

Synchronized Firing among Retinal Ganglion Cells Signals Motion Reversal

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SUMMARY

We show that when a moving object suddenly reverses direction, there is a brief, synchronous burst of firing within a population of retinal ganglion cells. This burst can be driven by either the leading or trailing edge of the object. The latency is constant for movement at different speeds, objects of different size, and bright versus dark contrasts. The same ganglion cells that signal a motion reversal also respond to smooth motion. We show that the brain can build a pure reversal detector using only a linear filter that reads out synchrony from a group of ganglion cells. These results indicate that not only can the retina anticipate the location of a smoothly moving object, but that it can also signal violations in its own prediction. **We show that the reversal response cannot be explained by models of the classical receptive field and suggest that nonlinear receptive field subunits may be responsible.**

INTRODUCTION

In order to initiate coordinated movements, the brain must compensate for both delays in the responses of neurons and delays in the movement of limbs to their intended targets. In principle, the only way to compensate for these delays is to use the past trajectory of an object's motion to make predictions about its future location. Such predictions are commonplace in everyday life, and even more salient examples come from athletics, such as a batter hitting a baseball or a wide receiver acrobatically catching a pass. Evidence of motion extrapolation exists for a variety of tasks: a batter tracking a fast-moving pitch (De Lucia and Cochran, 1985; Land and McLeod, 2000), human subjects hitting a moving object (Brouwer et al., 2003; Smeets and Brenner, 1995), extrapolating a trajectory (Pavel, 1990; Pavel et al., 1992), or tracking an object with smooth-pursuit eye movements (Barnes and Asselman, 1991; Becker and Fuchs, 1985; Robinson, 1965).

Animals also exhibit forms of motion extrapolation both in their eye movements (Klam et al., 2001; Medina et al., 2005) and in predictive firing of neurons in cortex and cerebellum (Fukushima et al., 2002; Heinen and Liu, 1997; Kettner et al., 1997; MacAvoy et al., 1991; Suh et al., 2000). Previous work has even shown that the retina makes a contribution to predicting the location of a moving object (Berry et al., 1999).

What happens, though, when the trajectory of a moving object suddenly changes in an unpredictable fashion? In sports, this leads to errors in motor coordination, such as a batter's swing missing a curveball when a fastball was expected, or a football player bobbling a fumbled football despite its slow movement. During smooth pursuit, sudden deviations in the trajectory lead to pursuit errors, and large localization errors often trigger catch-up saccades (Carpenter, 1988; Krauzlis and Lisberger, 1994; Medina et al., 2005; Robinson, 1965). One of the most profound motion discontinuities is a reversal of direction, which makes any previous extrapolation of the object's trajectory misleading. Motion reversal has long been known to evoke a characteristic change in EEG recordings from human subjects (Clarke, 1972; MacKay and Rietveld, 1968). These visually evoked potentials show peaks at 135–170 ms and ~260 ms after a reversal, and their source has been localized to the MT+ region of human cortex (Ahlfors et al., 1999). One experiment in monkeys suggested that single cells in areas MT and LIP might respond to motion reversals (Maimon and Assad, 2006). However, there has been no evidence of a specific response to motion reversal in earlier cortex, let alone subcortical structures.

Here, we show that motion reversal triggers an extra burst of firing in retinal ganglion cells. This firing occurs with a constant latency (~250 ms in salamander, ~190 ms in mouse), regardless of cell type or the location of the object in a cell's receptive field. As a result, the reversal evokes synchronized firing from a large population of ganglion cells, and this synchrony can uniquely distinguish the firing event from the response of the same cells to smooth motion. We suggest that this synchronized burst signals a violation in the retina's ongoing motion prediction. Such an error signal could be used by the brain to redirect attention or to reset central mechanisms of motion

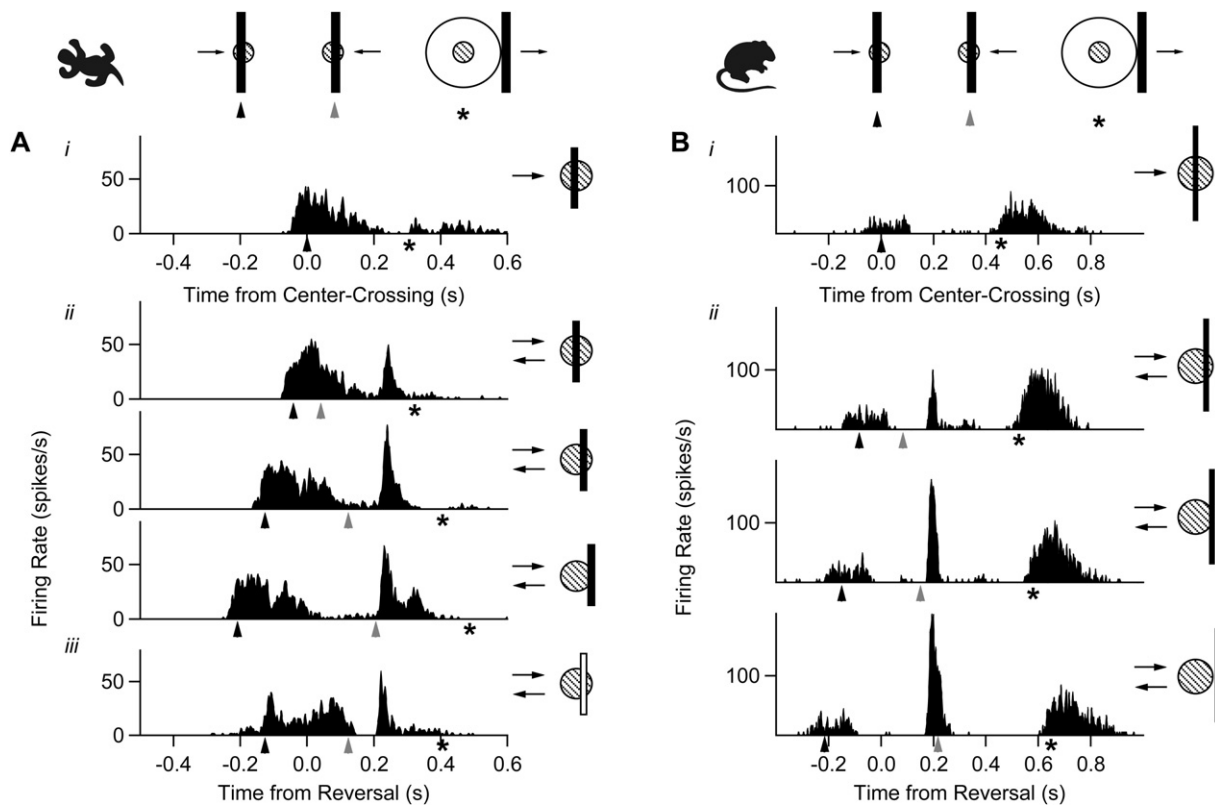


Figure 1. The Retina Responds Explicitly to Motion Reversal

(A) A salamander ganglion cell's response to a dark bar moving across the receptive field (*i*), a dark bar reversing at three different locations (*ii*), and a light bar reversing at one position (*iii*). The position of the bar relative to the receptive field center at time zero is shown by a pictogram (right). As shown by the key (top), arrowheads mark the time that the leading edge of the bar crossed the center of the receptive field (solid arrowhead) and the time it recrossed the same point after the reversal (gray arrowhead). Some cells, especially in the mouse, had a late response after the bar left the receptive field surround (asterisk, used for exits both to the right and left of the surround), which occurred regardless of whether there was a motion reversal or not.

