs phase lock with gamma oscillations (Chrobak and Buzsáki, 1998). Moreover, gamma band LFP and AP coherence has been found to be sensitive to motor task (Mehring et al., 2003) and preparatory activity during working memory (Pesaran et al., 2002) and related to selective visual attention (Fries et al., 2008). The autocorrelogram of multiunit activity in visual cortical areas of the cat is strongly modulated at 40–60 Hz and correlates with the gamma LFP, suggesting a functional link between AP generation and gamma oscillations (Gray and Singer, 1989). The multiunit–multiunit and the multiunit–LFP coherences are both increased during visual stimulation in the gamma (Womelsdorf et al., 2007) and during decision making in the beta frequency bands (Pesaran et al., 2008). Although AP–LFP phase coherency has been observed within the low-frequency LFP bands in the primary visual cortex of anesthetized primates (Montemurro et al., 2008), the feature-dependence of AP–LFP coherency in primary sensory areas is yet to be investigated. Notably, APs phase lock to 20–40 Hz LFP in the primate somatosensory cortex (Murthy and Fetz, 1996). The majority of pyramidal cells fire at the trough of the theta cycle and align to

the gamma (Bragin et al., 1995), and specific types of interneurons also phase lock to gamma (Tukker et al., 2007). Moreover, pyramidal cell firing in the hippocampus and entorhinal cortex of rodents also express a progressive theta phase precession consistent with the animal's position relative to the place field (O'Keefe and Recce, 1993) leading to the proposal that the AP phase relative to theta may encode the precise location of the animal (O'Keefe and Burgess, 2005; O'Keefe and Recce, 1993; Skaggs et al., 1996). All previous models of hippocampal phase coding assumed that phase precession is generated inside of the hippocampus as a result of interaction between two slightly different frequency theta oscillations (Blair et al., 2008; Lengyel et al., 2003; O'Keefe and Burgess, 2005); phase precession originating from the phase-coded sensory input and imposed on the hippocampus by the entorhinal cortex has not yet been proposed. Likewise, oscillatory interference models, relying on multiple oscillators, were proposed to explain grid cell behavior in the entorhinal cortex (Burgess, 2008; Burgess et al., 2007). However, as we show, coincidences of phase-modulated sensory input with a single propagating oscillation field are able to generate grid topography without further assumptions.

The mechanism by which local SMOs synchronize and generate coherent LFP relies on inhibitory interneurons (Lagier et al., 2004; Soltesz and Deschenes, 1993; Traub et al., 1996). To simplify terminology, we refer to LFP as the extracellular indicator of SMO; however, the performance of the model does not depend on the equivalence of LFP and SMO.

The aim of this study is twofold. First, we present a model that explains how feature-dependent phase coding may be employed in sensory information processing. Second, we investigate potential benefits of phase coding through examples, including generalization of 1-D phase coding to 2-D fields, information compression, selective reconstruction from combined representations, retaining stimulus invariants, and spatio-temporal transformations.

MATERIALS AND METHODS

COMPUTATIONAL MODEL

The core computational model consists of three layers, an input layer, a transformation layer, and an output layer. Each layer contains n neurons arranged in a 1-D array or a 2-D grid structure. The sensory input (1-D or 2-D) maps to the first layer. The first two layers are connected through n one-to-one unidirectional connections. In contrast, the second and third layers are connected through a single channel. Thus, the complete architecture is an n -to- n to one-to- n feed-forward network. For the sake of simplicity, we first consider a 1-D array of neurons to introduce the basic operations, and then we demonstrate that these operations are applicable to a 2-D grid as well.

I modeled neurons as deterministic state machines with binary AP or non-AP states and a real-valued baseline function $\varphi(t)$ representing the voltage of the SMO. The state of each neuron is defined by the combination of these two parameters. Although time is a continuous variable in the model, for numerical simulations we applied a discrete timescale ~ 0.2 ms ($T_\gamma = 0.01$).

We denote $\varphi_i(t)$ as a wave function with frequency γ and period $T = 1/\gamma$ corresponding to each neuron's SMO. We assume that $\varphi_i(t)$ is coherent among the neurons of a given layer with a constant phase offset $d_i\Delta\Phi$ proportional to the distance of the

neurons from the origin of the oscillation (d_i), like a propagating wave

$$\varphi_i(t) = \varphi_0(t + d_i\Delta\Phi), \quad (1)$$

where $\Delta\Phi$ has units of time/distance. For an n -dimensional grid of neurons ($n = 1, 2, 3$), this phase gradient causes the neurons to express a coherent but phase-shifted n -dimensional oscillation field, where the phase difference between grid points is proportional to the difference between their distances from the center of oscillation (Figure 2E). At grid points where the phase difference reaches 2π , neurons have a zero phase-lag difference. The phase gradient starts to increase from homogeneously distributed centers and attenuates at the edges where spreading oscillations derived from adjacent centers meet (Benucci et al., 2007; Rubino et al., 2006).

The sequence of events from encoding to decoding are as follows:

- (1) Encoding phase (layer 1): First we generate a 1-D input pattern \mathbf{S} , a stimulus vector of length n representing the intensity of the stimulus acting on each of the respective n neurons (Figure 1 stage 1). We now define \mathbf{A} as the pattern of APs induced by \mathbf{S} . \mathbf{A} is an $n \times n$ binary matrix with elements $\mathbf{a}_{i,t}$, representing the time t at which each neuron i receives an input, that is computed by converting the intensity vector \mathbf{S} to a latency vector such that a higher intensity input induces a shorter latency AP (Figure 1 stage 2) (Hopfield, 1995).

$$\mathbf{a}_{i,t} = f(\mathbf{s}_i) \quad (2)$$

The function f assigns each input value \mathbf{s}_i to a time \mathbf{t}_i , where

$$\mathbf{t}_i = \frac{n}{\gamma \cdot \mathbf{s}_i} \quad (3)$$

and γ is the frequency of SMO. The role of n is to scale the AP times to encompass multiple SMO periods. Importantly, the exact function in Eq. 2 does not matter as long as the latency is a monotonic function of the \mathbf{s}_i input. Moreover, it does not matter whether the input is sensory or it was derived from another brain region; the model is only concerned with the pattern of APs that is induced by \mathbf{S} .

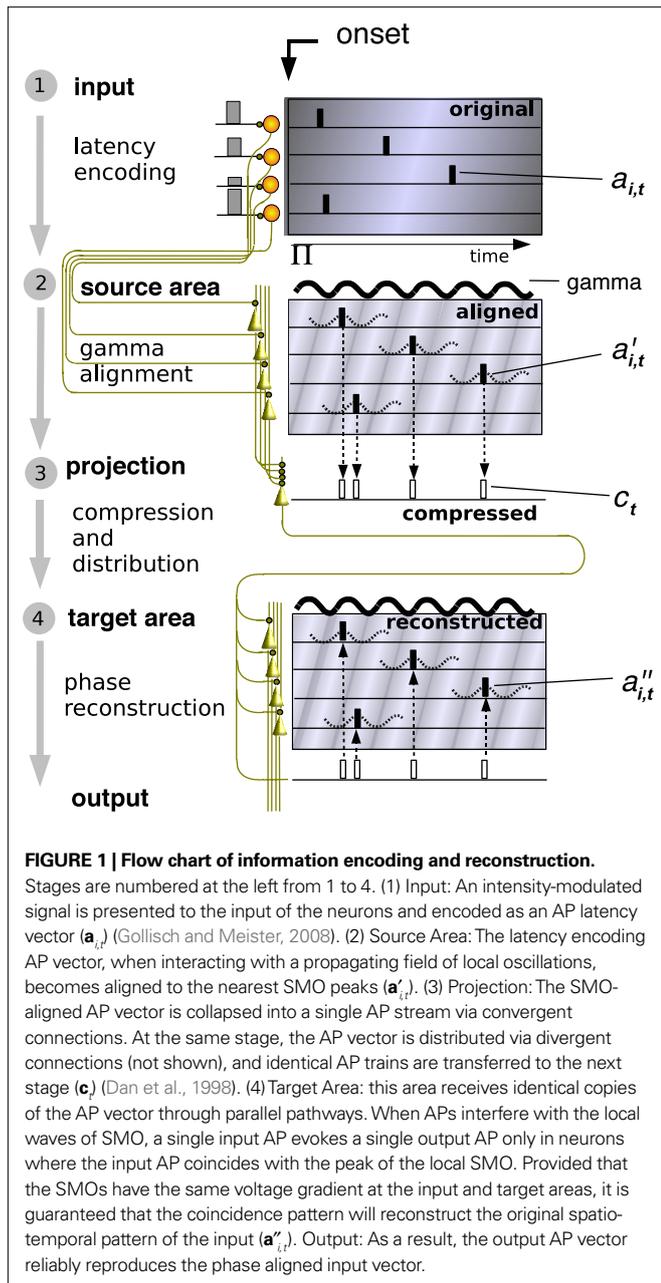
- (2) Transformation phase (layer 2): Now, we define \mathbf{A}' as the gamma-aligned pattern of APs in \mathbf{A} . More precisely, \mathbf{A} is transformed into \mathbf{A}' by delaying each AP until the neuron reaches the local maximum of its SMO. As a result, each i -th neuron generates an AP at a peak of $\varphi_i(t)$. We implement this delay by aligning each binary AP event to the peak of the respective neuron's intrinsic oscillation $\varphi_i(t)$ (Eqs. 4 and 5).

$$\mathbf{a}'_{i,\tau_i} = \mathbf{a}_{i,t} \quad (4)$$

where

$$\tau_i = \varphi_i^{-1}(\max[\varphi_i(t)]) \quad (5)$$

and \max is defined as the next local maximum of $\varphi_i(t)$ after the time of the received input $\mathbf{a}_{i,t}$. Here \mathbf{a}'_{i,τ_i} are the elements of \mathbf{A}' , a matrix with binary elements, where columns correspond to 0.1 ms time bins and rows correspond to specific neurons.



A “1” in position \mathbf{a}'_{i,τ_i} represents an AP for neuron i at time t while a “0” represents the lack of AP (Figure 1 stage 3).
 (3) Transfer (layer 2–3): Then \mathbf{A}' is collapsed into an array \mathbf{C} by summing across all neurons for each time bin.

$$\mathbf{c}_t = \sum_{i=1}^n \mathbf{a}'_{i,\tau_i} \quad (6)$$

Since the sum is a binary function, \mathbf{C} contains the APs derived from all neurons with their original time stamps. Although the neuronal specificity is lost in the sum, the precise timestamps of APs unambiguously relate the individual APs to specific locations through their phases. Thus, the spatial information is retained by the relative phases of APs (Figure 1 stage 3).

At the same step, \mathbf{C} was “cloned” to n copies and fed into the next layer of n neurons.

$$\mathbf{c}_j = \mathbf{C}, \quad (7)$$

where $j = \{1, \dots, n\}$. The information in \mathbf{C} is transferred from the input layer to the target layer. Since all the information from the input layer neurons is collapsed into a single channel (stage 3), this information is vulnerable to errors deriving from different conductivity delays. Therefore, a redistribution of \mathbf{C} over multiple channels is necessary to increase redundancy for error correction.

