Computational Modeling of Orientation Tuning Dynamics in Monkey Primary Visual Cortex

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Abstract

In the primate visual pathway, orientation tuning of neurons is first observed in the primary visual cortex. The LGN cells that comprise the thalamic input to V1 are not orientation tuned, but some V1 neurons are quite selective. Two main classes of theoretical models have been offered to explain orientation selectivity: feedforward models, in which inputs from spatially aligned LGN cells are summed together by one cortical neuron; and feedback models, in which an initial weak orientation bias due to convergent LGN input is sharpened and amplified by intracortical feedback. Recent data on the dynamics of orientation tuning, obtained by a cross-correlation technique, may help to distinguish between these classes of models. To test this possibility, we simulated the measurement of orientation tuning dynamics on various receptive field models, including a simple Hubel-Wiesel type feedforward model: a linear spatio-temporal filter followed by an integrate-and-fire spike generator. The computational study reveals that simple feedforward models may account for some aspects of the experimental data, but fail to explain many salient features of orientation tuning dynamics in V1 cells. A simple feedback model of interacting cells is also considered. This model is successful in explaining the appearance of Mexican-hat orientation profiles, but other features of the data continue to be unexplained.

Keywords: Cortical dynamics, orientation tuning, monkey, primary visual cortex, layers.

Introduction

How visual cells in the primary visual cortex (area V1) become selective to the orientation of a visual stimulus is an important question in visual neuroscience (see, for example, Das (1996); Reid and Alonso (1996); Sompolinsky and Shapley (1997)). The question is how V1 cells become strongly tuned for orientation since it is known that cells in the thalamus are not. Hubel and Wiesel, who first discovered the property of orientation tuning in V1 cells, proposed a simple circuitry to explain orientation tuning (Hubel and Wiesel, 1962, 1968). They suggested that the inputs from LGN cells with spatially aligned receptive fields are summed together by one cortical neuron. This spatial alignment of receptive fields generates a preference for a particular orientation in the cortical cell. We refer to this, and similar proposals, as feedforward models. An alternate class of models are feedback (or recurrent) models, which suggest that the feedforward tuning is initially weak and only provides a small orientation bias to the cortical cell. This bias is then amplified and sharpened by intracortical feedback (Wörgötter et al., 1991; Douglas et al., 1995; Somers et al., 1995; Ben-Yishai et al., 1995; Maex and Orban, 1996; Carandini and Ringach, 1997). Clearly, these models are not mutually exclusive; there is a continuum of models depending on the relative strength of the feedforward and feedback connections (Ben-Yishai et al., 1995).

In this work we study recent data on the dynamics of orientation tuning obtained by a cross-correlation technique (Ringach et al., 1997b). The method used for measuring tuning dynamics is a modification of the cross-correlation method and shares many of its properties (de Boer and Kuyper, 1968; Marmarelis and Marmarelis, 1978; Jones and Palmer, 1987; Victor, 1992; Reid et al., 1997). We believe such data may help to determine where the cortical circuitry is best represented along the continuum between feedforward models and feedback models.

As shown below, there are clear differences of dynamics between cell classes in V1. The aim of this paper is to consider these qualitative features, rather than fitting models to data. To achieve this goal, we simulated the experimental protocol on various types of receptive field models, including a simple Hubel-Wiesel type feedforward model: a spatio-temporal filter followed by an integrate-and-fire spike generator. The computational study reveals that simple feedforward models may account for some aspects of the experimental data, but fail to explain many salient features, including the Mexican-hat profile of the orientation distributions and the active inhibition seen at off-optimal orientations ("cross-orientation" inhibition). We then studied a feedback network of interacting cells. A similar model is used in the earlier work of Ben-Yishai et al. (1995). We find that the model with lateral coupling better approximates the behavior of some V1 neurons. However, some aspects of the experimental data remain unexplained.

