LINEAR ENSEMBLE-CODING IN MIDBRAIN
SUPERIOR COLLICULUS SPECIFIES THE
SACCADE KINEMATICS

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#### List of variables.

| Variable          | Description                                                                 | Value                      |
|-------------------|-----------------------------------------------------------------------------|----------------------------|
| $\hat{S} = [R, \Phi]$ | Saccade vector, with amplitude R and direction $\Phi$                       |                            |
| $\hat{S}_{0k} = [R_{0k}, \Phi_{0k}]$ | Optimal saccade vector for cell k                                          |                            |
| $r(t)$            | Radial trajectory along the saccade vector                                  |                            |
| $h, \dot{v}, \dot{r}$ | Horizontal, vertical and radial eye velocity                              |                            |
| $\dot{h}, \dot{v}$ | Horizontal, vertical eye acceleration                                      |                            |
| $n_k(t)$          | Instantaneous cumulative number of spikes of cell k                        |                            |
| $N_k$             | Total number of spikes in the burst of cell k                              |                            |
| $N_0$             | Maximum number of spikes for optimal saccade                              | $\pm 20$                   |
| $(u,v)$           | Collicular anatomical coordinates (in mm re. fovea)                        |                            |
| $(B_u, B_v)$      | Collicular magnification factors along u and v axes.                       | $(1.4, 1.8)$ mm            |
| $A$               | Shift in the SC mapping function                                            | 3.0 deg                    |
| $B_s$             | Magnification factor for simplified motor map                              | 1.0 mm, A=0                |
| $\eta, \eta_d$   | Normalization constants of SC population for static and dynamic model, respectively. | $2.1 \times 10^{-5}$ deg/spikes/s and $3.9 \times 10^{-4}$ deg/spike |
| $\tilde{m}_k$     | Movement contribution of cell k per spike                                  |                            |
| $\delta(t - t_{k,s})$ | Spike kernel of cell k at time $t_{k,s}$                                   |                            |
| $f_k(t)$          | Instantaneous firing rate of cell k (spikes/s)                             |                            |
| $F_0 / F_k$       | Peak (or maximum mean) firing rate of the population / mean firing rate of cell k | $F_c=800$ spikes/s |
| $P$               | Number of cells in the active population                                    | $\pm 425$                  |
| $\sigma_{pop}$   | Width of the population in the motor map                                   | 0.5 mm                     |
| $\rho_0$          | Cell density in SC motor map                                               | 86.2 /mm$^2$               |
| $\sigma_{bar}$   | Time constant of the burst in ms                                           |                            |
| $\sigma_0, \beta$ | Time constant of the burst at R=0 deg; burst-duration increment             | 3 ms, 0.07 ms/deg          |
| $\gamma$          | Exponent of gamma burst function                                           |                            |
| $\gamma \cdot T_0$ | Time-to-peak firing rate of gamma burst ($= \gamma \cdot \sigma_{bar}$)   | 30 ms                      |
| $S$               | Skewness (of eye-velocity profile, or of gamma burst)                      |                            |
| $V_{sk}$          | Asymptote of the brainstem nonlinearity                                    | 700 deg/s                  |
| $M_0$             | Angular constant of the burst-generator nonlinearity                       | 8 deg                      |
| $B$               | Forward gain of the linear brainstem burst generator                       | 80 deg/s                   |
| $\tau$            | Delay in the brainstem feedback loop                                       | 4 ms                       |
| $M_{H,V}(t)$      | Dynamic horizontal/vertical motor error (in deg)                           |                            |
| $K$               | Offset activity of SC population center-of-gravity                         | 5 spikes/s                 |
ABSTRACT

Recently, we proposed an ensemble-coding scheme of the midbrain superior colliculus (SC) in which, during a saccade, each spike emitted by each recruited SC neuron contributes a fixed minivector to the gaze-control motor output. The size and direction of this ‘spike vector’ depend exclusively on a cell’s location within the SC motor map (Goossens and Van Opstal, 2006). According to this simple scheme, the planned saccade trajectory results from instantaneous linear summation of all spike vectors across the motor map.

In our simulations with this model, the brainstem saccade generator was simplified by a linear feedback system, rendering the total model (which has only three free parameters) essentially linear. Interestingly, when this scheme was applied to actually recorded spike trains from 139 saccade-related SC neurons, measured during thousands of eye movements to single visual targets, straight saccades resulted with the correct velocity profiles and nonlinear kinematic relations (‘main sequence properties’ and ‘component stretching’). Hence, we concluded that the kinematic nonlinearity of saccades resides in the spatial-temporal distribution of SC activity, rather than in the brainstem burst generator. The latter is generally assumed in models of the saccadic system.

Here we analyze how this behaviour might emerge from this simple scheme. In addition, we will show new experimental evidence in support of the proposed mechanism.

**Keywords:** saccades – spatial accuracy – population coding – nonlinearity – main sequence – monkey
INTRODUCTION

Saccadic eye movements: kinematics.

Saccades are rapid eye movements that redirect the fovea fast and accurately to a peripheral stimulus of interest. In this paper we present a computational model on how a population of neurons in the midbrain superior colliculus (SC) encodes the metrics, kinematics, and trajectories of saccadic eye movements.

First, we will briefly describe the main characteristics of the saccade kinematics: the main-sequence, the shape of the velocity profiles, and straight oblique saccade trajectories (Fig. 1). Next, we will introduce the notion of ensemble coding of saccades by the midbrain SC.

Figure 1: Properties of visually-evoked saccades. (Left) The main sequence (NL). Dotted lines (L) indicate responses for a hypothetical linear system. (Center) Skewness of saccade velocity profiles increases with saccade duration. (Right). Component stretching. Here, an oblique saccade in a direction 60 deg re. horizontal ($\vec{S}_{60}$) has components with very different amplitudes, but velocity profiles (bottom) with equal durations and similar shapes. The pure horizontal saccade ($\vec{S}_{0}$) has a much shorter duration and higher velocity ($\vec{H}_{0}$) than the equally large horizontal component of the oblique saccade ($\vec{H}_{60}$).

Main-sequence kinematics. The kinematics of visually-evoked saccades have stereotyped characteristics: the amplitude-duration relation follows a straight line, while peak velocity depends in a nonlinear, saturating way on saccade amplitude. These relations are known as the ‘main sequence’ of saccades (Bahill et al., 1977). Westheimer (1954) recognized that these kinematic relationships betray a nonlinearity within the saccadic system, as for a linear system movement duration would be fixed for all...
saccades, and the peak velocity would increase linearly with the saccade amplitude (Fig. 1, left).\footnote{The saturation of peak velocity follows from the straight-line amplitude-duration relationship. For example, if velocity profiles are approximated by triangles, the amplitude-duration relation $D = a + b \cdot R$ yields for the peak velocity relation: $V_{pk} = 2R/(a + b \cdot R)$, which saturates at $2/b$ deg/s.}

The kinematic nonlinearity is typically assigned to a local feedback circuit in the brainstem, in which so-called medium lead burst cells embody the saccadic pulse generator (Luschei and Fuchs, 1972; Henn and Cohen, 1976; Van Gisbergen et al., 1981). It is thought that these cells are driven by a dynamic motor error signal, which is the difference between the desired endpoint and current eye position. Saccadic burst cells transform this motor error signal into an eye velocity output, and the majority of models of the saccadic system assume that the input-output characteristic of these cells underlies the nonlinear main sequence (Jürgens et al., 1981; Van Gisbergen et al., 1981; Scudder, 1988). However, there is actually very little experimental data to support this latter assumption. We will present evidence for an alternative scheme, which proposes that the nonlinearity of the saccade kinematics may be embedded in the spatial-temporal activation patterns of the midbrain SC.