(4) Reconstruction (layer 3): The final step is reconstruction. Here the task is to create a matrix \mathbf{A}'' by assigning each individual AP in \mathbf{C} to a neuron such that there is a topographic mapping between \mathbf{A} and \mathbf{A}'' , consequently recovering the original spatio-temporal structure of APs from the compressed code. To do so, we simultaneously broadcast the \mathbf{c}_j copies to each k -th neuron ($k = \{1, \dots, n\}$) in the target layer. Neurons in the target layer will only generate APs if the input \mathbf{c}_j coincides with a peak of their membrane oscillation $\phi_k(t)$ within a precision of $\Delta\tau$ (Figure 1 stage 4 and Movie 1 in Supplementary Material). Given that the gamma phase gradients are identical between the source and the target (but see Figure 1 in Supplementary Material), these coincidences must occur in neurons at the same grid position within the target layer as the grid position of neurons stimulated in the input layer. Consequently, the topography and timing of the coincidences between the \mathbf{C} and the SMO, which generates the APs at the target layer, will reproduce the topography and timing of the input APs. To implement this, we define a matrix \mathbf{B} with $n \times T/\Delta\tau$ binary elements, where 1-s represent the positive peaks of SMO and 0-s represent all other voltage values, with a $\Delta\tau$ precision:

$$\mathbf{b}_{i,\tau} = \begin{cases} 1, & \text{if } \text{mod}(t + d_i \Delta\Phi, \gamma) = 0 \\ 0, & \text{otherwise,} \end{cases} \quad (8)$$

where $\mathbf{b}_{i,\tau}$ represents the binary state of the i -th the neuron at τ time. Since the input layer SMO is identical to the target layer SMO, it is necessary that the intersection \mathbf{A}'' of \mathbf{B} and \mathbf{C} :

$$\mathbf{A}'' = \mathbf{B} \cap \mathbf{C} \quad (9)$$

will reproduce the original matrix \mathbf{A}' (Figure 1 stage 4). The output pattern \mathbf{A}'' can be transferred as input to other target areas. Equation 9 will be referred to as the “interference operator”, and Equations 8 and 9 as the “interference principle”.

SELECTIVE RECONSTRUCTION FROM COMBINED PHASE CODES

Let us define input images \mathbf{S}^A and \mathbf{S}^B and encode them separately as \mathbf{A}^α and \mathbf{A}^β . Next, align \mathbf{A}^α and \mathbf{A}^β to field gradients $\Delta\Phi_\alpha$ and $\Delta\Phi_\beta$, respectively, and compress the aligned \mathbf{A}''^α and \mathbf{A}''^β matrices into corresponding \mathbf{C}^α and \mathbf{C}^β representations. Next, we combine the two compressed representations as \mathbf{C}^γ by simply merging and sorting the AP times in ascending order:

$$\mathbf{C}^\gamma = \mathbf{C}^\alpha \cup \mathbf{C}^\beta. \quad (10)$$

Given that $\varphi^\alpha(t)$ has a phase gradient $\Delta\Phi_\alpha$ and $\varphi^\beta(t)$ has a phase gradient $\Delta\Phi_\beta$, the two different SMO fields must be able to selectively reconstruct the two AP patterns, $\mathbf{A}^{\prime\alpha}$ and $\mathbf{A}^{\prime\beta}$, both retaining the main features of \mathbf{A}^α and \mathbf{A}^β , respectively:

$$\mathbf{A}^{\prime\alpha} = \mathbf{B}^\alpha \cap \mathbf{C}^\gamma \quad (11)$$

and

$$\mathbf{A}^{\prime\beta} = \mathbf{B}^\beta \cap \mathbf{C}^\gamma. \quad (12)$$

With simulations (Figure 5) we demonstrated that given $\Delta\Phi_\alpha \neq \Delta\Phi_\beta$, \mathbf{A}^α can be recovered from the interference of \mathbf{B}^α and \mathbf{C}^γ without confusion with \mathbf{A}^β . Likewise, matrix \mathbf{A}^β can be recovered from the interference of \mathbf{B}^β and \mathbf{C}^γ without confusion with \mathbf{A}^α .

This mechanism provides the capacity to combine information from different sources within the same target layer and selectively recall them by different SMOs.

NUMERICAL IMPLEMENTATION OF THE MODEL

Numerical simulations were implemented using Matlab (Mathworks Inc., Natick, MA, USA). For the numerical implementation, instead of simulating spiking neurons with incremental time, we computed the state of all neurons in time at once using matrix operations. We remark that it was unnecessary to introduce a spiking neuron model since all the state transitions were deterministic and the interactions were linear. Thus, the state changes of neurons, i.e., the AP times, at a given layer as a function of the state changes at the subordinate layer were analytically solvable. Therefore, numerical simulations were performed by computing the complete AP time vectors as a function of the AP time vectors in the previous layer instead of deriving the membrane voltage V_t at every time step recursively from V_{t-1} .

Architecturally, the neuronal network consisted of three layers: input-layer, projection-layer and output-layer. The input layer contained n or $n \times n$ neurons, individually assigned to each pixel of the input image. We implemented the algorithm on feed-forward networks using three types of projections between layers: (i) A 1-D vector of n values (input vector) was projecting to a 1-D array of n input-layer neurons, which were connected through projection neurons to the same number of output-layer neurons. The connectivity between the values of the input vector and input-layer neurons was one-to-one; between the input-layer and projection-layer was n -to-one; between the projection-layer and output-layer was one-to- n . (ii) A 2-D input image, digitized as $n \times n$ pixels, projecting on a 1-D array of n^2 input-layer neurons. The architecture from the input-layer to the output-layer was similar as in (i), however, the output was visualized in 2-D to compare the reconstructed pattern with the input image. (iii) A 2-D ($n \times n$) input matrix was projecting on a 2-D grid of $n \times n$ input-layer neurons by emulating “receptive fields”. For this, the input matrix was partitioned into n subfields and the input-layer neurons were arranged into n non-overlapping groups that processed the n input subfields separately. Likewise, the output-layer neurons were arranged into n number of groups, emulating a topographical projection from the sensory organ to a primary sensory cortical area. The precise connectivity pattern between the input matrix and input-layer group was repeating across the groups but it was varied from simulation to simulation (see next paragraph). All neurons from a given group were

converging on a single projection neuron, which in turn, terminated on all the neurons of a given output-layer group. Thus, each input-layer group was interfacing with the corresponding output-layer group through a single projection neuron.

Operatively, the input was either a 1-D (n) or a 2-D ($n \times n$) matrix mapped to the input-layer neurons. The correspondence between the input matrix elements and the input-layer neurons was one-to-one. The 1-D simulations always used random vectors as input with the range of values from 0 to 2π and transferred to n neurons. As input to the 2-D simulations, we used a set of digitized images, including photos of people, animals, complex structures of rubber bands and grid patterns. They were selected to test the accuracy and geometric preservation of the reconstruction. Images were digitized with 256×256 pixel resolution and converted to grayscale. We down-sampled them to fit to the neuronal grid with a square geometry (16×16 , 18×18 , 32×32 , 36×36 , 60×60) and input them to the neurons. The input layer contained n or $n \times n$ neurons, individually assigned to each pixel of the input image. As described earlier, the projection-layer was different for the 1-D and 2-D network. For a 1-D solution, the projection-layer contained only one neuron (one channel). All the APs from the n input layer neurons were collapsed into a single channel, serving as a projection neuron. For the $n \times n$ solution, the projection layer contained n neurons (channels), thus the $n \times n$ grid of input layer was broken down into m subfields (receptive fields) and the APs from all the input neuron of a given subfield were collapsed into a single channel. The combined m channels served as a projection pathway. All the neurons at a given subfield of the output layer received a common input from a single projection neuron, containing the combined activity from the input subfield. Different mappings between the input and input-layer neurons were implemented by connection patterns, emulating different receptive field geometries (for details on modeling different receptive field architectures, see Supplementary Material).

The SMO was modeled by sinusoids. The phase gradient was either linear, emulating a propagating wave with a constant time lag of SMO between neighbor neurons, or non-linear, emulating oscillations deriving from randomly dispersed sources (for details on constructing non-linear SMO fields, see Supplementary Material below). APs were modeled as binary events. The duration of these events was the smallest time unit of the simulation ($\Delta\tau$), defined as a constant fraction of π , the half period of the SMO. Alignment to the nearest SMO peak was implemented by reassigning each AP to the nearest subsequent peak of SMO. Combining before transferring APs via the projection neurons was done by concatenating the AP time series. Coincidences at the output layer were detected when the latency difference between the incoming AP pulse and the peak of local SMO is $< \Delta\tau$. APs were assigned to neurons of the network where coincidences occurred. The output was the temporal pattern of APs on the grid of $n \times n$ neurons, converted to an image by an inverse of the image conversion applied to the image when it was converted to a temporal pattern. To quantify the efficiency reconstruction, we computed Pearson's correlation coefficients on the AP times between the input patterns and output patterns, neuron-to-neuron, as well as between the SMO-aligned input patterns and the output patterns. The implementation

of the algorithm for numerical simulations is described in the Supplemental Computational Methods. Demo codes written in Matlab are available at the following URL: <http://www.vis.caltech.edu/~zoltan/mcodes/phasecode.tgz>

ALGORITHM AND BIOLOGICAL MOTIVATIONS

The model that we introduce explains how oscillations might be used in the brain to coordinate APs to encode and reconstruct information between distant cortical areas. First, we distinguish four information processing stages and corresponding neuronal layers representing the results of each stage (**Figure 1**): (1) Input layer: latency encoding; (2) Source layer: gamma alignment; (3) Projection: compression and distribution, (4) Target area: reconstruction. The output of the target area may serve as an input to a connected area. The input layer may represent a group of sensory neurons or a group of cortical pyramidal cells. Postsynaptic to these neurons are neurons that encode the input and exhibit intrinsic SMO. Postsynaptic to encoding neurons are the projection neurons, such as sensory ganglia or long-range cortico-cortical connections. Postsynaptic to projection neurons are the target neurons, such as cortical layer-4 granule cells or layer 2–3 pyramidal neurons, which are the ultimate targets of sensory or cortico-cortical projections.

Each neuron was described by its grid location and layer, its connections, and its state. The state was defined by three variables: a continuous oscillating membrane potential implementing the SMO of the soma, a binary input, and a binary output, corresponding to AP or no AP. The probability of generating an AP was dependent on the input state and the actual SMO, which determined how close the membrane potential was to the threshold. In the absence of input, the membrane potential remained subthreshold, by definition. The SMO for each neuron was modeled by a sinusoid, and the grid of neuronal SMOs formed a 2-D field (SMO field) with a common frequency and a uniform phase gradient between adjacent neurons. Thus, the SMOs of neighboring neurons were near-synchronized, with a distance dependent phase delay, mimicking a spreading wave (Benucci et al., 2007; Bringuier et al., 1999; Grinvald et al., 1994; Prechtl et al., 2000). Each neuron generated an AP when the input, received from a presynaptic neuron (i.e., from an earlier stage), coincided with the neuron's own local SMO maximum, where the neuron's membrane potential is nearest threshold. We implemented a task in which these neurons encode and reconstruct an image using exclusively a feed-forward architecture in four steps corresponding to the four stages. For the sake of simplicity, we first illustrate these steps on a 1-D example, after which we extend it to 2-D examples using real images (see Section "Results").

Encoding

First we generate a 1-D input pattern **S** that represents intensity values of an image (**Figures 2 and 3A**). Each element of the **S** vector is assigned to the input of a neuron. Next, we compute the neuronal responses to **S**. We employ a latency encoding scheme (Gollisch and Meister, 2008; Hopfield, 1995), which converts the intensity values to AP latencies, inversely proportional to the intensity of stimulus component, generating a spatio-temporal activity vector **A** (Eqs. 2 and 3).