(B) A ganglion cell from mouse responding to the same moving dark bars.

extrapolation. We also find that the synchronized burst of firing helps the retinal representation of a moving object's location rapidly catch up to the object's true position. This acceleration is a consequence of the retinal representation switching from one edge of the object to the other after the reversal.

RESULTS

We used a multielectrode array to record spikes extracellularly from retinal ganglion cells in salamander and mouse (Segev et al., 2004) while presenting a variety of moving objects as stimuli (see Experimental Procedures). A moving bar (110 μm wide, traveling at 1.32 mm/s) evoked a strong, temporally distributed response as it moved through a ganglion cell's receptive field (Figure 1, top). However, when the bar reversed direction near the cell's receptive field (pictogram to the right), an additional sharp peak of firing was observed (Figure 1, bottom). While the cell's initial, distributed response was timed to the arrival of the bar on the

receptive field center (solid arrowheads), the sharp burst of firing following the reversal was not timed to the reentry of the bar on the receptive field center (gray arrowheads). Rather, the response occurred at a fixed latency (~ 250 ms in salamander, ~ 190 ms in mouse) after the bar reversed, regardless of reversal position. We therefore conclude that this response was triggered by the reversal itself. This response was seen in a large proportion of cells in both species (salamander: 278/745 = 37%; mouse: 17/39 = 43%). Many cells, especially those in the mouse, also fired spikes when the bar moved out of the surround (asterisk). The strength of this "shift effect" varied considerably within the population and did not correlate with the response to motion reversal (Barlow et al., 1977; McIlwain, 1966).

The reversal response appeared at a fixed latency not only for different reversal positions around a single cell, but also across different cells. Figure 2 shows spike time rasters for five cells in salamander and six in mouse responding to a bar reversing at one position (see pictograms). Because each cell had a different receptive field

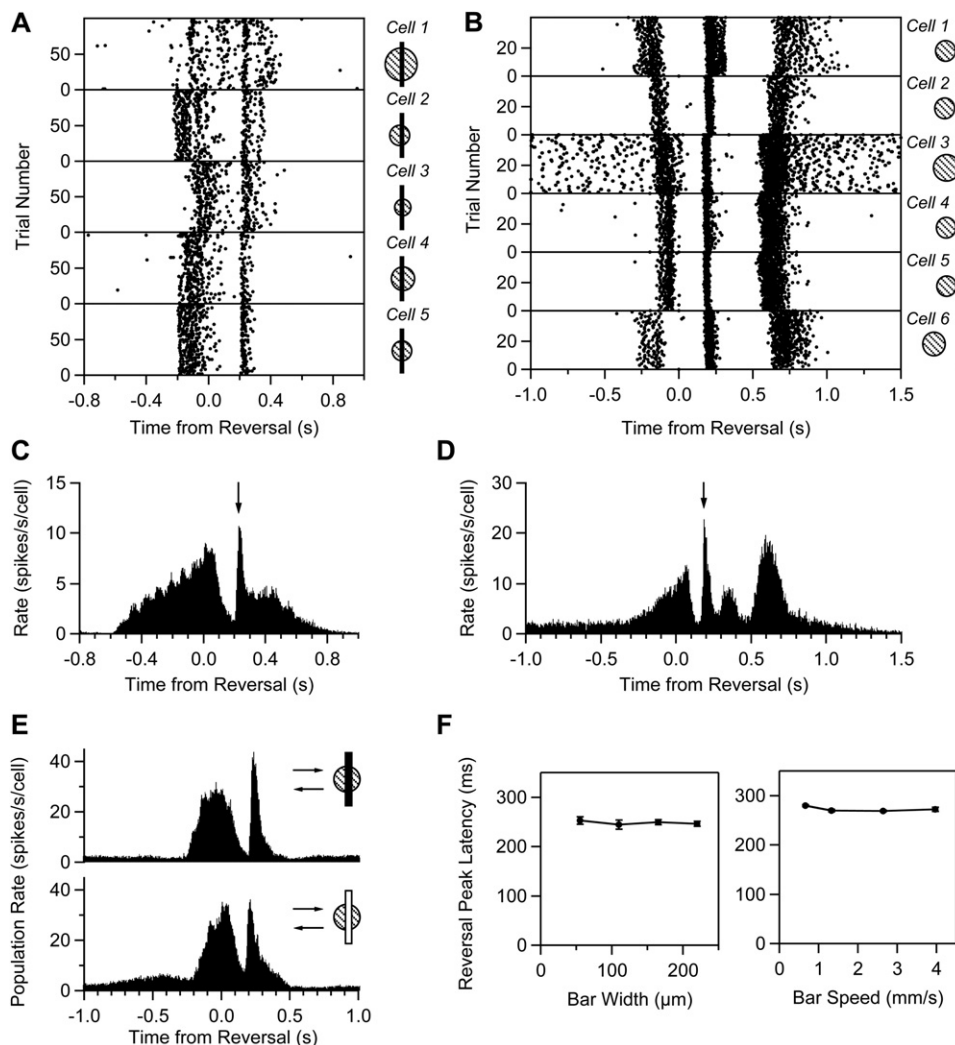


Figure 2. The Reversal Response Is Synchronous across Ganglion Cells

(A and B) Spike times (black dots) from five cells in salamander (A) and six cells in mouse (B) simultaneously recorded as a dark bar reversed near their receptive field centers (shown by pictograms to the right).

(C and D) Population average firing rate from multiple cells and reversal locations: (C) salamander, $n = 15$ cells at 9 positions each; (D) mouse, $n = 8$ cells at 11 positions each; reversal peak shown by arrow. The firing rates for each reversal location are aligned by the reversal time. In the salamander, the broad shoulder in the firing rate around ~ 400 ms is a smooth motion response from the bar reentering the receptive field. In the mouse, the peak in firing at ~ 350 ms is a smooth motion response, and the peak at ~ 600 ms is from the bar leaving the surround.

(E) Population average firing rate from 16 salamander cells responding to a dark bar (top) and a light bar (bottom) at the same reversal positions.