**Skewness.** A further characteristic property of saccades concerns the shape of their velocity profiles (Fig. 1, center). Typically, the duration of the acceleration phase of saccades (i.e. the time-to-peak velocity) is roughly constant across a wide range of saccade amplitudes. As a result, saccade velocity profiles are skewed. Van Opstal and Van Gisbergen (1989) applied gamma functions to parameterise velocity profiles for different saccade types (visually-evoked, slow saccades due to drugs or fatigue, remembered-target saccades, etc.) and showed that skewness is determined by the saccade duration, rather than by the saccade amplitude.

**Oblique saccades.** A third kinematic nonlinearity of saccades is ‘component stretching’ in oblique saccades (Evinger et al., 1981; Van Gisbergen et al., 1985; Smit et al., 1990). Oblique saccades evoked by single visual targets are approximately straight. This simple fact, however, implies that the horizontal and vertical velocity components of the saccade stay scaled versions of each other throughout the movement:

$$\dot{v}(t) = c \cdot \dot{h}(t) \quad (1)$$

with $c$ a constant. Given the fact that a small saccade has a much shorter duration than a large saccade, Eq 1 requires a mechanism through which the horizontal and vertical components are coupled, such that the shape of their velocity profiles becomes the same despite large differences in amplitude (Fig. 1, right). Different schemes have been proposed to achieve this coupling. For example, Grossman and Robinson (1988) and Nichols and Sparks (1996) proposed that the horizontal and vertical burst generators have their own independent feedback circuits, but are coupled in such a way that they scale each other’s gains. However, when Smit et
al. (1990) applied the cross-coupling model to measured oblique saccades; they demonstrated that the optimal coupling constants are not fixed, but vary in a complex way with saccade amplitude and direction.

A more obvious way to yield straight saccades results from the common source model (Van Gisbergen et al., 1985; Van Gisbergen and Van Opstal, 1989). In this conceptual idea, a nonlinear vectorial pulse generator produces a radial eye-velocity command, \( \dot{r}(t) \), from which the horizontal and vertical velocity components are subsequently derived by linear vector decomposition:

\[
\dot{h}(t) = \cos(\Phi) \cdot \dot{r}(t) \quad \text{and} \quad \dot{v}(t) = \sin(\Phi) \cdot \dot{r}(t)
\]

Note that from Eqn. 1 \( c = \tan(\Phi) \), and that in the common-source scheme component stretching and straight saccades are an emerging property of a shared nonlinear vectorial pulse generator. In cross-coupling models, however, stretching is an additional nonlinear design property needed to account for straight saccades. Although the common-source scheme fitted measured oblique saccade data better than the cross-coupled scheme (Smit et al., 1990), it assumed a vectorial pulse generator for which, at the time, no neurophysiological evidence was available. Here we will argue that the midbrain SC could serve this function.

**Superior colliculus.**

The SC is crucial for the generation of saccades. Its deeper layers contain a topographic map of saccade vectors, which is organized in oculocentric coordinates, as electrical microstimulation produces fixed-vector saccades, the amplitude and direction of which are determined by the site of stimulation within the map (Robinson, 1972). The stimulation parameters have a systematic effect on the saccade properties: low-frequency stimulation produces slower saccades than high-frequency stimulation (Stanford and Sparks, 1996), whereas at low current strengths the evoked saccade amplitude is reduced (Van Opstal and Van Gisbergen, 1990b).

Cells in the motor map fire a brisk burst of action potentials tightly coupled to the onset and duration of the saccade. Near the rostral pole of the SC cells are involved in the generation of small saccades, while at caudal sites cells encode large eye movements. The range of movement vectors for which an SC neuron is recruited is called its movement field (Sparks et al., 1976; Ottes et al., 1986).

Figure 2A-C shows a typical example of a SC movement field for a cell with an optimal saccade vector of \( \vec{S}_0 = [13,230] \) deg, together with the average saccadic eye movement evoked by micro stimulation at the recording site. Note that the evoked saccade (black traces) is almost indistinguishable from the optimal visually-evoked response (gray traces). In Fig. 2D it can be seen that electrically evoked saccades closely correspond with the optimal saccades encoded by cells near the stimulation electrode.

It is generally assumed that the location of the active population of cells in the motor map carries information about the amplitude (R) and direction (\( \Phi \)) of the upcoming saccade vector. There exists strong support
for the notion that activity in the SC motor map is also related to, and may even be responsible for, the kinematic properties of saccades (Berthoz et al., 1986; Van Opstal and Van Gisbergen, 1990b; Waitzman et al., 1991; Munoz et al., 1996; Stanford et al., 1996; Soetedjo et al., 2003; Matsuo et al., 2004). The precise nature of the collicular involvement in the dynamic control of saccades has been controversial, and will be the main topic of this paper.

Figure 2: Movement field of superior colliculus neuron pj9003. (A) Instantaneous firing rate (gray scale) as function of time, for saccades through the center of its movement field, sorted for different amplitudes (amplitude scan). Tick marks indicate spike-counting windows for saccade-related burst (20 ms before saccade onset to 20 ms before offset). Gray trace is the average eye position for the cell’s optimal visually-evoked saccade. Black trace corresponds to the average saccade elicited by micro-stimulation at the recording site. Note close correspondence. (B) Same for a direction scan through the movement field center. Velocity profiles of the saccades are superimposed. (C) Spatial extent of the movement field. Gray scale is number of spikes in the burst. Circles denote saccade end points shown in the left panels. Black trace is the average stimulation-induced saccade trajectory, which ends closely to the center of the movement field. (D) Close correspondence between the optimal saccade vectors of 13 different cells and the stimulation-induced saccades at the recording sites.

We will expand on our recent finding that the instantaneous activity of saccade-related cells in the SC faithfully reflects the instantaneous displacement of the eye in the direction of the planned saccade vector (Goossens and Van Opstal, 2006). This hypothesis was based on the
results of an experimental paradigm in which monkeys made saccades under open-loop conditions toward a briefly flashed visual target. In some trials, an unexpected brief air puff evoked a blink response that coincided with the saccadic eye movement. In those trials, the saccade trajectories were heavily perturbed, and highly variable. Moreover, peak eye velocities decreased by 40% or more, and saccade durations increased, often two-to threefold. Despite these strong perturbations, the saccade endpoint accuracy was virtually unaffected. In other words, the saccade displacement vector remained the same (Goossens and Van Opstal, 2000a).