Gamma alignment

The latency-encoded pattern **A** is transferred to the next layer of neurons exhibiting a coherent field of SMOs. The SMO field consists of local oscillations spreading through the grid of neurons as a linear or radial wave with a neuron-specific phase gradient (Eq. 1). A neuron of this layer generates APs only when the sustained input from a neuron at the previous layer coincides with its own next SMO maximum. As a result, the temporal pattern of **A** is transformed in an SMO aligned pattern **A'** (Eqs. 4 and 5). Since we considered gamma oscillation as one of the main frequency components of local SMOs, we refer to this operation as "gamma alignment" (**Figures 1 and 2B**), though phase coding does not depend on the frequency of SMO. Note, that with gamma alignment this model is critically different from synfire chain (Abeles, 1991) and AP-latency encoding (Hopfield, 1995). Gamma alignment is possible through rapid spike-time-dependent-plasticity (Bi and Poo, 1998; Cassenaer and Laurent, 2007; Markram et al., 1997).

Compression

A' is then collapsed across all neurons into one AP sequence **C** by eliminating the neuronal identity (**Figures 1 and 2C**, and Eq. 6). This AP sequence represents all the APs pooled across all of the neurons. Pooling can be implemented by convergent feed-forward synaptic connections on a single neuron. Although neuronal identities are seemingly lost by collapsing APs over neurons, the spatial origin of the AP is still referenced by the timing, which associates the AP with the original location of the neuron through the gamma alignment. A phase difference of the local SMO, to which the AP is aligned, is preserved by the small latency differences between APs after collapsing them. The resulting new sequence is "compressed" because information from all neurons were collapsed into a single sequence of APs of the same duration, and "compact" because it contains all the information about **A'**, including time and location.

Distribution

At the same stage, the collapsed AP vector is copied over the same number of projection neurons as in **A** (Eq. 7). Compression and distribution are accomplished in a single step within a feed-forward network by convergent and divergent synaptic connections, respectively. As a result, all APs from the original **A'** pattern are represented in all the projection neurons by pattern **C**, with a redundancy (**Figures 1 and 2C**). These packets of **C** patterns can be broadcasted to many neurons simultaneously or exchanged between distant cortical areas. Since all projection neurons carry the same AP sequence, any of them is able to broadcast the complete information indiscriminately to any brain area. Notably, after compression the information content is no longer sensitive to different conduction delays. Errors deriving from differential conduction delays are self-corrected by synchronization of APs in crossed-over feed-forward pathways, typical in thalamo-cortical, callosal and long-range associational projection systems.

Reconstruction

This is the final stage where the original information **A'** (the gamma-aligned representation of **A**) is reconstructed from the pattern **C** as input to this layer. The parallel pathway of projection neurons

broadcasts the complete **C** input to each neuron at the target layer, such as layer 4 of the cerebral cortex. The target layer neurons also express an SMO field similar to the one at the input layer. Here, we apply a principle similar to the input layer, except a neuron only generates APs if the input coincides with the neuron's SMO maxima within a $\Delta\tau$ time window (Figures 1 and 2D). Such coincidences between the input and the local SMO occur only in selected neurons, which are located at the same grid positions as the active input-layer neurons (Movie 1 in Supplementary Material). Since the relative timing of AP is preserved, the coincidences reproduce the original **A'** pattern due to the "interference principle" (Eqs. 8 and 9). Therefore, to achieve a near perfect reconstruction, it is assumed that the SMOs at the input layer and the target layer share similar spatial and spectral properties, including the frequency and phase gradient topography (but see isomorphic reconstruction and Figure 1 in Supplementary Material). Since the interference occurs at selected neurons, the input pattern **C** will activate a sparse set of neurons, with topographic positions similar to the original neurons. Therefore, it is necessary that the reconstructed pattern **A''** resemble closely to the gamma-aligned original pattern **A'** and consequently, the original pattern **A** as well. The reconstructed pattern can be combined with patterns deriving from other input structures and a new cycle of encoding, gamma-alignment, compression and reconstruction is initiated that propagates the now combined representation to a different brain area (Figure 5).

RESULTS

For the sake of simplicity and algorithmic clarity, we first used a 1-D encoding and reconstruction example. A vector containing 16 random real numbers from 0 to 1 was used as the input to the network. Next, this input was converted to latencies within the range of eight SMO cycles (Figure 2A). This pattern was aligned to the neurons' own SMO which slightly altered the precise temporal pattern of APs relative to the original latency code (Figure 2B). This spatio-temporal pattern was collapsed into a single AP train by removal of the spatial dimension. At the same time the resulting AP string was distributed over 16 neurons which transferred the code from one brain structure to another with redundancy (Figure 2C). One of these AP strings was applied as input to a target network, consisting of 16 neurons. In addition to the input, we rendered each neuron with an SMO that shared the same phase gradient and frequency as the SMO field at the input layer. Although each individual neuron received the same AP sequence as input, due to the interference principle (Eqs. 8 and 9) only those APs that "precisely" coincided with the neuron's own SMO were effective in generating output APs (Figure 2D). We quantify the precision of coincidence later. As a result the output-layer neurons reproduced the spatio-temporal pattern of the original gamma-aligned input with high fidelity.

To demonstrate the algorithm in 2-D we used a set of grayscale images sampled at neuron resolution and projected them on the input-layer neurons so that each neuron processed one pixel, allowing no interactions between neurons of the same layer (Figure 3). By varying the precise projection between the image pixels and neurons we were able to test different receptive field configurations (see Supplementary Material on modeling different receptive field architectures). Each set of input-layer neuron

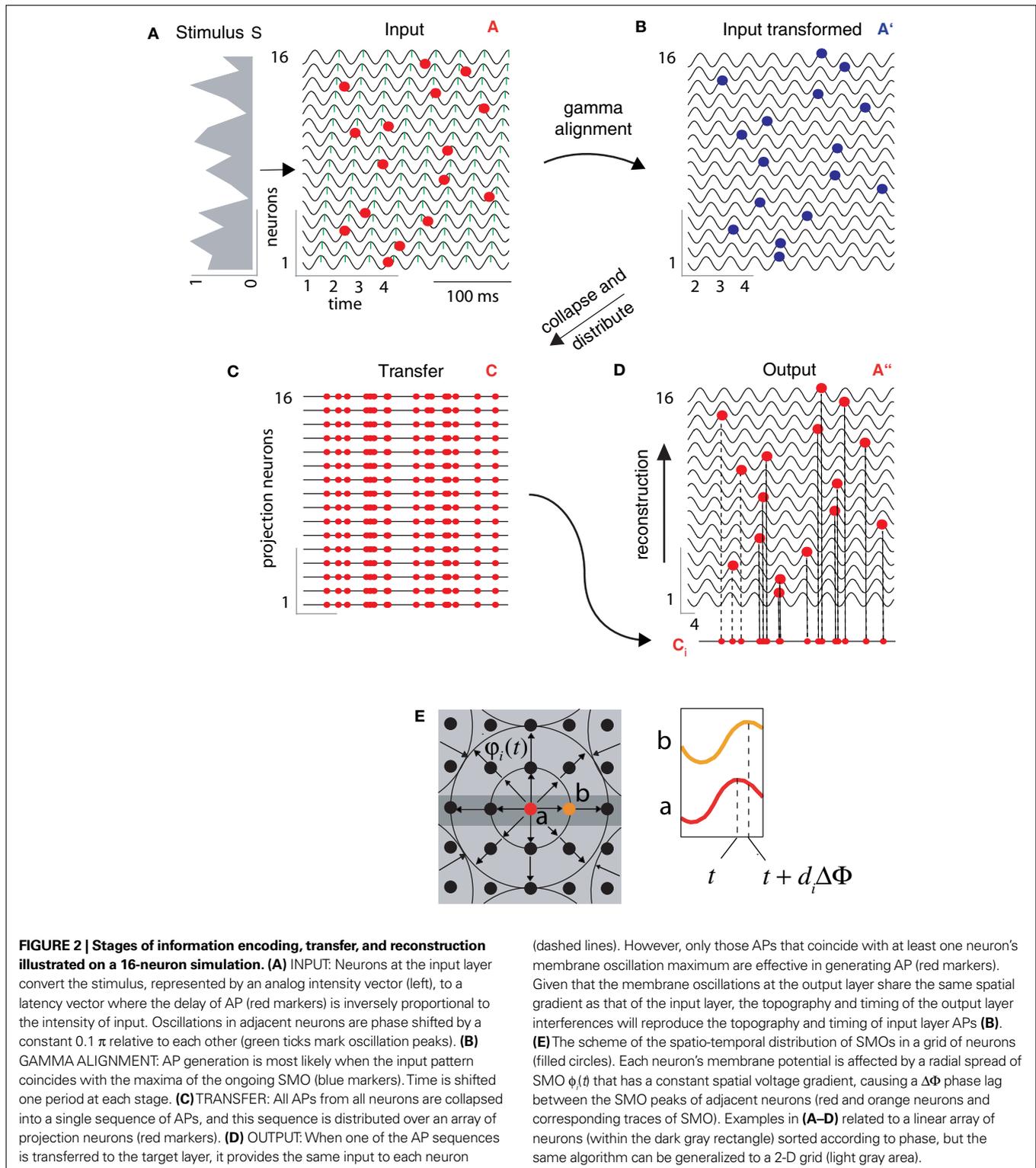
processed the receptive field independently throughout the four stages, and at the end, the reconstructed patterns were combined from each set. Finally, AP latencies were converted to grayscale values (Figures 3A,E). The close resemblance of the output image to the original implies that most of the information encoded from the input were correctly recovered from the phase of APs alone, thus phase coding is efficient.

BIOLOGICAL SIGNIFICANCE OF MODEL PARAMETERS

The efficiency of reconstruction was sensitive to a number of parameters, including the connectivity, the input size, the density of neurons, the SMO oscillation, and the time window of coincidence. The operative range of these parameters must be consistent with that measured in the living brain in order for the model to be biologically relevant. To understand how these parameters influence the reconstruction efficiency, we performed simulations. The independent variables were the number of neurons, the number of gamma cycles (i.e., the duration of reconstruction), the temporal resolution $\Delta\tau$ (equivalent to the time window of coincidence), the phase gradient $\Delta\Phi$ of gamma oscillation, the connectivity pattern, and the complexity of receptive fields (the number of pixels covered). The dependent variable was the reconstruction efficiency, quantified as the pixel-to-pixel Pearson's correlation coefficients either between the original image **A** and the reconstructed image **A''** ($r = r_{A,A''}$) or between the gamma-aligned image **A' A''** ($r' = r_{A',A''}$).