(F) Latency of the reversal peak as a function of bar width ($n = 11$ cells, salamander, left) and bar speed ($n = 13$ cells, salamander, right). Error bars are standard error of the mean (where visible).

center location, the initial response began at a different time. However, the reversal response was synchronized across all cells. The synchrony was even more apparent in the entire population (Figures 2C and 2D). When we aligned responses to the reversal time and calculated the average firing rate for a set of cells responding to different reversal locations, the distribution of initial firing due to the bar entering the receptive field was smeared, but the reversal response remained a sharp peak. Even large changes in the stimulus parameters failed to change the latency of the reversal response. Reversing bars of

opposite contrast, varying widths, and different speeds all elicited a response peaked at roughly the same latency (Figures 2E and 2F).

We next studied how the strength of the reversal response depended on the location at which the bar reversed on a cell's receptive field. When we measured the location of the bar's leading edge at the time of reversal, we found that reversals occurring before the cell's center coordinate had a strength that roughly followed the spatial profile of the receptive field center. However, reversals occurring beyond the center coordinate elicited

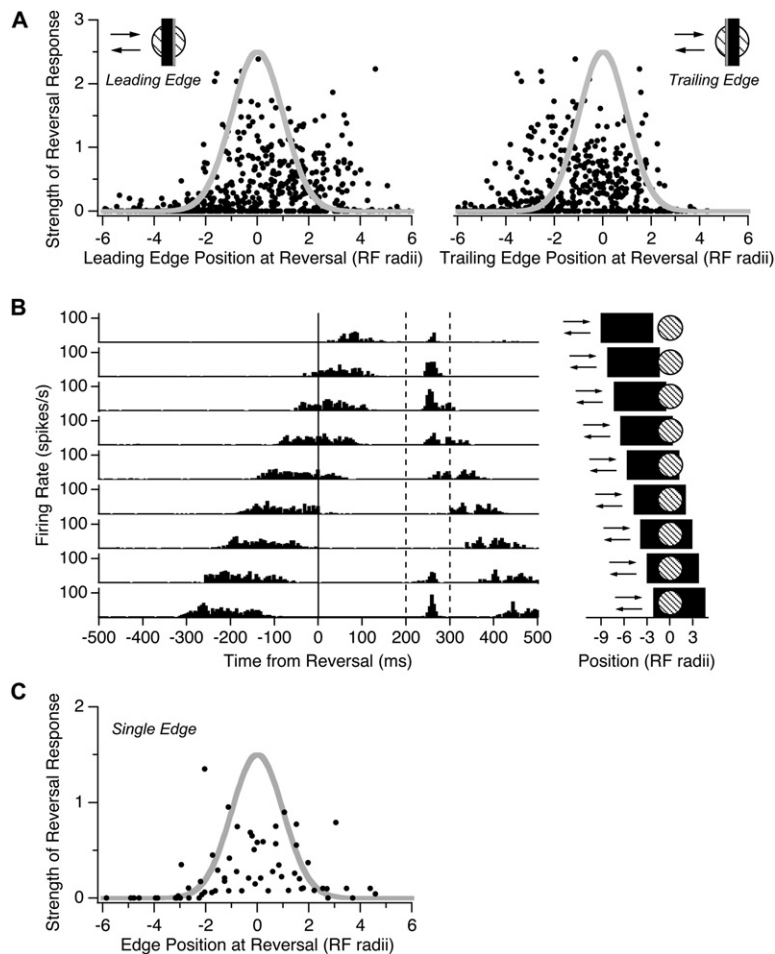


Figure 3. Dependence of the Response on Reversal Location

(A) (Left) Peak firing rate of the reversal response normalized by the peak firing rate during smooth motion and plotted against the distance of the bar's leading edge from the receptive field center coordinate, x_i , at the time of reversal ($n = 108$ cells, 338 total positions). Distances are expressed in units of the center radius of each cell, σ_i . (Right) Same data but with distances measured with respect to the location of bar's trailing edge at the time of reversal.

(B) Firing rate of a single ganglion cell during motion reversals of a wide ($440 \mu\text{m}$) bar at different locations (pictograms on the right). Dashed lines show the time window in which reversal responses were observed.

(C) Normalized peak firing rate at time of reversal plotted against the distance of a single-contrast edge from the receptive field center ($n = 8$ cells, 88 total positions).

responses from up to four center radii away (Figure 3A, left). The observation that both light and dark bars could trigger a reversal response in the same ganglion cell made us wonder if both the leading and trailing edge of the bar could drive firing after a motion reversal. If we instead considered the position of the bar's trailing edge, we found that responses for reversals beyond the center coincided with the spatial profile of the center (Figure 3A, right).

We investigated this effect further by using wider moving bars. Considered individually, many ganglion cells had a reversal response whose strength peaked around two different locations, corresponding to reversals of the leading and trailing edges. Figure 3B shows a cell with a peak in its firing rate at a fixed latency of ~ 250 ms after the reversal. This reversal response was elicited by the leading edge (top four traces) or trailing edge (bottom two traces), but no response occurred for reversals at intermediate positions. We repeated the analysis of Figure 3A for wider bars (220 and $440 \mu\text{m}$) and found the same asymmetry in the strength of the reversal response as function of the position of the leading edge at the time of reversal. However, when a single-contrast edge reversed its direction of motion, the strength of the reversal

response corresponded well with the spatial profile of the receptive field center (Figure 3C).

We also probed the retina with moving squares instead of extended bars. Motion reversal of a square object triggered a synchronous burst of firing in the ganglion cell population, just as for a reversing bar (Figure 4A). Because a square has a limited extent in the direction perpendicular to its motion, some reversals occurred in the surround above or below the center. These motion reversals did not trigger any firing (Figure 4B). For some reversal locations, the entire square was contained within the receptive field center and still triggered a burst of firing, indicating that stimulation of the surround is not necessary for a reversal response. Together, these observations suggest that synchronized firing can be generated when either the leading or trailing edge of a moving object reverses direction on the receptive field center but not on the surround.

Identifying Motion Reversals

Since the ganglion cells that respond to a reversing bar are not a small, specialized class, but rather a large fraction of the entire population, the brain faces a serious challenge in interpreting this message. Most spikes from

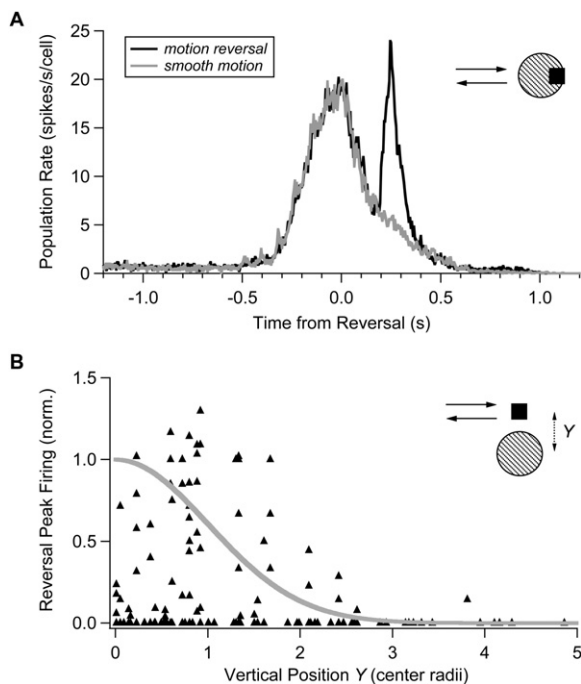


Figure 4. Ganglion Cell Responses to Reversing Squares

(A) Population average firing rate for responses to a smoothly moving square (gray; speed = 1.32 mm/s) and a reversing square (black; $n = 16$ cells at a total of 75 reversal positions).