Also the saccade-related activity in the SC was strongly affected by the blink perturbations: the mean- and peak-firing rate in the burst dropped substantially, while the burst duration increased to match the longer saccade duration (Goossens and Van Opstal, 2000b). However, the relation between instantaneous firing rate and dynamic motor error, which is nearly linear in control saccades (Waitzman et al., 1990), broke down in perturbed trials (Goossens and Van Opstal, 2000b; Keller and Edelman, 1994; Munoz et al. 1996). This indicated that the SC cells do not encode the instantaneous motor error.

Interestingly, the number of spikes in the saccade-related burst was the same for perturbed and control saccades. A similar observation had been made earlier by Munoz et al. (1996) for saccades that were briefly interrupted by stimulation in the brainstem omnipause region.

Our findings led to the hypothesis (Goossens and Van Opstal, 2006) that the instantaneous cumulative number of spikes in the burst, \( n(t) \), reflects the instantaneous eye displacement along the saccade vector, \( r(t) \). The data indeed showed that the saccade-related burst of all SC cells could be well described by this idea, both for control and for highly perturbed saccades. The consequence of this simple formulation was (i) that each spike in the SC contributed a fixed, tiny movement vector to the saccade, and (ii) that movement fields of SC cells would be dynamic. These predictions were fully explored in Goossens and Van Opstal (2006) and led to the formulation of the dynamic linear ensemble-coding model. Before introducing this model, we will first briefly review the concept of ensemble coding by the SC.

**Ensemble coding: static**

The idea of ensemble coding proposes that a large population of coarsely-tuned neurons in the SC motor map is recruited for a saccade, and that every cell generates its own movement vector. All contributions are then somehow combined to produce an accurate saccade vector. This idea was first forwarded by McIlwain (1982), and formulated into a quantitative model by Ottes et al. (1986) and Van Gisbergen et al. (1987). In their movement field model, Ottes et al. (1986) assumed that the shape and size of the recruited population, when expressed in collicular Cartesian coordinates \([u,v]\) (in mm relative to the foveal representation), is invariant for all saccades, and that this ‘mould’ of activity can be described by a Gaussian activation profile, centered in the motor map around the point image \([u_0,v_0]\) of the saccade vector \([R_0,\Phi_0]\), with a fixed width of \(\sigma_{\text{pop}}\) (in mm) and a peak mean firing rate, \(F_0\):
\[ F[u,v] = F_0 \cdot \exp \left[ -\frac{(u-u_0)^2 + (v-v_0)^2}{2\sigma_{\text{pop}}^2} \right] \] (3)

Here, the complex-logarithmic afferent mapping function \([R, \Phi] \rightarrow [u,v]\) is given by:

\[
u(R,\Phi) = B_u \ln \left( \frac{\sqrt{R^2 + 2AR \cos \Phi + A^2}}{A} \right) \quad \text{and} \quad v(R,\Phi) = B_v \arctan \left( \frac{R \sin \Phi}{R \cos \Phi + A} \right) \] (4)

The mapping parameters \([B_u, B_v, A] = [1.4 \text{ mm}, 1.8 \text{ mm/ rad}, 3.0 \text{ deg}]\) were determined from Robinson’s (1972) micro-stimulation data. The model describes the mean firing rate of a given cell in the SC motor map for any saccade vector with amplitude \(R\) and direction \(\Phi\), and accounts well for the observed asymmetric shape of SC movement fields (e.g. Fig. 2A, B). It also accounts for the fact that cells encoding small saccades have movement fields extending over a much smaller amplitude range than cells recruited for large saccades.

Van Gisbergen et al. (1987) applied this model in an attempt to understand how the neural population of Eq. 3 could then encode the planned saccade vector. To that end, they assumed that each recruited cell, \(k\), located in the motor map at \([u_k, v_k]\) would contribute a tiny movement vector, \(\vec{m}_k\), with a size, \(r_k\), and direction, \(\phi_k\), that was fully determined by its fixed efferent projections to the horizontal \(x_k\) and vertical \(y_k\) brainstem burst generators, according to:

\[
x_k = \eta \cdot A \cdot \exp \left( \frac{u_k}{B_u} \right) \cdot \left[ \cos \left( \frac{v_k}{B_v} \right) - 1 \right] \quad \text{and} \quad y_k = \eta \cdot A \cdot \exp \left( \frac{u_k}{B_u} \right) \cdot \sin \left( \frac{v_k}{B_v} \right) \] (5)

Note that the efferent mapping of Eq. (5), \([u,v] \rightarrow [R, \Phi]\) is the inverse of the afferent mapping of Eq. 4 (apart from the fixed scaling \(\eta = 1/(F_0 \cdot \sigma_{\text{pop}}^2 \cdot 2\pi \cdot \rho_0)\), with \(\rho_0\) the number of cells per mm\(^2\)). A cell’s preferred movement vector is hence given by \(\vec{S}_{\text{pol}} = F_0 \cdot \vec{m}_k / \eta\).

In the ensemble-coding model of Van Gisbergen et al. (1987) the saccade vector, \(\vec{S} = [R, \Phi]\), is then determined by linear summation of all cell contributions in the population, multiplied by their mean firing rates, \(F_k\):

\[
\vec{S} = \sum_{k=1}^{P} F_k \cdot \vec{m}_k \] (6)

with \(P\) the total number of cells in the population. The model of Van Gisbergen et al. (1987) explains how a large, invariant population (Eq 3) within a logarithmically compressed motor map (Eq 4) can represent saccade vectors with the correct amplitude and direction throughout the oculomotor range, and accounts for the shape of SC movement fields without any further adjustment of parameters (Fig. 2).

However, the model did not incorporate a mechanism to explain the saccade kinematics, nor did it account for component stretching in oblique saccades. Thus, it remained unclear how to combine the ensemble-coding scheme of Eq. 6 with either the common source model of Eq. 2, or with mutually coupled brainstem burst generators, although Smit et al. (1990) and Van Gisbergen and Van Opstal (1989) speculated that perhaps the vectorial pulse generator was embodied by the motor SC.
Lee et al. (1988) noted a further problem with the linear summation model. They argued, on the basis of local reversible inactivation experiments within the motor map, that vector averaging could better account for the observed data than linear summation. Thus, according to their proposal, the saccade vector was computed by:

$$\vec{S} = \sum_{k=1}^{P} F_k \cdot \vec{S}_{0k} / \sum_{k=1}^{P} F_k$$

(7)

Note that Eq 7 incorporates normalization by the total activity within the population. This requires a nonlinear mechanism for which it is not obvious how it might be implemented through fixed SC-brainstem connections.

**Ensemble coding model: dynamic**

The ensemble-coding model of Goossens and Van Opstal (2006) was proposed to capture the findings of the blink-perturbation paradigm (see above). In contrast to the static ensemble-coding model (Eq 6) of Van Gisbergen et al. (1987), they introduced dynamics into the motor map output, by assuming that each spike $s$ of each cell $k$ in the motor map, fired at time $t_{s,k}$ generates a tiny movement vector, here called a 'spike vector', which is given by:

$$\vec{s}_k(t) = \vec{m}_k \cdot \delta(t - t_{s,k})$$

(8)

Here, $\vec{m}_k$ is the effective connection vector of SC neuron $k$ to the brainstem burst generator (Eq 5), and the spike kernel $\delta(t - t_{s,k}) = 1$ for $t = t_{s,k}$ and 0 elsewhere.