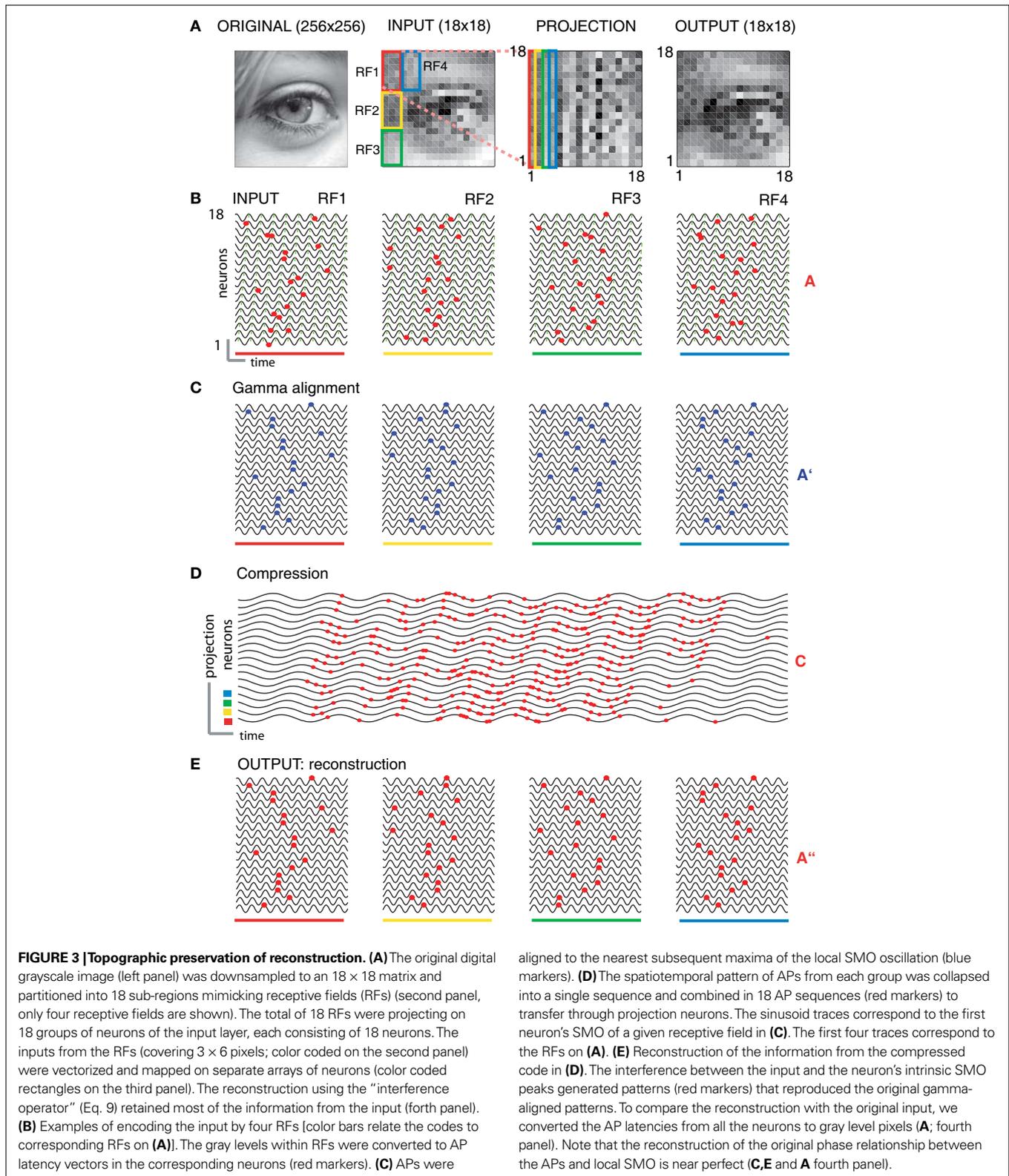
First, r and r' were tested against combinations of number of neurons and number of gamma cycles. The reconstruction efficiency is expected to inversely scale with the number of neurons since the more neurons' APs are combined in a single projection neuron, the more confusion occurs due to spurious interferences (aliasing errors). A few gamma cycles were expected to be sufficient to reconstruct the most of the information and certainly less gamma cycles than neurons are necessary. The steep exponential function [$f(x) = 22.45 \times x^{0.3} - 13.83$; $R^2 = 0.99$; root mean square = 0.78] of the significant correlations (six cycles over the increase of 800 neurons) suggests that as few as 64 neurons and three to four gamma cycles (75–100 ms) were sufficient to retain the information with $r > 0.9$ (Figure 4A); consistent with the reaction time across species.

Second, we systematically varied the temporal resolution $\Delta\tau$ between 0.1 and 2.5 ms and the phase gradient $\Delta\Phi$ between adjacent neurons from 0.012 to 5 ms. Since the reconstruction efficiency relies on the neuronal specificity recovered from the AP phases, the temporal resolution of phase, i.e., the precision of APs, is critical. This resolution determines whether neurons at the target area are able to discern the close succession of input APs and detect coincidences between APs and gamma peaks selectively. Likewise, the gamma phase gradient between adjacent neurons determines the specificity of AP-gamma coincidences with respect to the neuron's position and with respect to the wavelength of the spreading oscillation. Since $\Delta\tau$ and $\Delta\Phi$ are independent parameters of the model while both contributing to the precision of coincidences, we expected a tight relationship between them. Our simulations confirmed a monotonic relationship between temporal resolution and reconstruction efficiency from 0.1 to 1.5 ms and uncovered a surprising non-linear relationship between the gamma phase gradient and coding efficacy. At $\Delta\tau > 1.5$ ms no precise



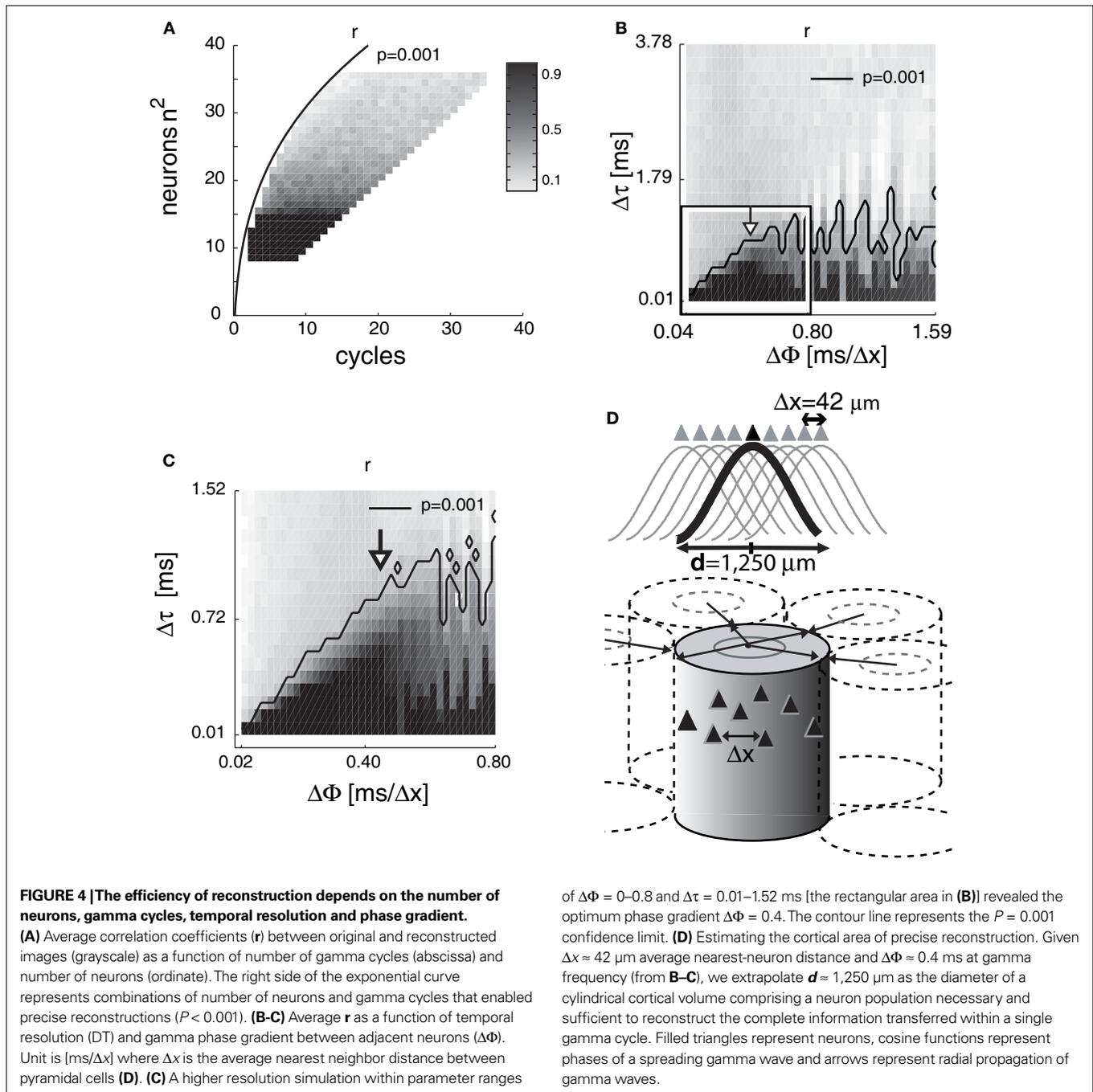
reconstruction was possible, thus setting a boundary for precise information processing consistent with empirical data (Shmiel et al., 2005). The lowest temporal resolution, or the largest time window of coincidence, which allowed a near perfect reconstruction ($r > 0.9$) was < 0.72 ms, as precise as the duration of an AP.

The first peak of the corresponding gamma phase gradient was $\Delta\Phi = 0.4 \pm 0.1$ ms, followed by a series of subharmonics from 0.6 to 1.59 ms (Figures 4B,C). Given that the average pyramidal cell density in layer 4 of the cerebral cortex is near 2.4×10^4 (Holmgren et al., 2003) and 2.5×10^4 cells/mm³ (Peters and Yilmaz, 1993),



the average nearest cell-to-cell distance derived by Monte Carlo simulations was $\Delta x = 42 \mu\text{m}$ (for details on computing neuron-to-neuron distance, see Supplementary Material), and that the optimal phase gradient was 0.4 ± 0.1 ms between adjacent neurons,

we estimated the speed of SMO propagation (v) and the diameter (d) of the cortical cylinder that would be covered by a radial wave of a single SMO cycle of f frequency and v speed. Assuming that $f_{\text{gamma}} \approx 40$ Hz and $T_{\pi} \approx 12.5$ ms, and applying $v = \Delta x / \Delta \Phi$ we