(B) Peak firing rate of the reversal response as a function of the vertical distance, Y , between the center of the square and a ganglion cell's receptive field center coordinate, x_i , expressed in units of the center radius, σ_i . Firing rate was normalized by the peak firing rate for smooth motion of the square. Gaussian spatial profile is shown in gray.

one of these cells signal the smooth motion of an object across the cell's receptive field, while other spikes from the same cell have a very different meaning: an object has reversed direction. This ambiguity must be resolved

by looking at the population of ganglion cells. With this idea in mind, we constructed a decoder which used the spikes from a group of cells to distinguish reversals from smooth motion. We can think of the decoder as a "reversal-detector" cell in a subsequent neural circuit, which receives input from many retinal ganglion cells. We model this detector with a linear filter $f(t)$ that acts on all input spikes $\{t_i\}$, and a threshold, θ . Whenever the contribution summed over all inputs exceeds the threshold

$$u(t) = \sum_i f(t - t_i) \geq \theta,$$

the decoder decides that a reversal has occurred.

After computing the optimal filter from the data (see [Experimental Procedures](#)), we evaluated the performance of the decoder by detecting reversals in single-trial spike trains (Figure 5). We randomly selected many groups out of 31 cells recorded in one experiment, and applied the filter to all the spikes in the ganglion cell population for 120 trials each of smooth motion and motion reversals. Different values of the threshold θ result in a tradeoff between more misses versus more false alarms; we chose many values of θ and quantified the total error using the area under the receiver operator characteristic (ROC) curve (Green and Swets, 1966). Performance depended strongly on the spatial separation of ganglion cell receptive fields in each group (Figure 5D). This effect occurs because reversal synchrony was easier to detect when smooth motion triggered the cells to fire at different times. For widely separated groups of 16 ganglion cells, the total error rate was 4.2%.

Can this simple decoder achieve even better performance with more cells? Because reversals anywhere up to two radii beyond the center coordinate in either the horizontal or vertical direction can trigger a burst of firing, we estimate that over 275 cells are available to the decoder and at least 100 of them have significant reversal

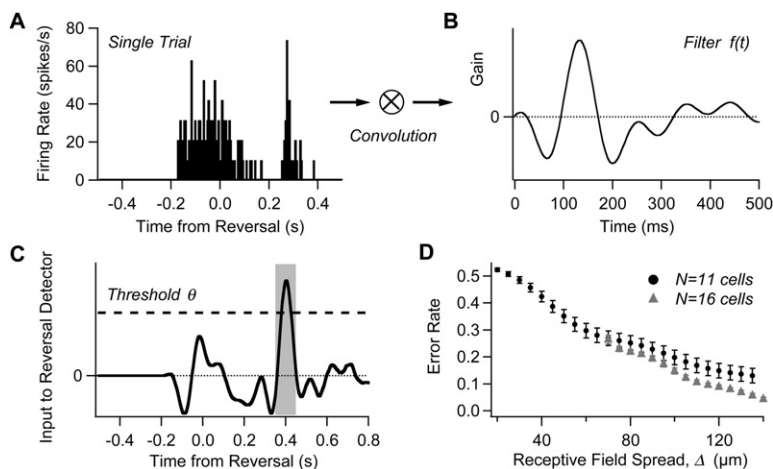


Figure 5. A Method for Detecting Motion Reversal Using a Population of Ganglion Cells

(A) The firing rate from a population of 12 cells on a single trial of motion reversal.

(B) Optimal filter, $f(t)$.

(C) Convolution of the firing rate from (A) with the filter from (B). Dashed line shows detection threshold. Shaded area represents target function for reversal detection, $\phi(t)$.

(D) Total error rate (from the area under the ROC curve) for detecting reversals among 120 reversal and 120 smooth motion trials plotted as a function of the ganglion cells' receptive field spread, Δ . Circles are for $n = 11$ cells, triangles are for $n = 16$ cells. Bars show the standard error of the mean compiled over many random selections of n cells. Chance value for a decoder that randomly selected reversal times is 95% error (see [Experimental Procedures](#)).

responses (see [Experimental Procedures](#)). To explore the performance possible with larger populations, we pooled over reversal locations, treating data from the same cell recorded at multiple reversal locations as multiple cells with receptive fields at different distances from point of reversal. Using 71 such cells with reversal responses, the decoder achieved 100% correct detection on all 120 reversal trials and 0% false alarms on all 240 nonreversal trials. While this analysis does not demonstrate that the brain actually does identify reversals in this manner, such identification is clearly possible using a biologically plausible mechanism and pooling over a realistic number of retinal ganglion cells.

Excellent detection is made possible not just by the synchrony of ganglion cell firing, but also by its temporal profile. If instead of filtering the spike trains with $f(t)$, we merely used the spike count in a single time window, the performance was very poor. For time windows between 5 and 40 ms, the spike count following the reversal exceeded all other time bins in no more than 26% of the trials. This means that error rates lower than 50% are not possible with a decoder that simply counts spikes in a single time window. The typical response of a ganglion cell involves initial firing due to smooth motion, followed by a pause, and then a sharp burst. The decoder's temporal filter has a shape resembling a second-derivative in time, which matches well to this temporal profile of ganglion cell firing. Of course, it may be possible to recognize a motion reversal using the *pattern* of cells that fire together in a small time window rather than just the total number of cells by performing a nonlinear operation on retinal spike trains, although such a decoding mechanism would be more biophysically elaborate than the form we propose.

Neural Image of a Reversing Object

We can gain more insight into the manner in which the retina tracks a moving object that reverses direction by considering the “neural image” of ganglion cell activity. The neural image is the spatial pattern of firing in the ganglion cell population as a function of time ([Berry et al., 1999](#)). We calculated the neural image for each point in time by first plotting the (normalized) firing rate of all the recorded ganglion cells as a function of the distance between the location of the moving bar and the cell's receptive field center coordinate ([Figure 6A](#)). Because we only recorded from a small fraction of all of the ganglion cells that respond to the bar's motion, we included data taken from the same cells at multiple reversal locations. These data were then smoothed to make our best estimate of the spatial pattern of activity in the larger population of ganglion cells (see [Experimental Procedures](#)).