Furthermore, $\vec{S}_{0k} = N_0 \vec{m}_k / \eta_d$ is the cell’s preferred movement vector (i.e. the center of its movement field), where the population scaling factor is determined by $\eta_d = 1/(N_0 \cdot \sigma_{pop}^2 \cdot 2\pi \cdot \rho_0)$. Here, $N_0$ is the fixed maximum number of spikes in the population.

The planned saccade trajectory, generated by the population of recruited neurons, is then determined by instantaneous linear summation of all spike vectors:

$$\vec{S}(t) = \sum_{l=1}^{P} \sum_{s=1}^{N_k} \vec{s}_s(t)$$

(9)

with $P$ the total number of cells in the population, and $N_k$ the total number of spikes (fired at times $t_{s,k}$) in the burst of cell $k$.

Several predictions of this simple model on the behaviour of SC cells were verified in Goossens and Van Opstal (2006), for which the most important ones are:
Movement fields of SC cells are *dynamic*; for saccades of any amplitude and direction the cumulative number of spikes in the burst of cell \( k \), \( n_k(t) \), predicts how far the eye has moved along either the cell’s preferred movement vector, \( \vec{S}_{0k} = [R_{0k}, \Phi_{0k}] \), or along the actual saccade vector, \( \vec{S} = [R, \Phi] \), i.e.:

\[
n_k(t) = a_k \cdot r(t)
\]

(10)

with \( a_k \) the slope in \#spikes/deg, and \( r(t) \in [0, R] \) the radial eye displacement along the saccade vector \( \vec{S} \).

The slope of Eqn. 10 is determined by \( a_k = \frac{N_k(R, \Phi)}{R} \), in which

\( N_k(R, \Phi) \) (the total number of spikes in the burst of cell \( k \) for the saccade) replaces the mean firing rate \( F_k \) in the static movement field description of Eqn. 3.

Eq 10 holds for fast, as well as for extremely slow (e.g. blink-perturbed) saccades to the same target.

With the additional assumption that the total number of spikes of the saccade population is fixed for all saccade amplitudes, and that this number controls the saccade offset (e.g. through a downstream threshold mechanism), the results of the micro-lesion experiments of Lee et al. (1988) could be explained without the need for the nonlinear vector averaging scheme of Eq. 7 (Goossens and Van Opstal, 2006).

To test the possibility that the SC population encoded the saccade kinematics, the result of Eq. 9 was fed into two independent, but *linear*, feedback circuits that represented the horizontal and vertical saccade burst generators.

When this simple model was applied to actual recordings taken from 140 saccade-related burst neurons during thousands of saccades, it was able to reproduce the metrics, straight trajectories, the main-sequence kinematics and velocity profiles of saccades to single visual targets across the oculomotor range (Goossens and Van Opstal, 2006). Note that in the reconstruction of these saccades the neuronal activity was not normalised in any way. Quite remarkably, only three free parameters (the forward gains, \( B_H \) and \( B_V \), and the delay, \( \tau \), in the brainstem feedback loops) sufficed to produce excellent correspondence between real saccades and model saccades, and between recorded activity patterns and actual activity patterns.

Because the applied SC-brainstem model is entirely linear, the fact that reconstructed saccades were straight and obeyed the main sequence had to result from the properties of the model’s input, i.e. the measured spatial-temporal activity patterns in the SC motor map. In other words, the saccade reconstructions strongly suggested that the motor SC might act as a *nonlinear vectorial pulse generator*.

The question as to which aspects of the input patterns could be responsible for these properties was not explored in the Goossens and Van Opstal (2006) study. Therefore, the present paper proposes a
mechanism through which the SC population could generate the nonlinear saccade kinematics.
METHODS

Simulations were performed in Matlab 7.4 (version R2007a) with a simplified description of the afferent collicular motor map:

\[ u(R) = B_u \cdot \ln(R) \quad \text{and} \quad v(\Phi) = B_v \cdot \Phi \]  
(11)

with \( B_u = 1 \) (isotropic map). This yields for the efferent mapping function:

\[ x = \exp(u) \cdot \cos(v) \quad \text{and} \quad y = \exp(u) \cdot \sin(v) \]  
(12)

The instantaneous firing rate of cell \( k \) (at location \( (u_k, v_k) \) in the motor map) during a saccade with coordinates \( [R, \Phi] \) (with point image \( [u_R, v_\Phi] \) in the motor map) was described by:

\[ f_k(u_k, v_k, t) = F_0 \exp \left( \frac{(u_k - u_R)^2 + (v_k - v_\Phi)^2}{2\sigma_{pop}^2} \right) \cdot s(t) \]  
(13)

with \( s(t) \) the cell’s temporal activity profile. In our simulations, we applied a normalised gamma burst to describe this temporal behaviour:

\[ s(t) = T_0 \cdot (t - t_{ON})^\gamma \cdot \exp(-\frac{(t - t_{ON})}{\sigma_{DUR}}) \]  
with \( T_0 = \sigma_{DUR} \cdot \gamma/e \quad \text{and} \quad t \geq t_{ON} \]  
(14)

Note that the amplitude of Eq. 14 is normalised to one, \( t_{ON} \) is the burst onset, and \( \sigma_{DUR} \) is a measure for the burst duration. The exponent \( \gamma \) determines the gamma-burst skewness by \( S = 2/\sqrt{(\gamma + 1)} \), and \( T_0 \cdot e = \sigma_{DUR} \cdot \gamma \) is the time-to-peak firing in the gamma burst.

The motor map consisted of a rectangular \( [u, v] \) grid of 51x51 neurons, in which \(-4.8 \leq u \leq +4.8 \) mm and \(-\pi/2 \leq v \leq +\pi/2 \) mm. We took a fixed population width of \( \sigma_{Pop} = 0.5 \) mm, and the scaling constant \( \eta_d = 3.9 \times 10^{-4} \) deg/spike was determined by tuning the model for a 20 deg horizontal saccade. For each saccade a population of about \( P=425 \) cells was recruited, which generated a fixed number of about 2540 spikes. The central cell in the population fired between 19-20 spikes.

In our simulations the linear brainstem was given identical gains of \( B_H = B_V = 80 \) deg/s, with a feedback delay of \( \tau = 4 \) ms, for both the horizontal and vertical eye-movement components. For simplicity, the plant transfer characteristics were ignored. Thus, the model’s saccade trajectories were completely reflected in the SC population output of Eqn. 9.

Figure 3 provides a schematic overview of the model described by Eqns. 11-14.
Figure 3: Schematic of the model used in the simulations. The SC is modeled as a rectangular grid of 51x51 cells (Eq. 11). Here, only one SC is shown, with the population centered at \([R, \Phi]=[15,45]\) deg. The linear brainstem contains two independent feedback loops, which in our simulations were taken to be identical.