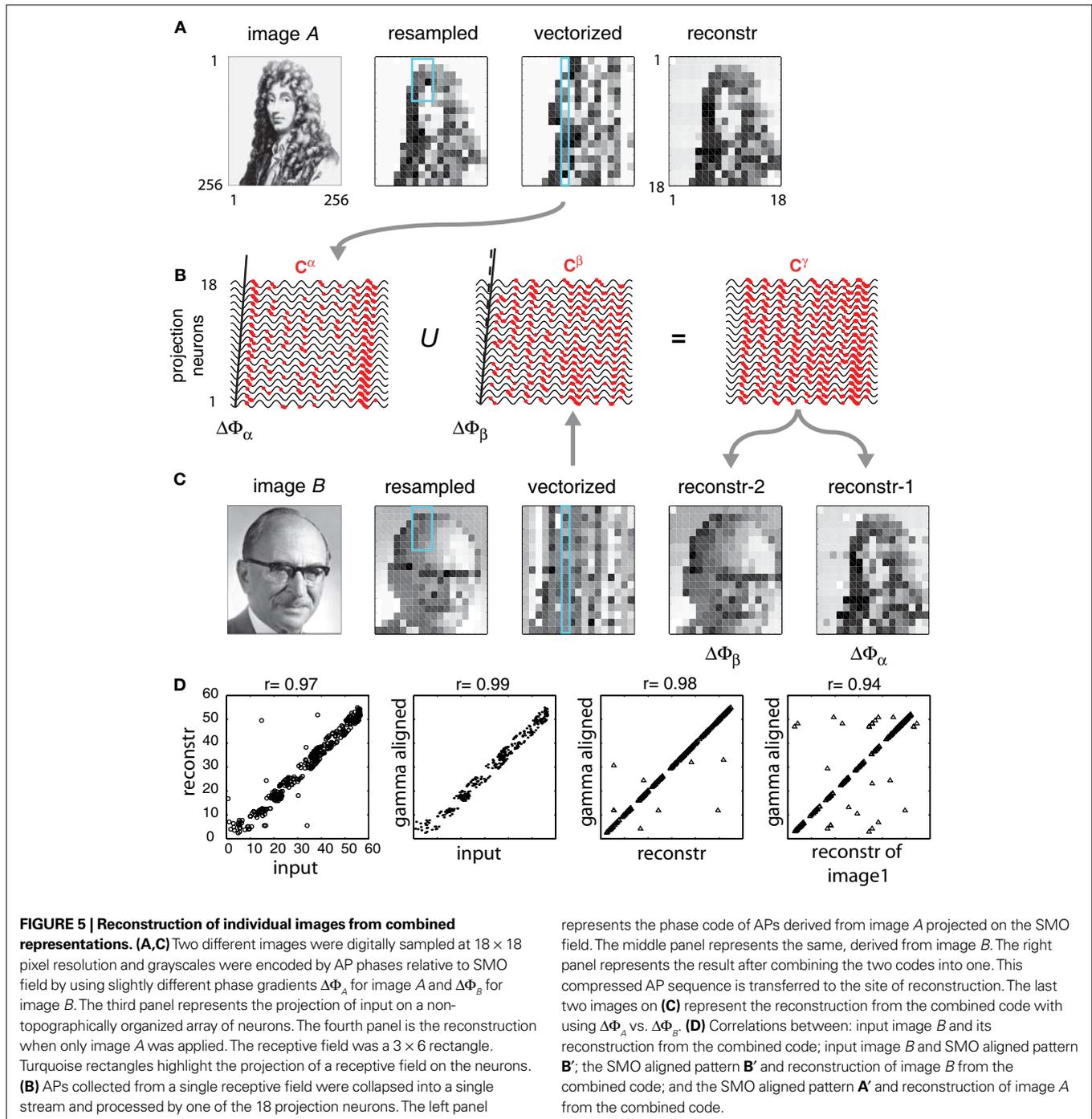
calculated $v = 0.1$ mm/ms as the speed of propagation, which is consistent with the 0.09–0.4 mm/ms expansion speeds of SMO measured from various species, cortical and allocortical areas (Benucci et al., 2007; Bringuier et al., 1999; Grinvald et al., 1994; Prechtl et al., 2000). To estimate d , we extrapolated how far a single radial oscillation period (T) would propagate as $\lambda = T \times v$. Given that $T = f^{-1} \approx 25$ ms and $v \approx 0.1$ mm/ms the estimated diameter range of integrative cortical units are $d = \lambda/2 \approx 1,250 \mu\text{m}$ (Figure 4D), consistent with the 900–1,000 μm center-to-center distance of iso-orientation columns in the cat V1 (Lowel et al., 1988; Peters and Payne, 1993). Considering that the reconstruction

efficiency is near perfect at as low as $\Delta\Phi = 0.1$ ms and does not improve beyond $\Delta\Phi = 0.8$ ms, a broader range of spatial domains ($578 \mu\text{m} < d < 4.6$ mm) could satisfy gamma frequency integration. This range is consistent with the 590 μm center-to-center distance of cytochrome oxidase blobs (Murphy et al., 1998), the $430 \pm 139 \mu\text{m}$ center-to-center distance of cortical biocytin patches in primate V1 and 600 μm in cat V1 (Lund et al., 1993) and 1.5–2.7 mm, the space constant ($d = 2 \times$ space constant) of the radial spread of stimulus induced transmembrane voltage change (Grinvald et al., 1994). The estimated d was also consistent with the reported range of cortical columns 600–900 μm (Jones, 2000) and

350–400 μm (Favorov et al., 1987), and the 400–500 μm projection field of layer 2/3 and layer 4 neurons in the rat barrel cortex (Lubke et al., 2003). Notably, the relationship between oscillation λ and column diameter is further supported by the observation that the dominant LFP in the primate motor cortex (area 4) is 20–25 Hz (Rubino et al., 2006), half of that of V1, while the center-to-center distance of cortical biocytin patches is twice of that of V1 (Lund et al., 1993). In summary, the operative ranges of model parameters satisfy the dynamic and structural constraints of the brain.

INFORMATION RECONSTRUCTION FROM COMBINED CODES

We thought it would substantially increase the performance of phase coding if neuronal architectures were able to combine codes of independent inputs into one compressed stream of APs and selectively reconstruct these representations at the target location by the same pool of neurons. Therefore, we asked whether the interference operator (Eq. 9) is able to recover multiple representations from a superimposed phase code of different input patterns. We used two different images, image S^α and image S^β and two slightly



different field gradients, $\Delta\Phi_\alpha$ and $\Delta\Phi_\beta$ (Figures 5A–C). First, processing image S^α generated the A^α latency code, transformed into A^α by gamma alignment to a field with gradient $\Delta\Phi_\alpha$. The pattern A^α was collapsed into C^α and reconstructed S'^α at the target using $\Delta\Phi_\alpha$ (Figure 5A). Likewise, we processed image S^β with the field gradient $\Delta\Phi_\beta$ and computed A^β and C^β (Figures 5B,C). Next, before the reconstruction, we combined the two compressed codes $C^\alpha \cup C^\beta = C^\gamma$, so that APs encoding image S^β were interleaved with APs encoding image S^α (Eq. 10). The combined code included both sequences of APs, derived from two different input images, without labeling them according to the neuron of origin (Figure 5B). Next, we transferred the combined code C^γ to the target, where the postsynaptic neurons successfully recovered image S'^α and image S'^β from the mixed code by applying field gradients $\Delta\Phi_\alpha$ and $\Delta\Phi_\beta$, respectively (Eqs. 11 and 12, Figure 5C). The reconstruction efficiency, quantified as the correlation between the gamma-aligned input and the recovered image, was near perfect ($r' > 0.9$, $P < 0.001$) when compared with the original input (Figure 5D). Since the variance of the correlation between the original and the gamma-aligned input was the same as the variance of the correlation between the original and reconstructed image, most of the differences between the original and reconstructed images were attributed to the gamma alignment rather than to the reconstruction.

The reconstruction from the combined phase code has intrinsic limitations. If the difference between the oscillation field gradients is small, the delay by which the two oscillation peaks reach adjacent neurons may be smaller than $\Delta\tau$, causing the neurons to confuse the input APs originating from different images. Spurious phase coincidences of the two fields may also occur if the two field gradients are very different. In order to avoid this, for a given phase gradients $\Delta\Phi_\alpha$ the second phase gradient $\Delta\Phi_\beta$ should be less than $\Delta\Phi_\alpha/(n \times \Delta\tau)$. Since n is relatively small (<9), especially when the architecture is partitioned into small receptive fields, this condition is easy to meet and does not compromise reconstruction.

Storing and retrieving multiple representations from the same pool of neurons by cueing them with slightly different SMO oscillations can be utilized as a very efficient mechanism for storing long-term memories. For the sake of algorithmic clarity, we postulated the independence of APs and SMO. However, in living tissue, large synchronized EPSPs and IPSPs may reset the field of SMOs, thus allowing selective cueing of information.

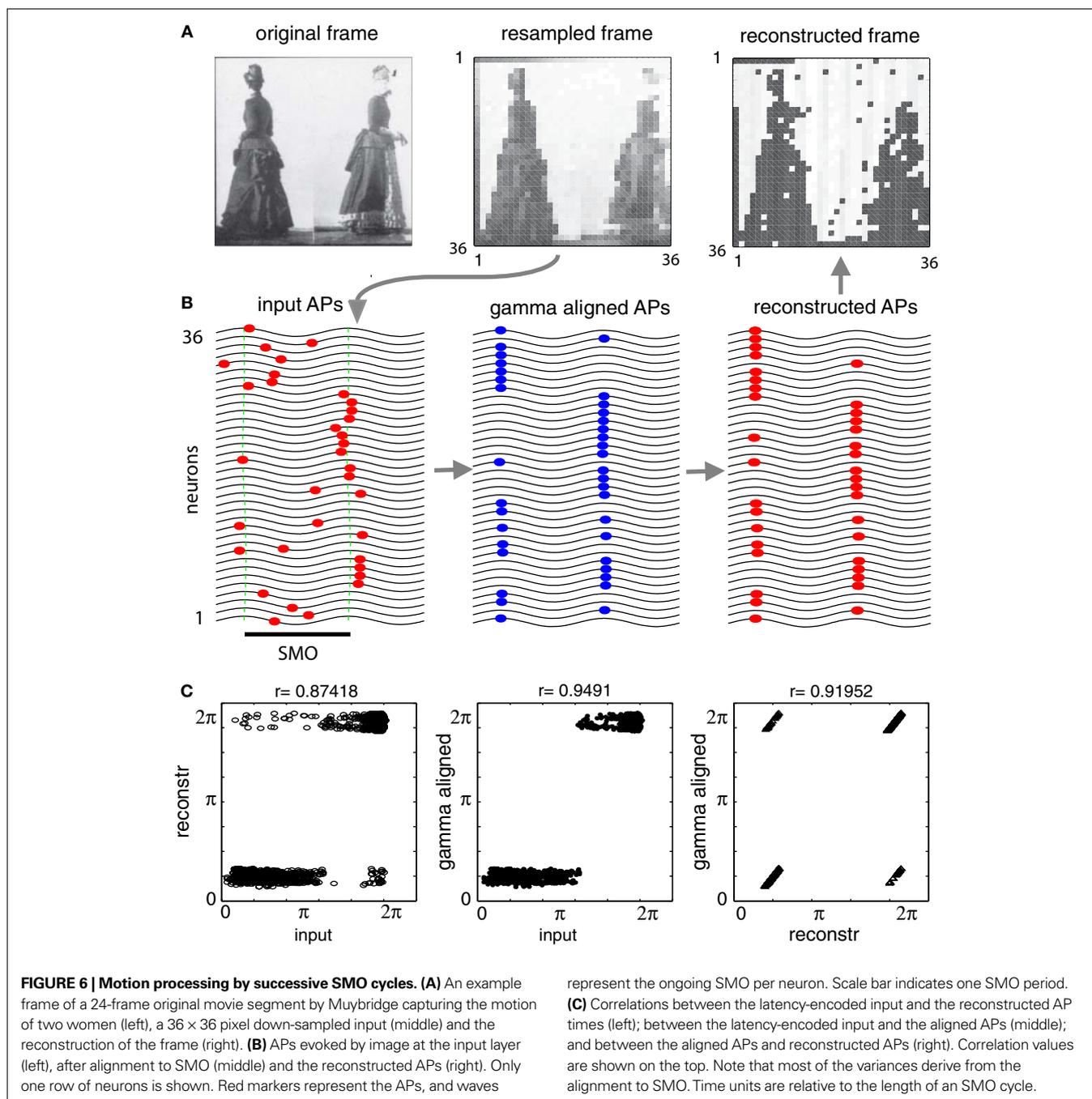
REAL-TIME MOTION PROCESSING WITH PHASE

Under natural conditions, images on the retina change rapidly either due to objects and the observer moving relative to each other or due to the observer's eye movements. A sequence of images, when presented with a frame rate $\geq 16 \text{ frame} \times \text{s}^{-1}$, is perceived as motion, (Wertheimer, 1912) and motion percept does not improve beyond $48 \text{ frame} \times \text{s}^{-1}$. This implies that under specific condition, single gamma cycles should be sufficient to process information from individual frames to generate a motion percept. It also implies that the human visual system does not benefit from a frame rate of higher than one frame per gamma cycle. We asked how phase coding would implement compression and reconstruction of a complete frame within a single gamma period with a rate of $40 \text{ frame} \times \text{s}^{-1}$, in order to remain consistent with human motion perception. According to our simulations, a single SMO cycle is capable of capturing

and retaining significant quality of the original image ($r = 0.87$, $P < 0.01$) and nearly complete representation of the gamma-aligned image ($r = 0.92$, Figure 6). In this simulation we used a 36×36 neuron architecture with 6×6 receptive fields converging to 36 projection neurons and reconstructed on a 36×36 neuron layer at the destination. For the SMO field we used a predefined set of 24 phase gradients. The SMO field gradients were generated by three randomly dispersed gamma sources slowly drifting frame-by-frame with Brownian motion. The input was a 24-frame segment extracted from an Eadweard Muybridge movie clip. Input frames were presented iteratively and each reconstruction was computed independently from the previous frame. The $\Delta\tau$ was 0.001, and a single gamma oscillation period was used for encoding. Because a faithful reconstruction of grayscale image requires $n > 4$ oscillation periods (Figure 4A), phase coding within a single oscillation period allowed to retain only the binary values per pixel (black-and-white) with no grayscale qualities processed (Figure 6A). Despite the black-and-white representation, the spatial reconstruction of the original frames was relatively good ($r = 0.874$, Figure 6C), which confirmed that sampling continuous motion by single SMO cycles is necessary and sufficient to encode a motion sequence. Notably, most differences between the original and the reconstructed image frames derived from the AP alignment (Figure 6C). The limited capacity to process textural details and motion at the same time is consistent with the anatomical and functional segregation of fast achromatic magno-cellular motion pathway and slow chromatic parvo-cellular pathway (Conley and Fitzpatrick, 1989; Fitzpatrick et al., 1985; Michael, 1988; Perry et al., 1984).