Long before motion reversal, the neural image travels along with the moving bar, despite the response latency of ganglion cells ([Figure 6B](#)). This spatial shift in the neural image results from motion anticipation, which is a simple form of prediction that the retina makes about the future location of a smoothly moving object ([Berry et al., 1999](#)). Immediately after the reversal, the neural image over-

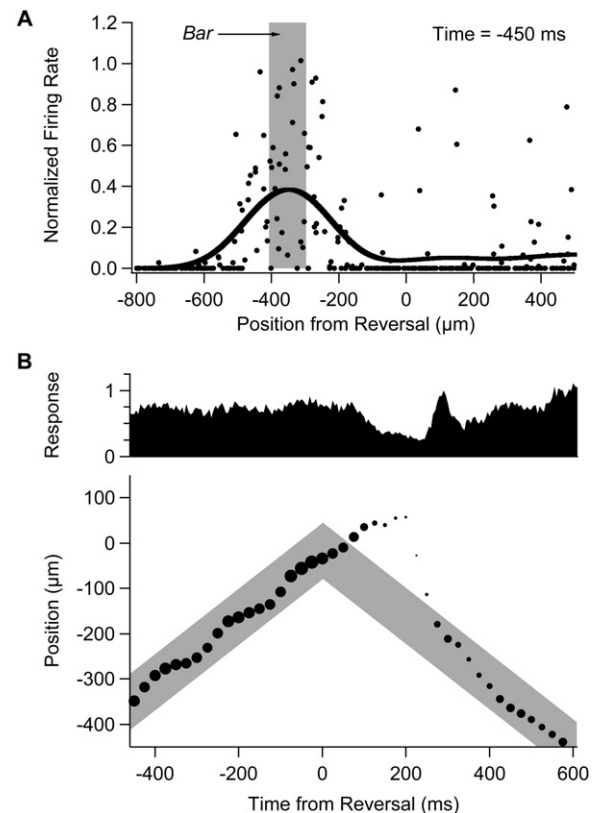


Figure 6. Neural Image of Retinal Activity

(A) Firing rate plotted against the distance between a ganglion cell's receptive field center coordinate, x_i , and the bar's reversal location (dots) along with neural image (black line). Firing rate values are normalized by the peak firing rate during smooth motion for each cell and are shown for a time 450 ms before reversal. The location of the moving bar is depicted by the shaded region.

(B) (Upper) Total response within the ganglion cell population plotted as a function of time from the motion reversal (normalized to unity at the time of the reversal response). (Lower) Location of the peak of the neural image (circles) plotted as a function of time from reversal; size of circles represents the peak firing rate; location of moving bar through time is depicted by the shaded region. (The slight oscillation in the location of the neural image during smooth motion is an artifact of sampling a discrete set of reversal locations. Note also that the peak firing rate [dots] need not track exactly the total response [upper panel] because of differences in the width of the neural image.)

shoots the reversal location. This is a manifestation of the predictive nature of retinal motion processing: because of its response delay, the retina does not yet know that a reversal has occurred and its previous anticipation of the object's location still operates. When information about the reversal becomes available, 100 ms later, the retinal response drops off. Then, at 250 ms after the reversal, the sharp, synchronous burst of firing occurs. Interestingly, the neural image rapidly accelerates in the new direction of motion at this time, nearly catching up to the bar's location. Retinal firing then briefly drops off, and finally, at 400 ms after the reversal, a smooth motion

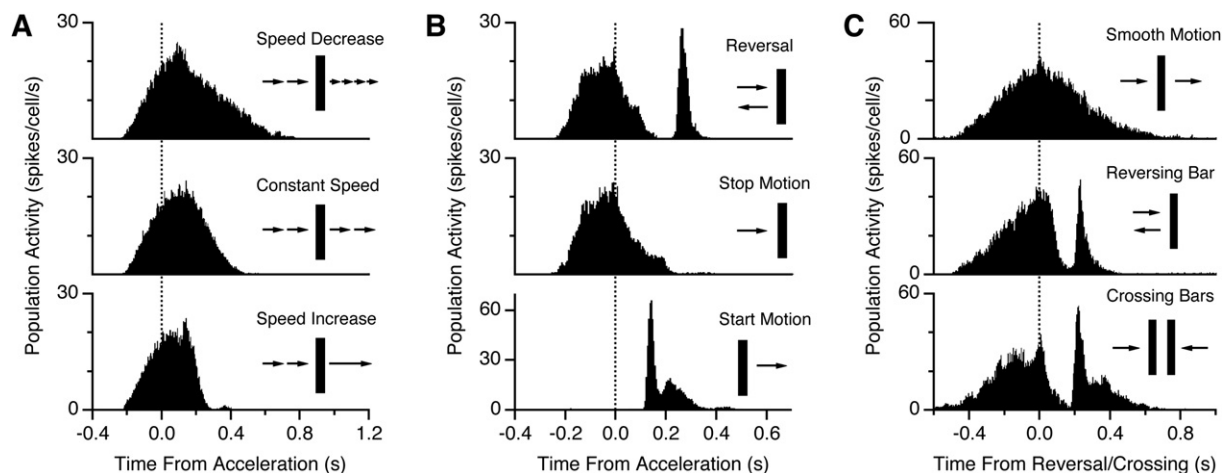


Figure 7. Response to Other Accelerations

(A) Population response of ganglion cells to motion that slows down by a factor of two (top), remains at the same speed (middle), or speeds up by a factor of two (bottom) without any change in direction.

(B) Population response to motion reversal (top), motion that abruptly stops (middle), or abruptly starts (bottom).

(C) Population response to a bar moving at constant velocity (top), a reversing bar (middle), or crossing bars (bottom).

Within each panel, data is taken from the same ganglion cells; across panels, data is taken from different retinas.

response emerges with correct anticipation of the bar's location. Again, the long delay before correct anticipation emerges is consistent with the interpretation that anticipation is a form of prediction that requires extensive integration over the object's past trajectory.

Why does the neural image catch up at the time of the synchronous burst of firing? This can be seen as a consequence of the spatial asymmetry of reversal locations that drive a response (Figure 3). When the leading edge of an object moves over a ganglion cell but the trailing edge is still within the receptive field center, the cell participates in the synchronous burst of firing. However, when the leading edge of an object reverses before it has reached the receptive field center, the cell does not fire. Therefore, when an object reverses, ganglion cells far away in the new direction of motion will fire but those far away in the old direction of motion will not fire. As a result, the neural image is significantly shifted in the new direction of motion, helping to catch up to the object's location. Another way of looking at the same data is to observe that during smooth motion, contrast gain control tends to localize ganglion cell firing near the leading edge of the object (Berry et al., 1999). But then, the reversal response can be triggered by either the leading or trailing edge. Thus, it helps to switch the location of the neural image to what will be the new leading edge of the object when it is moving in the other direction.

Generality of the Synchronized Response

Next, we asked whether this synchronized retinal response is unique to motion reversals or whether the retina conveys a more general signal of motion acceleration. To explore this question, we tested several other kinds of motion discontinuity. For both moderate acceler-

ation and deceleration in the same direction of motion, there was no extra burst of firing (Figure 7A), nor was there any extra firing for an object that abruptly stopped its motion (Figure 7B, middle). These data indicate that the response is not a generic signal of acceleration or motion discontinuity.