Figure 4 shows a movement field scan for a model neuron with an optimal saccade vector of \([R,\Phi]=[15,0]\) deg. Fig. 4A shows a classical static movement field description in which the cell’s activity is quantified by the total number of spikes in the burst for saccades in the optimal (horizontal) direction (left), and for optimal saccade amplitudes (15 deg) in different directions (right). Panel B shows the gamma-burst activity for this unit as function of time (Eq 15; \(T_0=11\) ms) for the same saccade vectors as in panel A. Note that these model responses correspond well with the actual data shown in Fig. 2.

Figure 4. Movement field of a SC model cell with an optimal saccade vector of \([R,\Phi]=[15,0]\) deg. (A) Left: Total number of spikes in the gamma-burst as function of saccade amplitude (direction 0 deg); right: number of spikes as function of saccade direction (amplitude 15 deg). As a result of the logarithmic compression of the motor map, the amplitude tuning curve is asymmetric, in contrast to the direction tuning curve. (B) Gamma-burst activity profiles for the amplitude (left) and direction (right) scans. Scale: firing rate in spikes/s.
RESULTS

As mentioned in Methods, despite the logarithmic compression of the afferent mapping function (Eq 4), the saccadic output of the total population has linear kinematics when the parameters describing the cell activity in Eqs 13-14, $F_{pk}$, $\sigma_{pop}$ and $\sigma_{Dur}$, are independent of the saccade amplitude, or of the cell’s location in the motor map. This property of the model is illustrated in Figure 5, which shows a number of horizontal saccadic eye movements (Fig 5A) with their corresponding velocity profiles (Fig 5B), together with the resulting main-sequence relations (inset Fig 5B).

![Figure 5: The model behaves as a linear system, when the temporal burst profiles have fixed parameters. (A) Eye position traces. (B) Velocity profiles scale linearly with saccade amplitude. Inset: main sequence.](image)

The reason for this overall linear behaviour is that the exponential nature of the efferent mapping function exactly cancels the logarithmic compression of the afferent map, and that this apparently also holds when a large population of cells is involved. Thus, if the population activity profile is taken invariant across the motor map (i.e. fixed values for $F_0$, $\sigma_{pop}$ and $\sigma_{Dur}$ for all cells in the SC), the ensemble-coding model of Eqn. 9 will not yield a nonlinear main sequence. Instead, saccade durations will be constant for all amplitudes, and the peak eye velocity will vary linearly with saccade amplitude (Fig. 5B, inset). Also the skewness of the saccade velocity profiles will be fixed (cf. Fig. 1).

Note that a fixed population size, $\sigma_{pop} = 0.5$ mm, together with a homogeneous cell density across the SC, implies that the total number of cells contributing to any visual evoked saccade is constant. And if the firing rates $F_{pk}$ and burst durations $\sigma_{Dur}$ are the same for each location in the motor map, the total number of spikes emitted by the population is the same too. Experimental evidence given in Goossens and Van Opstal (2006; their Fig. 7) indeed support the assumption that the size of the recruited population as well as the total number of spikes emitted by the population are the same for each saccade vector. The question therefore arises, whether the other burst properties are indeed independent of cell’s location in the motor map.
In Fig. 6A (top, left) we show that the number of spikes of cells in the motor SC for their optimal saccades is the same across the motor map, as the correlation between the spike count and optimal saccade amplitude is not significant. The bottom panel in Fig. 6A, however, shows an additional analysis for these cells, which indicates that the peak firing rate of SC cells varies systematically with their optimal saccade amplitude.

Figure 6: Burst properties of SC saccade-related cells. Cells (N=77) were selected for having at least five saccades into the center of their movement field (within 0.5σpop). Highlighted cells (n=32) are selected for producing N_0=20 spikes for their optimal saccade. (A) Top: number of spikes is not related to the optimal saccade amplitude. Bottom: Peak firing rate of the spike-density function, however, decreases systematically as function of a cell's optimal amplitude (B) Average spike-density burst profiles (peaks normalised) for the four clusters of cells show a clear increase of burst duration (and skewness) with saccade amplitude. Inset: average optimal radial saccade position - and velocity traces for the four cell groups. Note main-sequence behaviour and skewness.

Fig. 6B shows the normalised temporal burst profiles for the four selected amplitude clusters for which N_0=20 spikes (32 cells, asterisks in Fig. 6A). Note that the peak-firing rate occurs at about the same instant re. saccade onset, while the burst duration clearly increases with the optimal amplitude. Hence, also the burst skewness increases with saccade amplitude. The gamma function of Eq. 14 therefore seems to be a reasonable description for the shape of the saccade-related bursts.

Thus, in contrast to the simplified assumption made in the simulations of Figure 5, the burst characteristics of SC cells do appear to vary in a systematic way with their optimal saccade vector (i.e. with their location within the motor map). To investigate whether the nonlinear main
sequence properties may result from neural populations endowed with these characteristics, we incorporated these features into the spike-vector model.

To that end, we let the burst duration as well as a cell’s peak firing rate depend on the saccade amplitude, $R$, according to$^2$:

$$\sigma_{\text{Dur}}(R) = \sigma_0 \cdot [1 + \beta \cdot R] \quad \text{and} \quad F_{\text{pk}}(R) = \frac{F_0}{\sqrt[2]{1 + \beta \cdot R}}$$

in which $F_0$, $\sigma_0$ and $\beta$ are constants (see Fig. 7A). In accordance with the data shown in Fig. 6, cells near the rostral pole then tend to have shorter saccade-related bursts, with a higher peak-firing rate, than cells in more caudal areas of the motor map, while at the same time keeping the number of spikes for the optimal saccade constant. Fig. 7 shows these relations, as they were used in our model simulations. We took $\sigma_0=3$ ms, $F_0=800$ spikes/s, and $\beta=0.07$ ms/deg.

Figure 7: (A) Peak firing rate and burst duration (width, $\sigma_{\text{Dur}}$; left axis), and the number of spikes and burst skewness (right axis) of each model cell as function of its optimal saccade amplitude (Eqns. 14-15). (B) Examples of SC gamma bursts with the properties shown in (A).

Figure 7B shows the temporal activity profiles of the most active cell in the recruited population for a number of different saccade amplitudes. Note the drop in the cell’s peak firing rate and the increase in burst duration, keeping the total number at $N_0=19$ spikes. These burst properties resemble the measured burst profiles shown in Fig. 6B.

Figure 8 illustrates how the model generates saccades in different directions and with different amplitudes. In Fig. 8A the motor activity in the SC map can be seen at the time of maximum firing (30 ms after burst onset) for a saccade with an amplitude of 20 deg, and a direction of 120 deg.

$^2$ Alternatively, burst duration and peak firing rate could depend on the cell’s optimal saccade amplitude, $R_{0\text{k}}$. In that case, the burst properties depend exclusively on the cell’s location in the motor map, as $R_{0\text{k}}=\exp(u_k)$, rather than on the actual saccade, $R$ (Eqn. 10). See also the Discussion for this subtle, but important, conceptual difference.
deg (left and upward). Note that the activity transgresses the vertical meridian (the horizontal boundary at $v=\pi/2$ mm), as a small part of the population activity is also found in the left SC. This adheres to the ‘gluing problem’ (Van Opstal and Van Gisbergen, 1989; Tabareau et al., 2007). The insets in this panel show the horizontal/vertical eye position traces, the radial eye velocity, and the straight spatial trajectory of the saccade. Figure 8B shows that the model generates accurate saccades in all directions and across all amplitudes.