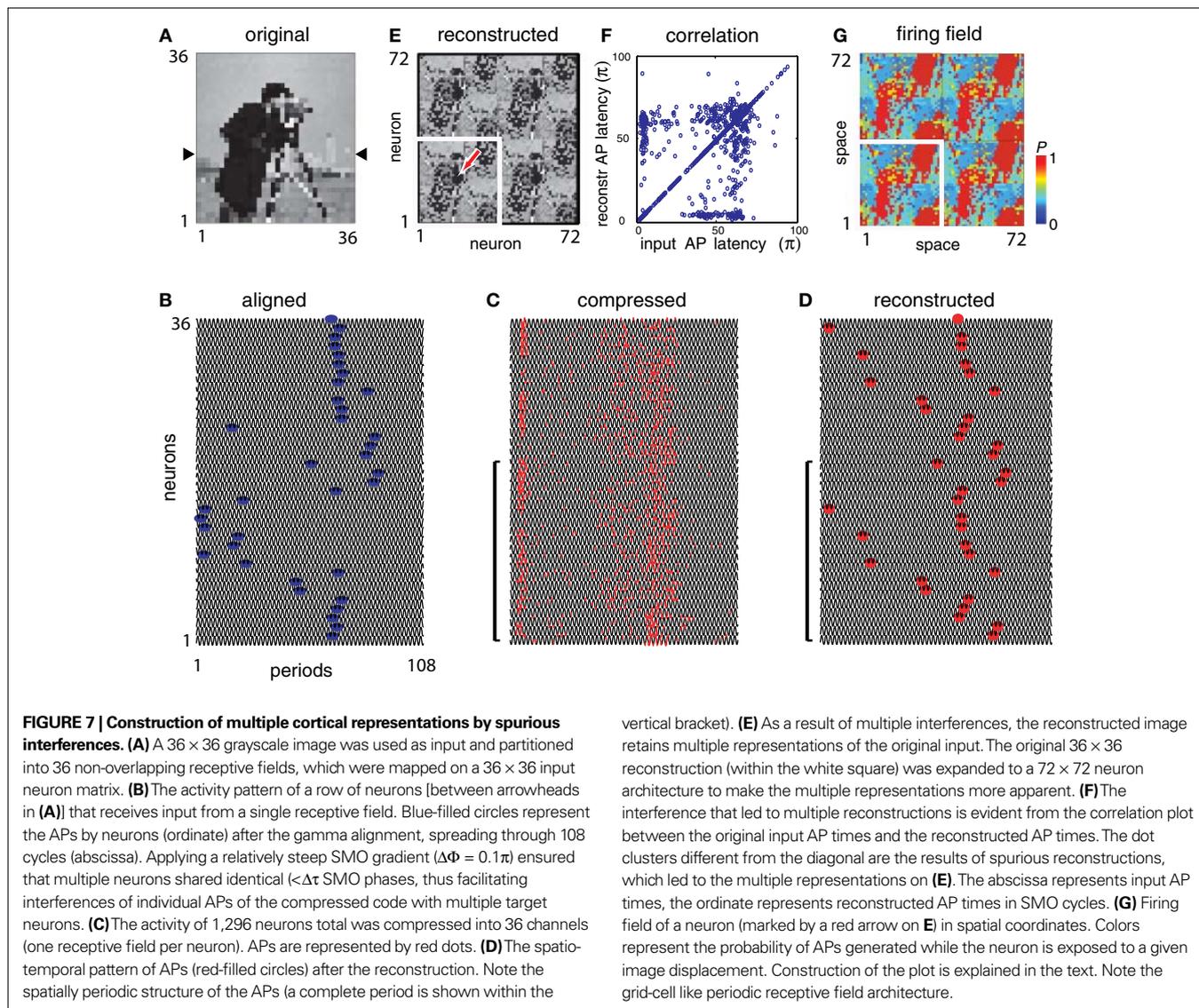
PHASE CODE GENERATES GRID REPRESENTATIONS

Since the discrimination time window is limited to one SMO cycle, the largest discriminative power is comprised in an area corresponding to λ (the distance that SMO travels in one oscillation). Beyond that diameter, representations likely repeat because spurious interferences will activate neurons that share the same SMO phase as they are multiples of λ apart. If the same input pixel is reconstructed by multiple neurons, separated by λ distances, as our model predicts, then it is necessary that different inputs are reconstructed by the same neuron, assuming finite input and neuron spaces. Consequently, individual neurons must endow multiple receptive fields, just like grid cells do in the medial entorhinal cortex (MEC) of rodents (Hafting et al., 2005). To provoke multiple cortical representations by spurious interferences, we performed pattern reconstructions on an image, tracking a prolonged interval of 144 gamma cycles with increased phase gradient ($\Delta\Phi = 0.1$) (Figure 7). The reconstruction, as predicted, yielded multiple representations of the original image (Figure 7E). This was due to systematic reconstruction errors caused by spurious interferences that occurred at neurons near λ distances apart within the neuronal matrix. The confusion of AP times across neurons of the output layer is clearly seen as clusters offset of the identity line of the input–output AP latency correlogram (Figure 7F). The consequence of these errors is twofold: (i) they generate multiple cortical representations of the same spatial location, as seen in Figure 7E; and (ii) these errors must make a given output layer neuron active not only when the input image is at the original position, but also a number of other positions that are multiple



of a constant distance away from the original position. In order to test this hypothesis, we kept track of the activity of an arbitrarily selected output-layer neuron (red arrow in **Figure 7E**) while circular shifting the original image matrix in 36×36 different positions. If the image on **Figure 7A** represents a square-shaped spatial environment from the rat's point of view, then shifting the image emulates an exhaustive exploration of this environment in a rat-centered coordinate system. For example, moving the image by 5 pixels north and 6 pixels east is equivalent to the rat moving 5 units south and 6 units west, where the unit is proportional to the rat's size. Next we plotted the firing probability of this

neuron according to the positions of the rat in the environment (**Figure 7G**). Confirming the prediction (ii), the firing probability revealed a multifold periodic receptive-field architecture, where receptive fields were constant distance apart, reproducing false-colored copies of the original image, evenly tessellating the space. This architecture is consistent with the periodic architecture of the spatial firing fields of MEC neurons (Hafting et al., 2005). Moreover, the model provides a physiological meaning to the constant spatial separation of the observed firing fields by relating it to the λ of SMO within the MEC. In addition to grid cells, according to (i), the model predicts that the actual spatial location



of the rat activates not one but multiple grid cells with shared firing fields, and these neurons must be anatomically arranged in a grid-like periodic architecture over the entorhinal cortical sheet (Figure 7E) that mirrors the periodic receptive field architecture of individual grid cells. Consistent with the proposed relationship between the grid cell geometry and phase coding, the frequency of SMO in MEC neurons, and thus λ , has been reported to scale with grid cell field spacing (Giocomo et al., 2007). We remark that the periodic structure of reconstruction necessarily occurs in 1-D but that the exact geometry in 2-D depends on the location of neurons where spurious interferences occur. Most likely, the spurious interferences generate confusions of different degrees. Neurons that are integer cycles apart share nearly identical SMO phases. Consequently, when APs are reconstructed from phases, these neurons generate confusions, which manifest in reproduction of the original image in nearly perfect copies. These neurons represent the main-grid points of the output representation (seen as the multiple images of the photographer in Figures 7E,G).

Other neurons generate only partial confusion. These neurons are usually half the distance of the main-grid points and they partially reproduce the original input (seen as phantom images of the photographer in the upper right and lower right corners of the area within the white frame in Figure 7E). Because the partial and complete confusion grids have the same period but with a half-period offset between them in both X and Y coordinates, the superposition of the two grid structures constitutes an elongated hexagonal grid topography (Figures 7E,G). Since there are also other ways of achieving a hexagonal structure, this may or may not be the mechanism by which the stereotypical grid cells in the MEC acquire precisely such geometry.

PHASE CODE EXHIBITS AP PHASE PRECESSION

Among the most prevalent evidence for the behavioral significance of systematic phase variation is the phase precession of hippocampal pyramidal cell firing relative to the local theta oscillation while a rat traverses through the place field of the neuron

(O'Keefe and Recce, 1993). These pyramidal neurons also exhibit theta frequency SMO (Leung and Yim, 1986). Since our model requires nothing more than that AP timing be dependent on a coherent oscillatory drive (SMO) that propagates in space, we investigated whether phase precession can occur without further assumptions. We implemented the spatial navigation condition by defining a 16×16 matrix of topographically arranged place cells where neuron a_{ij} at the i th column and j th row was driven by the input corresponding to the rat being at the location defined by its X_i and Y_j coordinates. Thus, the place field of the rat when moving through $X_i Y_j$ was modeled by neuron a_{ij} being exposed to a transient stimulus. Theoretically, this is equivalent to a transient stimulus, defined as a 2-D Gaussian, moving through the receptive field of a neuron (Figure 8A). This is similar to the real-time motion processing example in Figure 6, except that the SMO here represents theta instead of gamma. The complete path of the rat was modeled by a series of 20 frames consisting of 16×16 -pixel-resolution snapshots, in which the 2-D Gaussian advanced one pixel per frame. Since the diameter of the Gaussian was larger than the place field of a neuron, the rat's position evoked activity in adjacent place cells proportional to the amplitude of the Gaussian intersecting with their place fields. The square area outlined in Figure 8A covers the place fields of nine neurons. Multiple neurons were activated when the rat was crossing this area. The phase-coded APs of these neurons were combined into a compressed code and shared among the nine neurons before reconstruction. When we tracked the activity of these neurons while the location of the rat changed across frames, the compressed AP sequences displayed a systematic phase precession as a function of the distance from the center of the Gaussian, consistent with the empirical phase precession of hippocampal end entorhinal cortical neurons (Hafting et al., 2008; O'Keefe and Recce, 1993). We controlled only one parameter in these simulations, the phase gradient of the SMO field. Figures 8B–E shows that the direction of phase precession depended on the geometry of the SMO field applied to both input encoding and reconstruction. When unidirectional propagating waves (traveling waves) were applied, depending on the direction of propagation relative to the activation sequence of place cells, we observed different monotonic phase precession effects. Wave propagation in the direction opposite of the place cell activation sequence enabled progressive phase advancement, while wave propagation in the direction of place cell activation sequence enabled progressive phase delay (Figures 8B,D, respectively). By reducing the SMO phase gradient ten times decreased the slope of phase precession 10 times from 280° to 28° (Figures 8B,C). Furthermore, when a radial propagation of SMO field was applied (from a single or from multiple sources) the precession started with increasing phase advancement until the animal, i.e., the center of the Gaussian, reached the center of the neuron's place field, followed by a progressive phase delay as the animal was leaving the center of place field (Figure 8E). The dependency of phase precession angle on the SMO gradient relative to the activation sequence of the place cells suggests that hippocampal pyramidal neurons may be able to "read out" the rat's heading direction from the phase precession, assuming that the local SMO field gradient does not change rapidly over time (Huxter et al., 2008). In summary, phase precession derives naturally from phase coding,