This paper is organized as follows. We first describe the experiment and the most salient features of the data. A theoretical and computational study of the feed-forward model is then presented. We then study of the behavior of a small network of interacting oriented neurons. Finally, we discuss the relevance of our findings to the modeling of orientation tuning and cortical function.

The Experiment

The visual stimuli used are made up of sinusoidal luminance gratings,

$$I(x, y; \theta, \phi) = A\,(1 + \epsilon \sin[\,\omega(x \cos\theta - y \sin\theta) - \phi\,])\,. \tag{1}$$

Here, $(x, y) \in [-1, 1] \times [-1, 1]$ represents coordinates in visual space and θ and ϕ are fixed parameters whose values range from $-90°$ to $90°$ and from $0°$ to $360°$ respectively. The angle θ corresponds to the orientation of the grating in space while ϕ represents a spatial phase relative to the origin. The variable ω represents the spatial frequency of the grating. A is the mean luminance of the grating and $0 \le \epsilon \le 1$ is its contrast. Images of uniform luminance, A, with no spatial structure ($\epsilon = 0$), will be referred to as "blanks".

The visual stimulus used in the experiments of Ringach et al. (1997a) consists of a rapid sequence of randomly selected images, each of which is either a "blank" or a grating at one of N possible angles: $\theta_1 = -90°$, $\theta_2 = -90° + \Delta\theta$, $\ldots$,$\theta_N = -90° + (N-1)\Delta\theta$, and one of M possible phases: $\phi_1 = 0°$, $\ldots$, $\phi_M = 0° + (M-1)\Delta\phi$; where $\Delta\theta = 180°/N$ and $\Delta\phi = 360°/M$. Blanks are represented by a dummy orientation with index $N+1$; $\theta_{N+1} = blank$.

A random sequence of images, $\{I_i\}$, is generated from an associated random sequence of orientation angles and phases $\{(\theta_{n(i)}, \phi_{m(i)})\}$ where $n(i)$ is chosen from $\{1, \ldots, N+1\}$ with probability $1/(N+1)$ and $m(i)$ is chosen from $\{1, \ldots, M\}$ with probability $1/M$. θ and ϕ are piecewise constant functions in time, changing every T ms (the duration of one "frame" in the movie). In the physiology experiments T was 16.6 ms, $N = 18$, and $M = 8$.

The responses of cells to the visual stimulus are recorded and analyzed using the following cross-correlation technique. First, a correlation time $\tau \ge 0$ is selected. An $(N+1) \times M$ array of counters, C, is zeroed. We read through the recorded spike train. If a spike occurred at time t we find the orientation, θ_i, and spatial phase, ϕ_j, of the grating shown at time $t - \tau$. We then increment the corresponding counter, $C(i, j; \tau)$, by one. If the image was a blank, then $i = N+1$ and j is chosen randomly from $1 \ldots M$. We apply this procedure to each spike in the spike train. In this way, all the spikes are represented in the counter array.

Averaging over spatial phases, we define $Pr(\theta_i; \tau) = \sum_j C(i, j; \tau) / \sum_{i,j} C(i, j; \tau)$. $Pr(\theta_i; \tau)$ is the relative frequency at which a grating with orientation θ_i appeared in the stimulus sequence τ ms before a spike. Because the relative frequency approximates the probability distribution, in the following we call $Pr(\theta; \tau)$ the "probability" that a grating at angle θ appeared τ ms before a spike. The dynamics of orientation tuning are studied by measuring how $Pr(\theta_i; \tau)$ evolves with the correlation time τ. Blanks provide a dynamic baseline to which the responses to gratings can be compared. In our interpretation, if $Pr(\theta_i; \tau) > Pr(blank; \tau)$, for a fixed τ, then the orientation θ_i is excitatory relative to the blank. Similarly, if $Pr(\theta_i; \tau) < Pr(blank; \tau)$ then θ_i is inhibitory.