Figure 8 (A) Snapshot of the model’s activity in the motor map for a left-upward saccade at $[R,\Phi]=[20,120]$ deg. Insets in this panel show horizontal and vertical eye-position traces (top-right), eye-velocity profiles (bottom-right; note component stretching), and the associated 2D trajectory (bottom-left). The activity snapshot was taken at the moment of maximum firing of the gamma burst activity profiles (30 ms after burst onset). Note that activity is divided across the two motor maps. Grid superimposed on the anatomical $[u,v]$ coordinates shows iso-amplitude (running vertical, at $R=0,2,5,20,50$ deg) and iso-direction (running horizontal, $\Phi=-90,-60,-30,0,30,60,90$ deg) lines of the motor map (Eq. 11). The $u=0$ line in the center separates the two colliculi. (B) Targets (squares) and model saccade endpoints (black dots) for 63 locations. The target at $(20,0)$ deg (highlighted) was used to tune the model’s weighting constant to $\eta=0.00039$. Three example trajectories are also shown (gray lines).

The introduction of an amplitude-dependent increase of a cell’s burst duration, and a concomitant decrease in its peak firing rate (Eq. 15) had a dramatic effect on the kinematics of saccades. Yet, the trajectories of oblique eye movements remained straight. Figure 9A now clearly demonstrates that the duration of saccades increases with their amplitude. Figure 9B shows the saccade velocity profiles generated by the model, as well as the main-sequence properties of these eye movements. The main sequence behaviour of the model now resembles actual saccades quite well. In addition, the model also captures the duration-skewness relation of the velocity profiles. Thus, the mechanism described
by Eqn. 15, and illustrated by the recordings from a population of SC neurons in Fig. 6, may indeed underlie the nonlinear main-sequence properties of saccades, and constitute the mechanism by which the motor SC could act as a vectorial pulse generator.

Figure 9: Traces of eye position (A) and eye velocity (B) for simulated horizontal saccades to targets at R=[2, 5, 9, 14, 27, 35] deg eccentricity. Note the asymmetric shape of the velocity profiles, which is due to a fixed acceleration time (determined by $e \cdot T_0 = 30$ ms, Eq. 14). The main sequence for these saccades is shown in the insets of panel B.

An important prediction of the spike-vector ensemble-coding model is the linear relationship between the instantaneous cumulative number of spikes in the burst of a given SC cell, and the instantaneous displacement of the eye along the saccade trajectory (Eqn. 10).

To illustrate this feature of the model, Figure 10 shows these phase plots for a cell with an optimal saccade vector at [10,0] deg. The cell response is shown for saccades of different amplitudes, each yielding a different number of spikes, and their own temporal profile, according to Eqns. 13-14. The phase plots in Fig. 10A are straight, even for the saccades with a low number of spikes in their burst. This property results from the assumption that burst durations and peak firing rates depend systematically on the saccade vector rather than on the cell’s location in the motor map. Fig. 10B shows that the phase plots for very slow saccades (simulated by setting $\beta = 0.25$ ms/deg) of the same amplitudes are indistinguishable from those of fast saccades. The dynamics of the spike counts for the two conditions, however, are quite different, as is shown in Fig. 10C. Both properties of the model are in line with the data reported in Goossens and van Opstal (2006; their Figs 4 and 5) and with Eqn. 10.

It is important to realise that straight phase trajectories, regardless the saccade kinematics, are a consequence of the idea that the SC is directly involved in the motor program of the saccadic eye movement. This idea contrasts with other hypotheses, which assume that the SC cells encode only the desired vectorial displacement of the eye, and have no direct motor function (see Discussion).
Figure 10. (A) Cell activity during saccades into (gray symbols) and out of (black dots) the center of its movement field (centered at \([R, \Phi]=[10, 0]\) deg). Saccade amplitudes varied from \(R=[2, 5, 10, 15, 20, 25, 30, 35]\) deg. The cumulative spike count is shown as function of the instantaneous eye displacement; individual dots correspond to time samples, not spikes. The cell's burst duration (and hence its peak mean firing rate) depended on saccade amplitude, \(R\) (Eqn. 15, \(\sigma_0=3 \text{ ms}; \beta=0.07 \text{ ms/deg}\)). The phase plots are straight, even for the saccades with a low number of spikes, and their slopes depend systematically on the saccade vector (Eqn. 10). (B) Same plots for slow saccades, simulated by setting \(\beta=0.25 \text{ ms/deg}\). The phase plots of slow saccades are indistinguishable from the fast saccades in (A). (C) Cumulative spike counts of fast (gray) and slow (black) saccades follow different dynamics, because of their different saccade kinematics.
DISCUSSION

In this paper we have analysed our dynamic ensemble-coding model of the motor SC, in order to pinpoint the potential mechanism that may render the SC to act as a nonlinear vectorial pulse generator. We conclude, both on the basis of our recordings, illustrated in Fig. 6, and fully described in Goossens and Van Opstal (2006), as well as on the simulations with our model (Figs. 8-10), that the nonlinear kinematics of saccades are due to a gradient along the motor map of the burst properties of saccade-related cells. To summarise, saccade-related bursts can be characterised as follows:

(i) The total number of spikes for a cell’s optimal saccade is fixed across the motor map (Fig. 6A, top).

(ii) For a given saccade, the total number of spikes of a cell is determined by the amplitude and direction of that saccade according to its classical movement field description (Ottes et al., 1986; Eqs. 3-4; Fig. 3).

(iii) The temporal distribution of the spikes can be described by gamma functions, with their peak firing rates at a fixed time relative to the saccade onset.

(iv) The shape parameters of the burst are determined by saccade amplitude, such that the peak-firing rate decreases, while at the same time, the burst duration (and skewness) increases with the saccade amplitude (Eqn. 15; Fig. 6B).

Without the latter mechanism, the dynamic ensemble-coding model of Fig. 3 behaves as a linear system (Fig. 5).

Why a nonlinear main sequence?

We are well aware that we have grossly simplified the brainstem feedback loops by modeling them as linear systems (a gain with a feedback delay). The sole reason for this simplification, however, was to verify whether the spatial-temporal activity patterns in the motor SC could fully encode the saccade kinematics, without having to resort to nonlinear mechanisms like normalization of activity (e.g. center-of-gravity computation) or saturation of the brainstem pulse generator. Our previous study (Goossens and Van Opstal, 2006) showed that the linear ensemble-coding model produced realistic saccades with compelling accuracy, while needing only three free linear parameters to generate the saccade kinematics across the entire repertoire of collicular activity patterns. We therefore decided to explore the properties of this scheme in greater detail.