without making any additional assumptions, such as asymmetric synaptic potentiation (Mehta et al., 2000) or dual-oscillator drive (Blair et al., 2008; Burgess, 2008; Lengyel et al., 2003). Phase precession is controlled by a single parameter, the SMO field gradient. Furthermore, consistent with experimental data, phase precession in our model does not require multiple trials to develop but it does require SMO (Hafting et al., 2008). Since object locations in the environment, according to this model, are originally encoded by phase, we propose that phase precession (phase advancement and phase delay) is expressed in all cortical and limbic structures, including the MEC (Hafting et al., 2008) and primary sensory areas of the neocortex where the spatial representations are continuously updated due to behavior.

MODELING STIMULUS INVARIANTS

In order to represent stable objects in a dynamic world, neurons must encode object features invariantly with respect to a few common transformations such as position, rotation and time. In order to address whether or not phase coding is able to retain some of these features in spite of spatial or temporal discontinuities, we modeled four different types of invariant stimulus conditions: (i) space-time invariant 1-D stimulus, (ii) 2-D stimulus embedded in random background noise, (iii) 2-D transposition-invariant stimulus and (iv) character recognition. We classified these invariants with respect to either temporal or spatial transformations. Among the invariants we tested, (i) and (ii) were time invariant, while (iii) and (iv) were invariant to spatial transformations. (Implementations of the ii, iii and iv types of invariant preservations are described in the Supplementary Material.)

For (i) we constructed a 1-D random vector \mathbf{R} containing 32 values $[\mathbf{r}_1, \dots, \mathbf{r}_{32}]$. Next we defined "frames" as the possible subsets of this vector containing 16 adjacent values. We then began an iterative cycle, where first we inputted the first frame $[\mathbf{r}_1, \dots, \mathbf{r}_{16}]$ to a network of 16 neurons $[\mathbf{s}_1, \dots, \mathbf{s}_{16}]$ and performed all four phases of information processing from encoding to reconstruction. During each of the next 16 cycles, we incremented the frame position by one step $[\mathbf{s}_1, \dots, \mathbf{s}_{16}] = [\mathbf{r}_p, \dots, \mathbf{r}_{p+15}]$, where $i = \{2, \dots, 17\}$, and performed all four phases of information processing. Thus in each iteration, 15 out of 16 frame values remained identical to the values in the previous frame, but each of the 15 values was projected onto a different neuron at each exposure (Figure 9A). Presenting the neurons with only part of the complete pattern in each cycle, such that each neuron's input value changes every cycle, allowed us to address how a constant (time-invariant) spatial pattern is represented by the phase code in the array of neurons over time (Figures 9B–E). When we superimposed the temporal patterns of APs of the compressed codes (transfer stage) from all of the 17 successive trials, we observed a marked conservation of AP latencies along with sudden transitions of gamma re-alignments (Figure 9D). The conservation of time-invariant patterns is reflected by the temporal consistency of APs over time. This consistency was quantified by correlating the AP times (wrapped around 3π) from one frame with the AP times of the next frame and comparing it to the correlation obtained when the stimulus frames were uncorrelated random patterns (i.e., when there was no overlap between successive frames). The return plots in Figure 9F summarize these correlations. As seen on the left return plot,

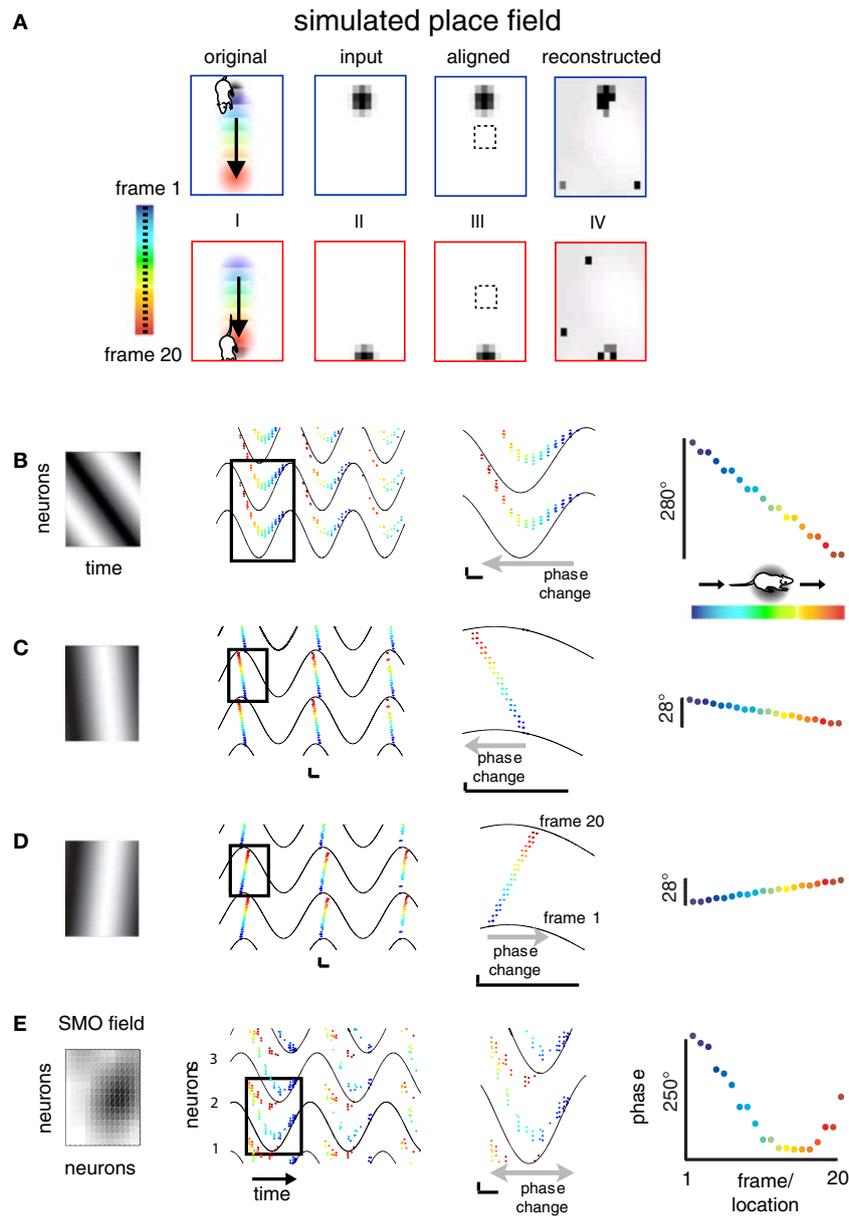
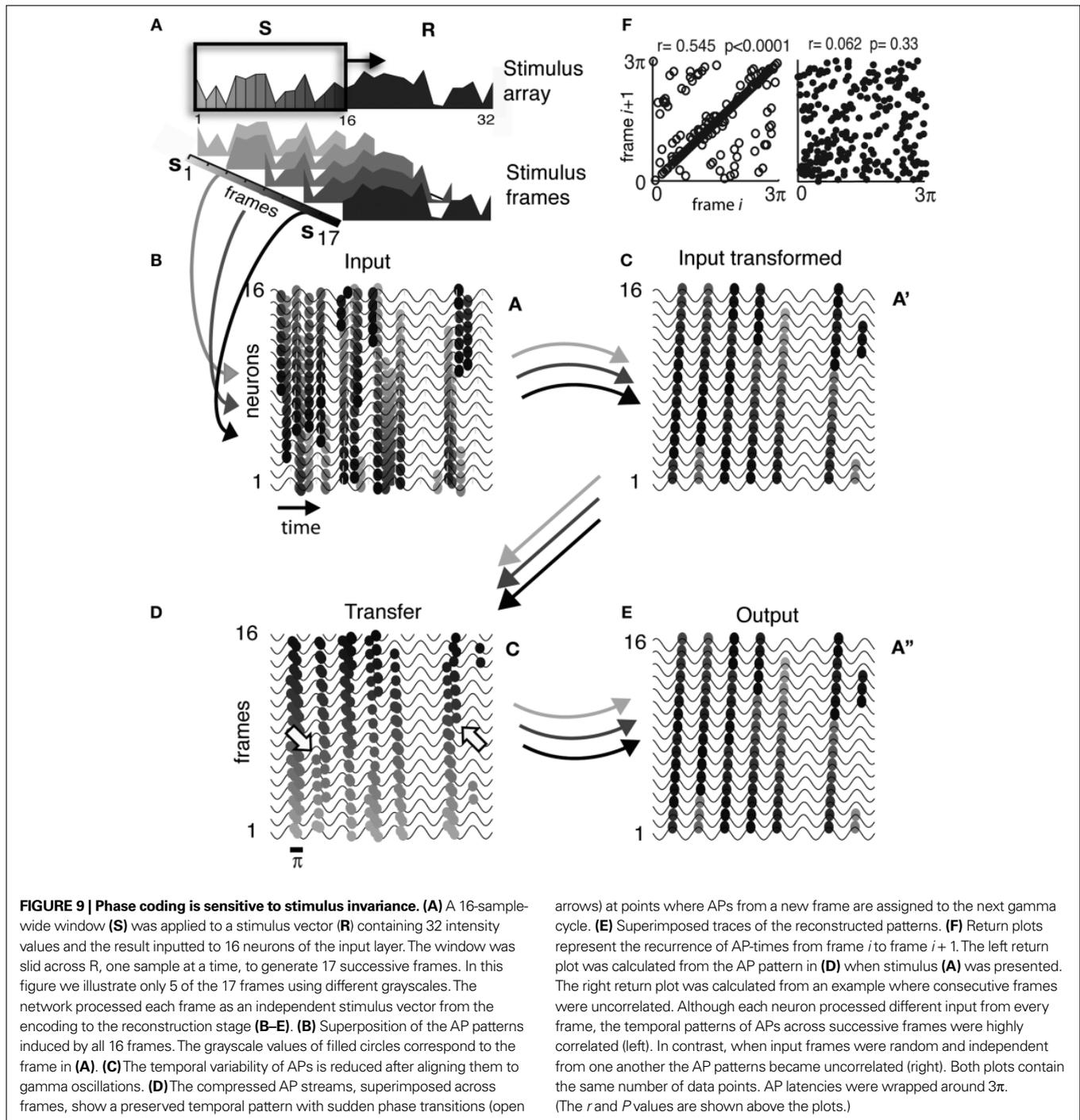