However, when a stationary object suddenly began moving, there was a synchronized burst of firing in the retinal population (Figure 7B, lower). While this firing pattern resembles the response to motion reversal, there are several salient differences: the latency is much shorter (140 versus 260 ms), the peak firing rate is roughly twice as large, and the latency depends systematically on the speed (data not shown). Furthermore, different sets of ganglion cells exhibit responses to motion onset and motion reversal. These facts suggest that different circuit mechanisms may be responsible for the response to motion onset, although a more unified picture may emerge from further studies. We also found that bars that moved at a constant velocity and crossed elicited a response that was nearly identical to that for a reversing bar (Figure 7C). So, a synchronized burst of firing is not unique to motion reversal. However, the synchronized firing patterns elicited by both motion onset and crossing motion possess important differences that distinguish them from the reversal response (see below).

Receptive Field Dynamics

In order to gain greater insight about the reversal response, we asked whether the spatiotemporal dynamics of a ganglion cell's receptive field could account for the phenomenon. Previous work has shown that ganglion cell responses to a smoothly moving bar are well predicted by a linear-nonlinear (LN) model with gain control

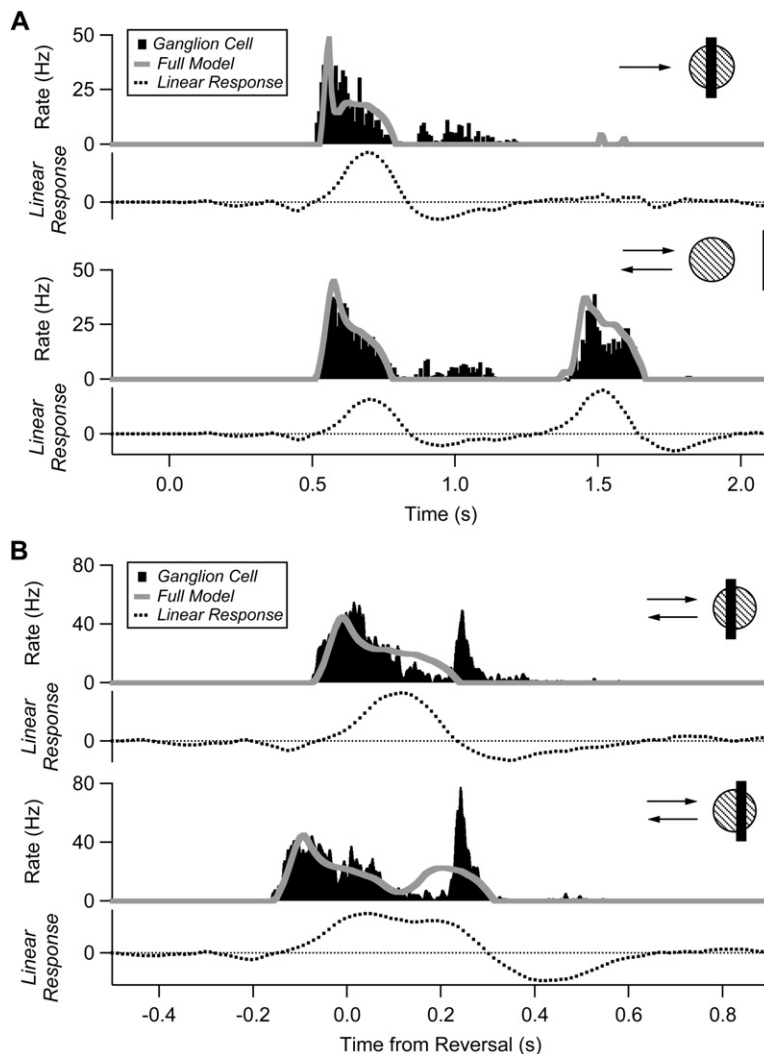


Figure 8. The Reversal Response Cannot Be Explained by a Classical Receptive Field Model

(A) Response of a single cell to smooth motion (top) and reversal far from the receptive field (bottom). Peristimulus time histogram (PSTH) is shown in black, the convolution of the linear receptive field and the stimulus is shown as a dashed line, and the best fit prediction from the LN model with gain control is shown as a gray line. Model parameters as in [Berry et al. \(1999\)](#); $\tau = 97$ ms; $\alpha = 360$ Hz; $\beta = 310$; $\theta = 0.043$.

(B) Same cell responding to reversals near the receptive field.

([Berry et al., 1999](#); [Shapley and Victor, 1981](#)). The linear part of the model describes the basic spatial and temporal filtering of the receptive field through a convolution operation ([Rodieck and Stone, 1965](#)). A nonlinearity then truncates negative values to produce a firing rate as a function of time. Adding gain control to this model enables it to mimic the ability of ganglion cells to anticipate the leading edge of a moving bar ([Berry et al., 1999](#)).

For each cell, the model was fit to firing rate traces for a bar which reversed far enough away from the receptive field center such that each pass of the bar elicited its own smooth motion response ([Figure 8A](#), lower). The same parameters were used for all other reversal locations (see legend). The model provided a good fit for responses to smooth motion, but failed to predict the reversal response for any reversal position ([Figure 8B](#)). The failure of the LN model lies in the initial spatiotemporal filtering, as this convolution often had a negative or zero value at the peak of the reversal response ([Figure 8B](#)). Across the population ($n = 85$ cells at 267 total reversal positions), the convolu-

tion value at the time of the reversal peak varied widely and on average was not significantly different than zero ($p = 0.87$). This analysis indicates that more sophisticated models of receptive field dynamics are needed to explain the reversal response (see below).

DISCUSSION

We have found that an abrupt reversal of motion evokes an extra burst of firing in a large fraction of all retinal ganglion cells in the salamander and mouse. This firing occurs with the same latency when either the leading or trailing edge of an object reverses on a cell's receptive field. Because the latency is constant for all cells and all reversal locations, and because either edge can drive a response, motion reversal evokes a synchronous burst of firing in a population of roughly 100 ganglion cells. To our knowledge, this mode of firing has not previously been described.

Circuit Mechanism

As shown in Figure 8, the reversal response cannot be explained by models of the classical receptive field, even if they include gain control. However, the reversal response embodies several kinds of invariance to properties of the moving object, such as polarity, size, and speed. These invariances are not only useful for the neural code, but also have implications for the circuit mechanisms responsible. The fact that both bright and dark objects can evoke a reversal response suggests that the retinal interneurons involved in this processing have ON-OFF response characteristics, as has been described in many types of amacrine cells (Miller et al., 2006; Pang et al., 2002; Sakai and Naka, 1987a, 1987b), or perhaps come in analogous ON and OFF populations, as found for bipolar cells (Ghosh et al., 2004; Pang et al., 2004) and starburst amacrine cells (Famiglietti, 1991; Zhang and Wu, 2001). We also found that the reversal response can be triggered entirely by a single-contrast edge. Presumably, these two facts are related: the trailing edge of a dark moving object locally resembles a bright moving edge. Thus, if the reversal response can be triggered separately and similarly for either edge, the latency would be independent of the object's size.