In almost every model of the saccadic system, the brainstem local feedback loops are described by a saturating nonlinearity that mimics the amplitude-peak velocity relationship (e.g. Van Gisbergen et al., 1981; Jürgens et al., 1981; Scudder, 1988). The question is justified why the burst generator would contain this nonlinearity in the first place. It is not likely that it reflects a mere passive neural saturation, or neural fatigue, for several reasons: First, it has been shown that slow saccades (e.g. saccades to remembered targets in darkness) obey their own nonlinear main sequence (Smit et al., 1987). In addition, abducens oculomotor neurons reach firing rates that are comparable to those of the pontine
burst neurons. Yet, models of the saccadic system invariably employ linear transfer characteristics to describe these other types of neurons. Moreover, even though oculomotor neurons have nonlinear characteristics (as they are recruited beyond a threshold, essentially behaving as rectifiers), it is generally assumed that the output of the total neural population varies approximately linearly with changes in eye position.

Interestingly, a recent theoretical study by Harris and Wolpert (2006) suggested that the main-sequence properties of saccades could reflect an optimal control strategy of the system, as it has to cope with several conflicting constraints. The function of saccades is to redirect the fovea as fast and as accurately as possible to a peripheral target. However, the properties of internal noise within the system (assumed to increase with activity levels; this is e.g. visible in the movement field scans of Fig. 2), a low spatial resolution in the peripheral retina, and a penalty for overshooting the target (as commands then have to cross hemispheres), require a trade-off between movement duration and accuracy. Their analysis indicated that the optimal trajectory to satisfy the constraints obeys the main-sequence relationships.

Earlier theoretical work by Harris (1995) had indicated that, given the main-sequence properties of saccades, another optimal strategy of the system to acquire the target on average as fast as possible would be to undershoot the target by about 10%. Indeed, the phenomenon of saccadic undershoots is well established. Even when an experimental manipulation (e.g. magnifying lenses) produces accurate saccades on average, the system rapidly adapts to the new situation by developing undershoots (Henson, 1978).

Thus, there appears to be a strong case against mere passive mechanisms, like neural saturation, that determine the motor control of saccades. Instead, the main sequence could result from a deliberate built-in optimal control strategy.

If so, one might wonder whether the optimal nonlinear controller could further benefit from embedding the main-sequence nonlinearity (and the undershooting strategy) at a vectorial encoding stage, rather than at the brainstem level of the horizontal and vertical saccade components. According to the common-source model of Van Gisbergen et al. (1985), an obvious benefit of a nonlinear vectorial pulse generator is that oblique saccade trajectories will automatically be straight without the need for an elaborate cross-coupling scheme (Smit et al., 1990).

Of course, straight saccade trajectories by themselves would be optimal in the sense that they constitute the shortest possible path between the start- and end points. Note that cross-coupling schemes such as proposed by Grossman and Robinson (1988), and later also by Nichols and Sparks (1996), could work as well as the common-source model to produce straight saccades. However, having (at least) two saturating nonlinearities in the brainstem leads to coupling coefficients that depend in a complicated way on the saccade amplitude and direction (Smit et al., 1990). In other words, generating straight saccades with coupled and unequal nonlinear horizontal and vertical burst generators requires an entire map of horizontal and vertical coupling coefficients. Moreover, these coefficients depend also on the saccade type (visually-evoked vs.
remembered in darkness), which seems to be hardly an efficient and plausible network design.

In contrast, in the common source model component cross-coupling (Fig. 1) is an emergent property, which requires no additional tuning of the horizontal and vertical burst generators. Indeed, in our model (Fig. 2) the horizontal and vertical feedback loops are independent, yet the saccades are straight and the components show the appropriate amount of stretching. Moreover, fast and slow saccades are produced by the same mechanism, and therefore do not require separate sets of tuning parameters.

Note that the vectorial pulse-generator model does not imply that the horizontal and vertical saccade components always have equal durations. For example, when the gains of the horizontal and vertical feedback loops are different, saccade trajectories will curve toward the faster component. Patients suffering from Niemann-Pick type C disease appear to suffer from a deficit that selectively affects their vertical saccades (Rottach et al., 1997). It seems as if the gain of their vertical saccade generator may be decreased to only 10% of that of the horizontal system. Consequently, their oblique saccades are heavily curved toward the horizontal. The model of Figure 3 can faithfully reproduce the responses of these patients, by simply letting $B_V = 0.1 \cdot B_H$ (data not shown).

**Burst properties depend on the actual saccade vector.**

In our simulations, the SC burst properties were assumed to depend on the actual saccade amplitude, $R$ (Eqn. 15). Note that this is not a trivial assumption, as the burst duration and peak-firing rate could instead have depended on the cell’s optimal saccade amplitude, $R_{0k}$. In that case, the burst properties would have read:

$$\sigma_{DUR}(u_k) = \sigma_0 \cdot [1 + \beta \cdot \exp(u_k)] \quad \text{and} \quad F_{PK}(u_k) = \frac{F_0}{\sqrt{1 + \beta \cdot \exp(u_k)}}$$

where $u_k = \ln(R_{0k})$. Simulations with SC cells obeying Eqn. 16 resulted in saccades that followed a virtually identical main sequence as with Eqn. 15 (not shown).

Conceptually, however, Eqns. 15 and 16 describe quite different mechanisms to embody the main-sequence nonlinearity. According to Eqn. 15 the temporal distribution of spikes in the burst, as well as the peak firing rate are determined by the actual saccade, $\tilde{S}$, in which the cell is participating. Hence the temporal burst properties, as well as the number of spikes, are fully determined by the cell’s input. Such a cell will fire a briefer and more intense burst of spikes when it takes part in saccades smaller than its optimum saccade, than when it participates in saccades that exceed its optimal amplitude, but for which it fires the same number of spikes. In this way, the distribution of firing rates within the active cell population is circular symmetric about its center in the motor map even though there is a rostral-to-caudal decrease in the firing rate of cells for their respective optimal vectors.

In the case of Eqn. 16, however, the number of spikes is determined
by the input, but the temporal firing properties of the cell are only determined by its location within the motor map, \([u_k, v_k]\), and hence by its optimal saccade, \(S_0k\). Such a cell will fire identical spike trains for different saccades for which the number of spikes is the same. This would mean, however, that the distribution of firing rates within the active cell population would be skewed, since Eqn. 16 implies that the rostral cells within this population have higher firing rates than caudal cells located at the same distance from the center. Our recordings support the scheme of Eqn. 15 (not shown; manuscript in preparation).

A second difference between the two proposals is that the phase plots of \(r(t)\) vs. \(n_k(t)\) (see Fig. 10) will be straight for cells obeying Eqn. 15, but they will be systematically curved for cells that are ruled by Eqn. 16 (not shown).

**Center-of-gravity weighting combined with nonlinear feedback?**

The experiments described in Goossens and Van Opstal (2006) suggested that the deep layers of the SC are involved in programming a saccadic motor command, and hence that the instantaneous activity (quantified by the cumulative number of spikes) relates linearly to the instantaneous eye displacement along the saccade trajectory (Eq. 10; Fig. 10).

