



Published in final edited form as:

J Comp Physiol A Neuroethol Sens Neural Behav Physiol. 2010 September ; 196(9): 613–628. doi:10.1007/s00359-010-0547-z.

Calcium-dependent control of temporal processing in an auditory interneuron: a computational analysis

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Abstract

Sensitivity to acoustic amplitude modulation in crickets differs between species and depends on carrier frequency (e.g., calling song vs. bat-ultrasound bands). Using computational tools, we explore how Ca^{2+} -dependent mechanisms underlying selective attention can contribute to such differences in amplitude modulation sensitivity. For omega neuron 1 (ON1), selective attention is mediated by Ca^{2+} -dependent feedback: $[\text{Ca}^{2+}]_{\text{internal}}$ increases with excitation, activating a Ca^{2+} -dependent after-hyperpolarizing current. We propose that Ca^{2+} removal rate and the size of the after-hyperpolarizing current can determine ON1's temporal modulation transfer function (TMTF). This is tested using a conductance-based simulation calibrated to responses in vivo. The model shows that parameter values that simulate responses to single pulses are sufficient in simulating responses to modulated stimuli: no special modulation-sensitive mechanisms are necessary, as high and low-pass portions of the TMTF are due to Ca^{2+} -dependent spike frequency adaptation and post-synaptic potential depression, respectively. Furthermore, variance in the two biophysical parameters is sufficient to produce TMTFs of varying bandwidth, shifting amplitude modulation sensitivity like that in different species and in response to different carrier frequencies. Thus, the hypothesis that the size of after-hyperpolarizing current and the rate of Ca^{2+} removal can affect amplitude modulation sensitivity is computationally validated.

Keywords

Adaptation; Temporal processing; Roex filter; Evolution; Temperature

Introduction

Shifts in auditory filtering may be accomplished by changing intrinsic biophysical properties of single units (Fortune and Rose 1997; Zakon 2003; Hildebrandt et al. 2009). For example, the frequency of electrical resonance in certain hair cells is determined by the kinetics of the BK conductance, with the expression of different BK splice variants in different hair cells shifting the resonant band across the papilla (Art et al. 1995). Beyond the cells of a single papilla, however, intrinsic biophysical filters can be employed in different dendritic compartments of single auditory cells (adjusting response to different inputs) or auditory cells in different organisms, producing changes in sensitivity for different species and hybrids (Zakon 2003). This paper examines (1) how an intrinsic mechanism can contribute to the

different temporal filters exhibited in response to mating versus predator sounds in an identified auditory cell in crickets; and (2) if the mechanism would be capable of producing the shifts in sensitivity found between species.

In crickets (Gryllidae), amplitude-modulated calling songs are produced by repeated closure of the forewings (Bennet-Clark 1989). Song amplitude modulation structure is species specific, meaning temporal sensitivity plays an important role in mediating mate choice (Walker 1957; Pollack and Hoy 1979; Shaw 2000; Shaw and Herlihy 2000). At least one mechanism for auditory temporal filtering in this system is that underlying selective attention, where sensitivity and phonotaxis are directed to only one of several otherwise attractive songs (Pollack 1986; Pollack 1988). Neural correlates of selective attention have been shown in omega neuron 1 (ON1), a pair of intrasegmental auditory units in the prothoracic ganglion. ON1s are post-synaptic to auditory receptors and, in part, transfer input to contralateral ascending neurons to enhance binaural contrast (Selverston et al. 1985; Horseman and Huber 1994). When presented simultaneously with a pair of calling songs of different amplitude, the response of ON1 is biased toward the higher amplitude song, even though both songs elicit excitation when presented alone (Pollack 1988). Such selectivity is mediated by the cell's mechanisms for adaptation (cf. Sah and Faber 2002; Bond et al. 2005): excitation increases intracellular $[Ca^{2+}]_i$, activating an outward, hyperpolarizing conductance and subsequently decreasing the response to a maintained stimulus (Sobel and Tank 1994; Baden and Hedwig 2007). This feedback mechanism shifts the cell's I/O curve to higher amplitude stimuli, with the lower amplitude song now below threshold on the curve. The result is selective attention that is sensitive to both the amplitude and the timing of the elements of the two songs, with songs occurring during "after-hyperpolarization" (ahp) less likely to elicit a response. The mechanism is thus a form of temporal filtering (Harris and Dallos 1979; Frisina 2001): filtering pulses that are too close together in time (Sobel and Tank 1994; Farris et al. 2004; Marsat and Pollack 2004; Baden and Hedwig 2007).