How will $Pr(\theta_i; \tau)$ behave as a function of τ? For $\tau = 0$, for example, we do not expect to see any influence of the visual stimulus on the output train of action potentials since there is a time delay between the retina and the evoked activity of cortical cells. Therefore, $Pr(\theta; 0)$ should be a uniform (flat) distribution. Similarly, for very large correlation times τ we also expect to obtain a uniform distribution for $Pr(\theta; \tau)$; this is because cells have only finite memory. For intermediate values of τ one would expect to see a smooth peaked distribution which represents the preference of the cell for particular orientations. There will be one value of τ for which the cell will be better tuned than at any other time; we will refer to this value as the "optimal τ".

Experimental Results

In this section we describe some of the salient features of the experimental results in Ringach et al. (1997a, 1998) that we would like the mathematical models to explain. No obvious differences were found between the dynamics of simple and complex cells. Both classes of neurons exhibited the range of behaviors described below. The perspective plots in Fig. 1 illustrate typical results obtained from orientation selective cells in monkey V1 at different values of the correlation time τ, with τ given beside each graph. Each plot represents the measurements from a single cell. The graphs within the perspective plots show the distributions $Pr(\theta;\tau)$. The dashed line represents the baseline given by the probability of a blank, $Pr(blank;\tau)$. The range of values for the correlation times τ was chosen to bracket the response period for each cell. The x-axis represents the orientation θ between -90° and 90°, and the y-axis represents the value of $Pr(\theta;\tau)$. The vertical scale bar in the lower graph of each perspective plot has a height of $1/(N+1)$, which is the value for a uniform distribution on the orientation domain.

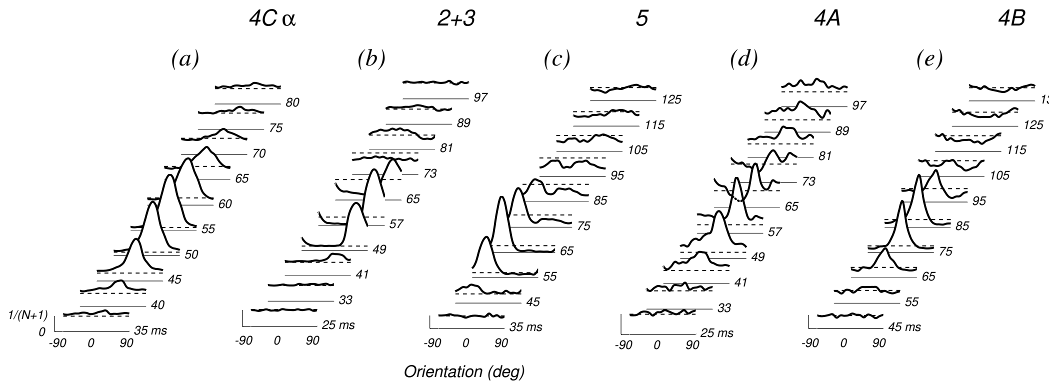

Figure 1: Examples of orientation tuning dynamics from monkey V1 cells. (a) Orientation tuning dynamics from a cell in layer 4Cα. The graphs within the perspective plots show the distributions $Pr(\theta;\tau)$ at different values of τ. The dashed line represents the baseline given by the probability of a blank, $Pr(blank;\tau)$. The x-axis represents the orientation θ between -90° and 90°, and the y-axis represents the value of $Pr(\theta;\tau)$. The vertical scale bar in the lower graph of each perspective plot has a height of $1/(N+1)$, which is the value for a uniform distribution on the orientation domain; (b) Orientation dynamics from a cell in layer 2+3 ; (c) Orientation dynamics from a cell in layer 5; (d) Orientation tuning from a cell in layer 4A. The profile at 57 ms is "transparent" with the profile at 65 msec indicated by a dotted line; (e) Orientation tuning from a cell in layer 4B.