However, alternative theories have been forwarded, that assign quite a different role to these cells: instead of encoding a motor command, they are thought to encode the saccade goal, imposed by the retinal target location (Port and Wurtz, 2003; Krauzlis et al., 2004; Walton et al., 2005). According to this theory, SC cells are not involved in generating the actual saccade trajectory and kinematics, but their weighted and scaled output (the center-of-gravity of the population, Eq. 7) represents the instantaneous desired eye-displacement signal toward the goal. This saccade goal corresponds to the eye-centered location of the target, and is then thought to drive mutually coupled nonlinear feedback loops in the brainstem (e.g. Port and Wurtz, 2003; Walton et al., 2005).

We wondered, whether the observed relations between SC firing patterns and eye movements (like in Fig. 10) could also be obtained for such a scheme. Obviously, when the center-of-gravity computation is taken literally (Eq. 7), even a single spike in the motor map would already encode the final saccade goal, which is biologically implausible. Moreover, micro stimulation at low current intensities has been shown to yield saccades with smaller-than-optimal amplitudes (Van Opstal et al., 1990), which cannot be explained by an ideal center-of-gravity computation. A small change in the formulation of Eq. 7, however, could capture this phenomenon. Suppose, that the output of the SC yields a dynamic estimate of the saccade goal according to:

\[
\hat{S}_G(t) = \frac{\sum_{k \in Pop} f_k(t) \cdot \hat{S}_{0k}}{K + \sum_{k \in Pop} f_k(t)}
\]

in which \(f_k(t)\) is a cell’s instantaneous firing rate, given by Eq. 13, \(\hat{S}_{0k}\) is
the cell’s optimal saccade vector, and K (in spikes/s) is a fixed constant. As burst activity profiles we took gamma bursts with a fixed duration, by setting $\alpha_0 = 4.5$ ms (i.e. no implicit main-sequence properties in burst duration and peak firing rate like in Fig. 7, Eqn. 15). When the total activity in the population is low, K dominates the numerator and the desired saccade vector will be small. When the activity is high, K may be neglected, and Eqn. 17 approaches the center of gravity of the population.

A similar model has been proposed earlier (Van Opstal and Van Gisbergen, 1989) to explain the results of micro stimulation and saccade weighted averaging to double stimuli.

The dynamic goal specified by Eqn. 17 serves as the input to the nonlinear local feedback model of the brainstem proposed by Jürgens et al. (1981) and recently applied by Walton et al. (2005). The kinematic nonlinearity determines the horizontal and vertical eye-velocity components by:

$$\dot{e}_{H,V}(t) = V_{pk} \cdot [1 - \exp(-\frac{M_{H,V}(t)}{M_0})]$$  
and  
$$M_{H,V}(t) = S_{H,V}(t) - \int_0^t \dot{e}_{H,V}(\tau) d\tau$$  (18)

where $M_{H,V}(t)$ is the dynamic motor error for the horizontal and vertical eye-movement components, given by the difference between the desired horizontal/vertical eye displacement of the goal, and the actual horizontal/vertical eye displacement since the saccade onset, encoded by resettable integrators in the feedback loop. Parameters $V_{pk}$ and $M_0$ determine the amplitude-peak velocity nonlinearity of the main sequence. Typical values are $V_{pk}=700$ deg/s and $M_0=8$ deg. To obtain straight oblique saccades and component stretching, the horizontal and vertical burst generators would need to be coupled (see Introduction and above), but we have not incorporated this feature here. Instead, we limited the simulations to horizontal saccades.

**Figure 11:** Simulations with the center-of-gravity model (Eqns. 17-18). (A) Phase plots for cells with different optimal amplitudes are not straight. Inset: brainstem burst generator nonlinearity. (B) The nonlinear model also fails to explain the invariant phase plots for fast (gray line) and slow (dotted, black) saccades. Insets: gamma bursts for fast and slow 10 deg saccades (left); dynamic burst generator nonlinearity (right). See text for explanation.
Figure 11A shows the results of simulations with this model for cells with different optimal saccade amplitudes, ranging from $R_0=2$ to 50 deg. The burst duration of the SC cells was taken constant at $\alpha_{Dur} = 4.5$ ms, and $K=5$ spikes/s. Note that although all curves increase monotonically, and reach approximately the same number of spikes for the optimal saccade, they are clearly not straight (cf. Fig. 10A).

The experimental data of Goossens and Van Opstal (2006) indicated that the phase relationships are also preserved for strongly perturbed, slow saccadic eye movements (e.g. Fig. 10B). To test the nonlinear vector-averaging model for this property we simulated lower firing rates in the SC for a site encoding a 10 deg rightward saccade, by varying burst-duration parameter $\beta$ (Eq. 15) between 0.01 and 0.25 ms/deg, where high values for $\beta$ correspond with low SC firing rates. Since the number of spikes in the bursts was kept constant between 18-20 spikes (in line with the experimental data, and Fig. 6A, top), the burst durations increased accordingly, as the peak-firing rates dropped (Fig. 11B, left-hand inset). Note that without any changes in the properties of the brainstem burst generator, saccades would maintain the same kinematics regardless the SC burst profiles, as the dynamic goal signal of Eqn. 17 is virtually unaffected by the SC firing rates. However, since it is well established that lower levels of SC activity tend to correlate with slower saccades (see Introduction), we incorporated the assumption of Sparks and coworkers (Lee et al., 1988, Nichols and Sparks, 1996), that the asymptote of the burst generator ($V_{pk}$ in Eq. 18) co-varied with the SC firing rates according to $V_{pk}(\beta)=V_{pk}/(1+\beta)$ (Fig. 11B, inset lower right). However, despite this imposed kinematic relationship, the phase plot between the cumulative spike count of the SC cell and the actual eye displacement was not invariant to the temporal changes in the SC burst.

To summarise, the vector averaging scheme cannot capture the full repertoire of SC firing rate vs. eye movement correlations, simply because that model denies a direct collicular role in motor control.

**Conclusion**

From the simulations in this paper we conclude that the tight relationship between SC firing rates and saccadic eye displacement for both fast and very slow blink-perturbed saccades can only be explained if the nonlinear main sequence is embedded in the SC spatial-temporal activity patterns. We believe that our recordings, exemplified by the data in Fig. 6, together with these simulations indicate that the saccade-related burst in the SC specifies a *motor* command, rather than the spatial goal for the upcoming saccade.

The mechanism by which the SC incorporates the main sequence, and hence acts as a nonlinear vectorial pulse generator, appears to be by a gradient in the burst properties across the motor map, from brief and intense in the rostral zone, to less intense and of longer duration in the caudal zone. Yet, the number of spikes for each cell’s optimal saccade remains constant across the motor map and the distribution of firing rates within the active population follows a symmetrical, Gaussian profile.

Passive mechanisms like neural saturation or fatigue are not required to explain the observed results. Rather, the organization as described in
this paper could result from an optimization principle that allows saccades to acquire the target as fast and as accurately as possible, given the conflicting constraints within the system.

The question as to which mechanism ensures that for a given cell the number of spikes, and their temporal distribution, depends on the actual saccade amplitude, and hence on the neuronal input (Eqs. 13-15), will have to be explored in future studies.

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