The fact that the response latency is roughly constant as a function of the speed of the moving object is challenging to explain, and places strong constraints on the circuit mechanism. One possibility is that the detection of a motion reversal involves the re-excitation of a retinal interneuron during the object's return path at a fixed time delay with respect to when it was first excited by the initial path of the moving object. If such interneurons tile the retina and ganglion cells pool over many such subunits, excitation could follow a motion reversal at the same latency for a variety of speeds, even if driven by a different set of subunits at each speed. Furthermore, the subset of ganglion cells that receive input from these interneurons would fire synchronously, as we observe. An appealing candidate for such an interneuron is the bipolar cell, or perhaps even the axonal terminal of the bipolar cell, as these cells are likely to serve as motion-sensitive subunits in the receptive fields of Y-type ganglion cells (Demb et al., 2001; Shapley and Victor, 1979; Victor, 1988). Why might the bipolar cell be re-excited at a fixed time delay? One speculation is that motion in the initial direction leaves inhibition in its wake, and that the interneuron requires a roughly fixed amount of time to recover from this inhibition before it can be excited again (Dong and Werblin, 1998; Roska et al., 1998).

The case of crossing bars is important because the spatiotemporal pattern of photoreceptor activation on either side of the crossing point, taken alone, is identical to that of reversing bars. The observation that crossing bars elicit nearly the same response as a reversing bar (Figure 7C) therefore implies that the computation of motion reversal must be a fairly localized operation, consistent with the re-excitation model. Our experiments involving moving bars of different widths have implications

for the spatial scale of the reversal computation. Both the leading and trailing edges of a relatively narrow moving bar can drive a reversal response (110 μm width, compared with a typical receptive field diameter of $\sim 250 \mu\text{m}$). But smooth motion of the same bar does not elicit separate responses from each edge. These data indicate that the receptive field of the reversal-sensitive subunits must be significantly smaller than the ganglion cell dendritic field and suggest that their size is less than $\sim 100 \mu\text{m}$. Again, these observations point to individual bipolar cells as a possible locus of reversal detection, although sharp excitation could also be generated by a narrow-field amacrine cell inhibiting a sustained amacrine cell and thereby transiently removing tonic inhibition from the ganglion cell.

The model of re-excitation of a retinal interneuron after recovery from inhibition also has implications for the locations that can trigger a reversal response. In this model, it is not the location at the time of reversal that determines the response. Instead, it is location of an edge at the time that interneurons recover from inhibition. Because of this additional delay, reversal locations somewhat beyond the boundary of the receptive field center can trigger a response via this mechanism. This may explain why we sometimes see quite strong responses at a distance of over two radii from the center coordinate. What is clear from our data is that the surround cannot directly generate a reversal response by itself, as there are several conditions that are not effective: (1) motion reversals that occur before reaching the center; (2) reversals with the trailing edge more than one surround radius (which is four to five center radii) past the center; and (3) reversals of a square above or below the center. Thus, all of our data is consistent with the idea that reversals are generated by the center mechanism, but possibly with a delay relative to the time of reversal.

Synchrony in the Neural Code

There are many ways that the ganglion cell population might encode a motion reversal. We find that the retina, rather than simply staying silent, sends a positive signal to the brain. The retina could contain a specialized class of ganglion cells that signal motion reversals. Instead, it employs the same ganglion cells that anticipate smooth motion and uses a distinct firing mode within the population: namely, a synchronized burst, which can be readily detected by subsequent neural circuits. Synchronized firing has been proposed as a special event in many neural populations (Abeles, 1991; Hatsopoulos et al., 1998; Meister, 1996; Vaadia et al., 1995), although the role of synchrony for population codes is an ongoing topic of research. This retinal reversal response is one of the few examples in which synchrony encodes a qualitatively different property of the world (Ishikane et al., 2005; Neuenschwander and Singer, 1996). By multiplexing the reversal signal onto optic nerve fibers using a synchrony code, the retina can send different kinds of visual messages using fewer optic nerve fibers (Meister, 1996).

What is the purpose of the retina's response to a motion reversal? During smooth motion, anticipation corrects for the retina's response latency, and as a result, the peak firing rate of a ganglion cell represents the object's true, current location (Berry et al., 1999). Immediately after a reversal of motion, the retina cannot anticipate the object's location, which can lead to large localization errors. The synchronized burst of ganglion cell activity can therefore serve to identify a violation in the retina's ongoing prediction of a moving object's location.

Why might the brain need to know that retinal motion prediction has been violated? One intriguing possibility comes from a consideration of how an animal coordinates its motor output with moving objects in the environment. Motion anticipation corrects for neural delays in the retina itself, but this degree of anticipation is not sufficient for coordinated movement. There are additional neural delays in sensory-motor pathways downstream of the retina, and there are motor delays for limbs to reach their intended locations. Thus, subsequent neural circuits in the brain may need to perform additional extrapolation of a moving object's trajectory. Evidence from human psychophysics indicates that in many behavioral contexts, both hand and eye movements embody such extrapolations (Barnes and Asselman, 1991; Brouwer et al., 2003; Land and McLeod, 2000; Pavel et al., 1992; Smeets and Brenner, 1995).

Although the neural mechanisms that underlie central motion extrapolation are not known, the only way, in principle, to make such an extrapolation is to use the past trajectory of an object's motion. However, when there is a sudden reversal of motion, the entire preceding trajectory will be not just irrelevant, but actually misleading about the future location of the moving object. As a result, the central mechanism of motion extrapolation would benefit from being "initialized," so that it can begin to accumulate a prediction of the object's future location that uses only the relevant portion of the trajectory. Perhaps the synchronous burst produced by the retina following a motion reversal may play a role in initializing central motion extrapolation.

Different Kinds of Synchronized Firing

Why might the retina respond to motion onset and reversal but not to other accelerations at the speeds we tested? One possibility is that less severe accelerations lead to a small enough error in the retinal representation of the object's location such that no explicit violation signal is needed. For instance, when motion suddenly stops, the neural image overshoots the actual position of the object (Figure 6), but this overshoot is less than 100 μm , which is smaller than the receptive field size of an individual ganglion cell. In contrast, when the object reverses its direction of motion, the localization error at the time of the reversal response is more than 250 μm compared with the center of the bar. As the size of the retina's localization error should depend on the speed of motion as well as the magnitude of the acceleration, it will be fruitful to analyze

the neural image following a variety of different motion discontinuities.

Another related possibility is that synchronized firing is only needed to initialize mechanisms of central motion extrapolation. In the case of motion onset, such central mechanisms would not be engaged prior to the discontinuity, but for other kinds of mild acceleration, they would already be operating. In this context, it is important to consider how the brain might specifically recognize the appropriate motion discontinuities. What the brain needs is a detector that is triggered by either motion onset or motion reversal, but not by smooth motion or other mild accelerations. In fact, if retinal spike trains are imported into the same reversal detector as described above (Figure 5), we get excellent discrimination between motion onset and smooth motion (0 errors in 240 trials using all 71 cells). The decoder achieves the same performance for discrimination between motion that suddenly starts (detection) and motion that suddenly stops (no detection). Such a discontinuity detector could thus serve as the trigger for initializing central motion extrapolation mechanisms.