Fig. 1(a) illustrates the response obtained from a cell located in layer $4C\alpha$, one of the main input layers of V1. For $\tau < 40$ ms and $\tau > 75$ ms the distributions are uniform. The distribution develops into a unimodal distribution and peaks at $\tau = 55$ ms. The distribution's peak occurs at the same angle as the preferred angle of the cell as found using conventional drifting grating stimuli. Notice that, in this case, the probability of the orientation orthogonal to the preferred one and the probability of a "blank" behave in a very similar way. In fact, this behavior does not hold for all layer 4C cells; there are cells for which the blanks and the orthogonal orientation do not have similar dynamics. More intricate dynamical responses are observed outside layer 4C (Ringach et al., 1997b); see Figs. 1(b–e).

Consider the responses depicted in Fig. 1(b). For $\tau < 40$ ms and $\tau > 90$ ms the distributions are uniform. The response develops and peaks at $\tau = 57$ ms. The distribution's peak occurs at the same angle as the preferred angle of the cell as found using conventional drifting grating stimuli. The distribution relaxes to a uniform profile around $\tau = 73$ ms, at which time the blank is more probable than any angle. We observe an "inversion" of the probability profile at $\tau = 81$ ms. At this time the probability of the preferred orientation of the cell is lower than that of any other angle; instead, the most probable angle $\tau = 81$ ms before a spike is the angle *orthogonal* to the preferred angle. Many cells show such "inversions", and this is one of the properties we would like the models to explain.

Another important aspect of the behavior shown in Fig. 1(b) is the dynamics of the probability of a "blank". Initially, for $\tau < 57$ ms, the probability of a blank behaves in a very similar way to the probability of the orientation orthogonal to the preferred angle. However, this is not the case for larger values of τ. For example, for 57 ms $\leq \tau \leq$ 73 ms the probability of an angle that is not close to the preferred angle is *less* than the probability of a "blank". This suggests active inhibition at off-optimal orientations. We note that this behavior also "inverts" at $\tau = 81$ ms. An explanation for the dynamics of this baseline is another goal of the mathematical modeling.

The results shown in Fig. 1(c) show a response pattern similar to that seen in Fig. 1(b). A unimodal distribution develops and peaks at $\tau = 65$ ms followed by a slight "inversion" of the distribution at $\tau > 115$ ms. However, we also see that for 75 ms $\leq \tau \leq$ 105 ms the distribution is clearly bimodal; it has local maxima at the preferred orientation and at the orientation orthogonal to it and has local minima flanking the preferred orientation. We refer to these profiles as "Mexican-hat" shaped — observe the shape at $\tau = 75$ ms. Such distributions are characteristic of many of the well-tuned cells found in V1. The dynamics of the "blank" baseline in Fig. 1(c) is quite similar to that in Fig. 1(b). In particular, for 65 ms $\leq \tau \leq$ 85 ms off-preferred angles appear to be inhibited – their probabilities are below the blank baseline.

In Fig. 1(d) we present another example of a cell that develops a Mexican-hat distribution profile. Early in the dynamics, for $\tau = 41$ ms, the distribution is quite broad. The distribution then sharpens and by $\tau = 57$ ms the cell is much better tuned. A Mexican-hat profile is evident at $\tau = 65$ ms. Finally, in Fig. 1(e) we present another example of a well-tuned cell that develops a Mexican-hat profile at $\tau = 95$ ms and which later, at $\tau = 115$ ms, exhibits an "inversion".

The dynamical features shown in Fig. 1 are typical of V1 cells. The goal of this study is to explore what kind of models may explain the phenomena of "inversions", the development of Mexican-hat shaped profiles, and the dynamics of the "blank" baseline.