FIGURE 8 | Phase code exhibits phase precession. (A) The input to hippocampal place cells was modeled by generating a 20-frame translation of a 2-D Gaussian representing the rat's position (rendered with colors) relative to the environment (rectangle). The area was evenly tiled with place fields individually mapped on a 16×16 grid of input layer neurons. The size of the Gaussian was chosen such that 9–12 neurons were simultaneously exposed to the rat's location, but the composition of neurons driven by a given input frame changed with the rat's location. The four blue and red framed panels represent the four processing stages of the initial and terminal frames, respectively: (I) the rat's location, represented by a 2-D Gaussian, (II) the discrete sampling of the input by the grid of neurons (input), (III) the input phases in grayscale after APs are aligned to the SMO, and (IV) the reconstructed activity pattern of neurons encoding the rat's own location. The outlier pixels in both frames are reconstruction errors. **(B–E)** Effects of SMO field topography on phase precession. Left panels represent the SMO (**(B–D)** in time-neuron and **(E)** in neuron space). Second and third panels represent snapshots of the activity of three projection neurons at different magnifications. These neurons in the model represent hippocampal pyramidal cells with adjacent place fields within the

outlined area [dashed rectangle on **(A)**]. The APs evoked by consecutive frames are projected between the SMO waves of adjacent neurons and color-coded according to the input frame in **(A)** corresponding to the rat's location. The abscissa represents the time of the APs. The fourth panels represent the phase of APs during a single run as a function of the rat's location. **(B)** When a traveling wave SMO field gradient was applied the neurons expressed a monotonically advancing AP phase precession spanning 280° theta phase angle. **(C)** By reducing the phase gradient of SMO ten times the AP phase precession angle decreased proportionally to 28° . In both examples **(B,C)** the traveling direction of SMO was set to be the opposite to the place cell activation sequence. **(D)** We reversed the direction of phase precession relative to **(B,C)** from monotonically increasing advancements to monotonically increasing delays by changing the SMO phase gradient such that it traveled in the direction of the activation sequence of neurons. **(E)** A bidirectional AP phase precession was enabled by a radially propagating SMO field, where the AP phases advanced continuously until the animal reached the place field center [outlined area in **(A)**] and developed an increasing delay as the animal exited. Phase scale bars represent $\frac{1}{4} \pi$. Arrows signify the directions of phase precessions.



invariant stimulus frames evoke AP patterns in the compressed code that are highly correlated across frames, despite phase transitions and despite the fact that each neuron receives a different input from each frame. In contrast, uncorrelated frames evoke APs that are uncorrelated across successive frames. This conservation of the AP pattern for an invariant stimulus has implications for visual information processing. For example, we predict that V1 projecting LGN neurons should generate self-similar and stimulus-dependent AP sequences for slowly moving images. These AP sequences are

expected to be invariant of position shifts of the whole image on the retina within a certain range of eye movements.

DISCUSSION

We presented theoretical support for phase coding in the CNS with the following two questions in focus: How can the precise spatio-temporal structure of a stimulus be encoded by the phase of APs, and how then can it be recovered from the phase of APs? The model outlined above interprets phase locking as a mechanism by which

the phase of APs encodes the precise spatiotemporal structure of the visual input using gamma oscillations as a reference. Utilizing the phase the brain is able to keep different stimulus features separated within the same code (impossible with rate coding) until these features are separated by individual neurons upstream, consistent with sparse coding (Figure 5). At the same time the phase code allows simultaneous information to be bound together by referencing them to the same gamma cycle, which allows a smooth tracking of sensory input, such as that illustrated on the example of motion processing at a physiological rate (Figure 6). Aside from the capability of modeling thalamo-cortical and cortico-cortical information processing, phase coding demonstrates a remarkable flexibility in reproducing known functional features of the allocortex, such as the grid cells of the entorhinal cortex and the phase precession of hippocampal pyramidal cells (Figures 8 and 9, respectively). Moreover, phase coding allows the extraction of stimulus features that are invariant with respect to spatial and temporal transformations (Figure 9 and Figure 2 in Supplementary Material).

Remarkably, phase coding does not contradict with rate coding. The two encoding schemes are compatible (Ahissar et al., 2000), they may coexist (O'Keefe and Burgess, 2005) and/or may complement each other (Gollisch and Meister, 2008; Kayser et al., 2009) simply because the likelihood of coincidences between the presynaptic AP and postsynaptic SMO scales with the frequency of APs. Moreover, modulation of firing rate is concomitant with changing the phase structure of APs (Margrie and Schaefer, 2003). Immediate predictions made based on phase coding are summarized in Table 1.

Nevertheless, the robustness of the model depends on a number of limiting factors, which remain to be investigated. One is the dependency of phase coding on the need for compression before transmission. A key assumption we made for the sake of theoretical clarity in the 1-D model was the n -to-1 dimensionality reduction

from the n input layer neurons to a single channel and a 1-to- n dimensionality expansion by information reconstruction at the target layer. This extreme information compression was feasible within the parameter space specified in Figure 4 (Section "Biological Significance of Model Parameters"). However, considering parameters closer to the physiological condition, such as a large n with intrinsic noise in membrane potential affecting AP generation, combined with certain variance of oscillations, and heterogeneous axonal composition within the projection pathways, the n -to-1 reduction is neither necessary nor feasible. This has two consequences. First, a substantial saving in axonal volume can still be achieved with an n -to- m mapping, where $n > m \geq 1$, which would not be possible without phase coding. Second, by clustering neurons into m groups, where instead of n a smaller $n' = n/m$ neurons converges on a single channel, the need for m projection neurons would still substantially improve the reconstruction efficacy, consistent with the known convergence of sensory input on single neurons that constitutes for receptive fields (Figure 3).

Another limiting factor is the precision of neural code. The phase coding model requires neurons to resolve ~ 1.5 ms coincidences between input and intrinsic SMO under the assumption that SMO is 40 Hz. Whether or not the CNS is able to maintain this high precision has been argued (Shadlen and Newsome, 1998; Shmiel et al., 2005). Since the ~ 1.5 ms precision would guarantee that neighboring neurons do not confuse the origin of two input APs coinciding with their slightly different SMO, this precision is most likely to be verified in the cross-correlograms of nearby neurons. Another factor of precision is the frequency of SMO. In structures where the predominant SMO has a less than 40 Hz frequency, such as the 5–7 Hz in the entorhinal cortex (Giocomo et al., 2007), the required precision is proportionally less, only $\Delta t < 12$ ms. Thus ~ 1.5 ms precision is the most extreme requirement for precision.

Table 1 | Predictions based on the phase coding model. Phase coding, a common oscillatory reference with a field gradient, and the AP-SMO coincidence are the three cornerstones of the model. They are also highly interdependent, which enables to make these predictions.

#	Predictions
1	Gamma LFP must show a phase gradient ($v = -0.1$ mm/ms) between adjacent neurons within a cortical column. This can be conceived as propagating waves or radial spread of gamma activity from multiple generator foci.
2	The phase gradient is dependent on the wavelength of characteristic LFP at a given region, and this wavelength is proportional to the average columnar center-to-center distance (Giocomo et al., 2007; Lund et al., 1993). Thus, the radius of cortical columns must scale with the wavelength of dominant LFP oscillations. A supportive evidence is that grid cell spacing has been reported to scale with the theta frequency SMO in layer-II entorhinal cortical neurons (Giocomo et al., 2007).
3	Among simultaneously recorded neurons, those that are exposed to their preferred stimulus should exhibit a more precise phase lock of APs with the local gamma recorded from the same electrode than neurons exposed to a suboptimal stimulus.
4	The precision of AP to AP cross-correlation between excitatory neurons in the superficial cortical layers should be an order of magnitude higher (~ 1.5 ms) than the precision of the autocorrelations (15–25 ms).
5	High magnitude of zero time-lag cross-correlations should be most expressed between layer-4 granule cells of the neocortex because they are the recipients of the compressed code. In contrast, supra granular layer pyramidal neurons should generate much smaller (but more precise; see Section "Discussion") cross-correlations because their outputs represent the reconstructed sparse code. The average firing rate of these neurons is expected to be also lower than that of granule cells in layer-4.
6	Intervention of SMO propagation is expected to disrupt information reconstruction, thus altering perception. Similarly, disruption of spike timing relative to the gamma oscillation must perturb perception.
7	According to our simulations, an interval of four gamma cycles is sufficient to retain more than 90% of the information, thus 100 ms processing time after the first APs reaching V1 (~ 60 ms) should be sufficient to make complex perceptual decisions (Thorpe et al., 1996).

The third limiting factor is the coherency of oscillations between the source and target area. Phase coding is extremely sensitive for the degree of coherence between source and target. A number of physiological mechanisms have been suggested to maintain high cross-structural coherence, such as the thalamo-cortical loop (Llinas and Ribary, 1993; Steriade et al., 1996), entrainment of cortical gamma rhythms by hippocampal theta (Sirota et al., 2008), cortical theta reset (Rizzuto et al., 2003), and gap-junctions (Traub et al., 1996), but direct evidence for an overall synchronizing mechanism that would ensure precise coherence between sensory structures and corresponding cortical areas is lacking.

The fourth limiting factor is the noise tolerance of phase coding. We argued that the compression of the phase code before transference through the projection pathways improves the noise tolerance, and we suggested that compression combined with a feed-forward parallel distribution would improve reliability by implementing an error correction mechanism.

The last limiting factor is the lack of direct evidence for phase-shifted fields of SMO. This pre-assumption was made based on combining two sources of evidence. One is the direct observation of propagating membrane potential changes by intrinsic optical signal and voltage-dependent Ca²⁺ imaging (Benucci et al., 2007; Bringuier et al., 1999; Grinvald et al., 1994; Prechtl et al., 2000) and propagating LFP waves (Prechtl et al., 2000; Rubino et al., 2006). The other source of evidence derives from direct intracellular observations of single neuronal oscillations, such as subthreshold membrane oscillations (Giocomo et al., 2007; Hutcheon and Yarom, 2000; Llinas et al., 1991). Based on the high correlation between LFP and SMO of individual neurons, the conjecture of phase-shifted field of SMO is plausible.

In conclusion, we demonstrated through various examples that the benefit of encoding information by the phase of APs relative to the intrinsic neuronal oscillations is multifold:

- (i) Allows different types of sensory data to be converted to a common coding scheme in which information is referenced by the field of local oscillations and decoded independent of the neuronal distance from the original location and time in the brain, given that the reference

oscillation field has the same spectral properties as at the source.

- (ii) Protects the code against potential distortions due to the variance of conductivity during transmission in long projection pathways.
- (iii) Enables deployment of the same compressed information over large areas of the cerebral cortex and selective “reading-out” of the local information (**Figure 5**) without point-to-point mapping between the neurons of the input and target areas.
- (iv) Effectively compresses the spatio-temporal pattern of APs before transmission (n -to-1 and 1-to- n).
- (v) Enables invariant temporal relations to be encoded independent of the delay and duration of sensory information processing. Similarly, it enables invariant spatial relations to be encoded independent of the absolute spatial location of the agent (**Figure 9** and **Figure 2B** in Supplementary Material).
- (vi) It provides an intrinsic biological mechanism for stimulus segmentation.
- (vii) It enables information reconstruction at single-neuronal precision (sparse coding) without confusion of information between neighboring neurons.
- (viii) It implements spatial and temporal transformations between cortical representations (spatio-temporal receptive fields), which cannot otherwise be achieved by simply changing the topographic mapping between interconnected cortical areas (**Figure 1** in Supplementary Material).

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at <http://www.frontiersin.org/systemsneuroscience/paper/10.3389/neuro.06/006.2009/>

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