At the same time, this detector is unable to reliably distinguish motion onset from motion reversal. However, if we use the same form of decoding algorithm, but with a different linear filter that is optimized for distinguishing start from reversal, we can again achieve excellent performance (0 errors in 240 trials). This demonstrates that the ganglion cell population does in fact convey information about which kind of motion discontinuity has occurred, and it suggests that a more sophisticated decoding process, using either two successive simple stages or a single complex mechanism, can perform this discrimination.

A similar issue arises in how the brain might interpret the response to crossing bars. Although this response seems very similar to that for a reversing bar, the retina still provides information that can distinguish between crossing and reversing bars: the crossing bar stimulus will generate synchronized firing in two populations of ganglion cells on either side of the crossing point, while a simple reversal will only generate synchronized firing in a single group of cells offset from the reversal point in the new direction of motion. Thus, the brain can discriminate between reversal and crossing by pooling over an even larger population of ganglion cells.

These analyses indicate that synchronized firing in the population of retinal ganglion cells has the potential to encode qualitatively different events in the visual world. While the connection between synchronized firing and central mechanisms of motion extrapolation is highly speculative, it may help to make sense of this surprising aspect of the retinal code as well as motivate further experiments.

EXPERIMENTAL PROCEDURES

Recording

Pieces of retina obtained from larval tiger salamanders (*Ambystoma tigrinum*) and mice (strain C57/BL6) were perfused continuously with

oxygenated Ringer's medium. Ganglion cell spikes were recorded extracellularly from a multielectrode array. Salamander recordings were made at room temperature, and mouse recordings were made at 36°C. Details of the recording and spike sorting are described elsewhere (Segev et al., 2004).

Visual Stimulation

Visual stimuli were presented on a computer monitor running at 120 Hz (Puchalla et al., 2005). Moving bars were presented on a gray background and were 110 μm wide, traveling at 1.32 mm/s unless otherwise noted. Bars reversed at 9–15 locations near the receptive fields of the ganglion cells with 55 or 110 μm separations between locations. Parameters of other stimuli are noted in figure legends. Moving squares were 165 \times 165 μm in size and also traveled at 1.32 mm/s. Squares reversed at a grid of 7 \times 7 locations over the multielectrode array.

Receptive Fields

Spatiotemporal receptive fields were measured by reverse correlation to random flicker presented at 60 Hz. In experiments with moving bars, we mapped receptive fields with flickering black and white strips 22 μm wide and oriented parallel to the moving bar stimuli. In experiments with moving squares, we mapped receptive fields with flickering squares 55 μm on a side. A 1D or 2D Gaussian was fit to the spatial profile of the center to identify a center coordinate x_i and center radius σ_i for each cell i [29]. The uncertainty in the center coordinate ranged from 0.95 to 7.6 μm with an average of 2.3 μm ; the uncertainty in the center radius averaged 3.3 μm . Thus, retinal locations measured in units of center radii had a combined uncertainty of $\sim 3\%$ (Figure 3 and Figure 4). The spread of receptive fields in a group of cells, Δ , was defined as the standard deviation of the set of center coordinates, $\{x_i\}$. For making pictograms, we displayed a circle with a size given by the boundary of the receptive field center. The boundary was defined as the point at which the spatial profile changed polarity.

Reversal Response

Cells were classified as reversal responsive if they had a peak in their firing rate, as measured by the peristimulus time histogram (PSTH), that was greater than 10 Hz between 200 ms and 300 ms after the reversal in the salamander and between 150 ms and 250 ms in the mouse. We chose this relatively conservative criterion to exclude cells with low firing rates and cells with receptive fields too far from the location of motion reversals. We also required that the latency of the reversal response remained constant for at least three different reversal positions. This was to make sure that we did not mistake a smooth motion response that happened to come at roughly the right time for a reversal response. Our criteria are likely to disqualify some cells that actually respond to motion reversal, mostly because we only sample a discrete number of reversal locations, so our estimate of the fraction of cells participating is a lower bound. All population averages and decoding analysis include only reversal responsive cells; figure legends indicate whether population averages include all tested locations for reversal responsive cells (Figure 2) or only locations with a firing rate greater than 10 Hz (Figure 4).

Decoder

Reversals were detected using a decoder consisting of a linear filter applied to spikes from all the ganglion cells, followed by summation and a threshold operation. The optimal filter $f(t)$ is a function of time that, when convolved with the average population firing rate accumulated over all trials, $r(t)$, most closely resembles a target function $\phi(t)$, which represents the detection of the reversal event following the reversal response. The target function was chosen to have a value of one for a 100 ms window beginning 350 ms after the reversal, and

zero everywhere else (Figure 5C). We solved for this condition in the frequency domain:

$$\tilde{f}(\omega) = \tilde{\phi}(\omega) / \tilde{r}(\omega)$$

where

$$\tilde{f}(\omega)$$

is the Fourier transform of $f(t)$.

For correct detection, the decoder had to exceed threshold within the time window defined by the target function. All other threshold crossings were counted as errors (false alarms). In addition, any threshold crossings that occurred during other stimulus trials, such as smooth motion, were counted as false alarms. Because we measured error rates on a per trial basis, we allowed no more than one false alarm per trial. As false alarms could occur anywhere over a 1 s period in each trial (except for the 100 ms wide target region), the error rate for a decoder that selected random times would be 90% for reversal trials only and 95% for reversal and smooth motion trials combined. For purposes of cross-validation, the optimal filter was constructed using half of the data, then tested on the other half, and vice versa.

Pooling over Many Cells

If we assume that reversal responses can arise from locations up to two radii away on either side of the center coordinate in either the horizontal or vertical direction (Figure 3 and Figure 4), then there is a circular region of approximately two center radii that can sense the reversal of even a very small object. Assuming a center radius of 125 μm and a density of 1400 cells/mm² in salamander (Segev et al., 2004), there are ~ 275 ganglion cells in this area. If 278/745 = 37% of them have a reversal response, then the decoder can pool over ~ 100 cells. Reversal of larger objects would engage even more ganglion cells.

Neural Image

We first calculated the PSTH over 75 stimulus trials for each ganglion cell in 25 ms bins. Then, for each time step, we plotted each cell's firing rate versus the distance between its receptive field center coordinate, x_i , and the location at which the moving bar reversed direction. The firing rate was normalized by each cell's peak firing rate in response to smooth motion. Data were pooled over 9–15 different reversal locations and smoothed with a Gaussian filter that had a width of 20 μm (Figure 6A). For each point in time, the location and amplitude of the peak activity of the neural image was found (Figure 6B). In these experiments, the neural image was found to be a unimodal distribution at all points in time, so that its peak was unambiguous.

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