A Feed-forward model

We first present a feed-forward model that has been considered extensively in the literature as a good candidate to explain the spatio-temporal properties of simple cells (Movshon et al., 1978; Citron and Emerson, 1983; Jones and Palmer, 1987; McLean and Palmer, 1989; DeAngelis et al., 1993a; McLean et al., 1994; Reid et al., 1997). It consists of a linear spatio-temporal filter followed by an integrate-and-fire spike generator with leakage:

$$\frac{dv}{dt}(t) = -\lambda(v(t) - v_r) + DC + \int_0^t \iint H(x, y, t - s) I(x, y, s) \; dx \; dy \; ds. \tag{2}$$

$H(x, y, t)$ is the impulse response of a linear spatio-temporal filter, corresponding to a linearized model of the input from the LGN, incorporating inputs from both on-center and off-center cells (Palmer and Davis, 1981; Ferster, 1986; Heeger, 1992b,a). $I(x, y, s)$ is the image shown at time s and λ is the rate of leakage ($1/\lambda$ is the membrane time constant). The leakage has been set to zero in most of the computations presented here; although we do discuss the effects of having nonzero leakage in various cases. DC represents DC input and v_r is the resting potential. A spike is generated when the membrane voltage variable $v(t)$ reaches the firing threshold of -50 mV, the voltage is then reset to $v_r = -70$ mV (Connors et al. (1982); McCormick et al. (1985); Somers et al. (1995); Carandini et al. (1996)). In the model we also mimic the effects of an inhibitory reversal potential: the voltage is not allowed to decrease past a lower barrier of -90 mV (see appendix).

We assume that the kernel H is space-time separable, hence

$$\frac{dv}{dt}(t) = -\lambda(v(t) - v_r) + DC + \int_0^t G(t - s) r(s) ds. \tag{3}$$

where $r(s)$ is the spatial dot product of the image at time s and a spatial kernel,

$$r(s) = \iint K(x, y; \theta_K, \phi_K) \; I(x, y, s) \; dx \; dy. \tag{4}$$

The spatial kernel is taken to be a windowed Gabor function (Marcelja, 1980; Daugman, 1985),

$$K(x, y; \theta_K, \phi_K) = K_0 \; W(d) \; e^{-\frac{d^2}{L^2}} \; \sin(\, \omega(x \cos\theta_K - y \sin\theta_K) - \phi_K \,). \tag{5}$$

This two-dimensional spatial kernel is parametrized by an orientation angle θ_K and phase ϕ_K (the preferred orientation and phase of the simulated cell). In the above, ω is the spatial frequency of the carrier, and $d^2 = x^2 + y^2$. The function is windowed to reflect that in the experiment the stimuli are shown in a circular patch: $W(d) = 1$ if $d < 1$ and 0 otherwise. The length-scale L determines the rate of decay away from the center of the receptive field. Fig. 2(a) shows the Gabor function when $\omega = 3\pi$, $\theta_K = 0$, $\phi_K = 0°$ and $L\omega = 4.2$. We use these parameters for the rest of the article. (Ringach et al., 1997c) found Gabor-like receptive fields with multiple subfields in monkey V1.

In our simulations we use a biphasic temporal kernel:

$$G(t) = \begin{cases} a_e (t/\tau)^5 e^{-t/\tau} & \text{for } 0 \le t \le .05 \\ a_e (t/\tau)^5 e^{-t/\tau} - a_i \left(\frac{(t - .05)}{\tau}\right)^3 e^{-(t-.05)/\tau} & \text{for } t \ge .05 \end{cases} \tag{6}$$

where $a_e = 1.67$, $a_i = 16.7$, $\tau = .01$ sec (see Fig. 2(b)). The temporal kernel is chosen so that $\int_0^\infty G(t) \; dt = 1$ and $\max(G)/\min(G) \sim -5$, with its maximum occurring at 50 ms, its minimum around 80 ms, and G decaying to zero around 150 ms (Gielen et al., 1981). Such a choice of G introduces both time lag and memory effects. G models a magnocellular cell; it has both a positive and a negative response and has a positive mean.