Several lines of evidence suggest that the temporal properties of selective attention in ON1 are specialized for processing amplitude modulation. For example, ON1 responses differ for calling song and ultrasound inputs (Sabourin et al. 2008). In particular, ON1 band-pass filters the amplitude modulation of sounds with lower carrier frequencies (i.e., near calling songs) to slow amplitude modulation rates, but exhibits broadly tuned low-pass filtering of amplitude-modulated ultrasound (i.e., bat-like sonar; Marsat and Pollack 2004). Although this can partly be explained by receptor (presynaptic) sensitivity (Sabourin and Pollack 2010), mechanisms in ON1 (postsynaptic) are clearly implicated, as excitation-induced ahp is enhanced for song-like carrier frequencies compared to that generated by ultrasound input (Pollack 1988). Importantly, this difference in ahp may be due to dendritic tonotopy in internal Ca^{2+} ($[Ca^{2+}]_i$): ultrasound stimuli elicit less $[Ca^{2+}]_i$ in certain dendritic compartments than calling songs do (Baden and Hedwig 2007). In addition, Tunstall and Pollack (2005) showed that sensitivity of ON1 to amplitude modulation in a slow-singing species is more narrowly tuned to slower modulation rates than that for a fast-singing species. Furthermore, the extent of adaptation was greater in the more narrowly tuned species. Taken together, these data suggest that processing of amplitude modulation in ON1 is influenced by the properties of Ca^{2+} -dependent feedback, which may be specialized to increase the binaural processing of functionally relevant sounds (Marsat and Pollack 2004, 2005).

Thus, we use a computational model based on omega neurons in Gryllids (Sobel and Tank 1994; Farris et al. 2004; Baden and Hedwig 2007) to first show that simulating the Ca^{2+} -dependent mechanisms of selective attention is sufficient to simulate the *in vivo* responses to amplitude-modulated stimuli. Second, we test the prediction that variance in these mechanisms is sufficient to produce differences in temporal filtering. In particular, a biophysical parameter

analysis shows that sensitivity to amplitude modulation can be shifted (similar to the effect of different carrier frequencies) by varying the size of the Ca^{2+} -sensitive ahp conductance and the rate of Ca^{2+} removal. In addition, the model allows for these Ca^{2+} -dependent shifts to be compared to those produced by changing the speed of spike-generating mechanisms using temperature. Third, from a functional point of view, we extend the interpretation of the results beyond carrier-dependent filtering, as the model provides evidence that variance in these mechanisms could contribute to general shifts in temporal sensitivity, including those in ON1 that are found between species (Tunstall and Pollack 2005).

Methods

Although measurements of ON1 from several different laboratories are available, they are mostly collected in current clamp or through Ca^{2+} imaging. We are not aware of any voltage clamp approaches with ON1, making few data available for certain biophysical components. Whether the components are taken from ON1 or other cells is thus noted. However, parameter values were calibrated to fit experimental data (see below).

A modified Hodgkin and Huxley (1952; Huguenard and McCormack 1994) conductance-based model simulated the membrane potential of ON1 in response to afferent input. The model integrates voltage and Ca^{2+} -sensitive conductances to produce the membrane response:

$$C \frac{dV}{dt} = - (I_{\text{Na}} + I_{\text{K}} + I_{\text{Ca}} + I_{\text{ahp1}} + I_{\text{ahp2}} + I_{\text{Leak}}) + I_{\text{stim}}. \quad (1)$$

The current components are: spike-generating sodium (I_{Na}) and potassium (I_{K}); voltage-sensitive calcium (I_{Ca}); two Ca^{2+} -sensitive after-hyperpolarizing currents (I_{ahp}); a leak and the stimulus current. C is membrane capacitance.

Spike-generating conductances (I_{Na} and I_{K}) were treated as generic and based on characteristics of vertebrate interneurons (Traub et al. 1991; McCormick and Huguenard 1992). Parameter values were ultimately chosen, however, to model action potentials recorded from ON1 in vivo (Farris et al. 2004; see below). For all parameters, their units, initial value of variables and value of constants, see Table 1.

The simulated currents and gating variables are as follows. The rate of change of the activating and inactivating variables is described by a first-order differential equation:

$$dx/dt = (x_{\infty} - x) / \tau_x \quad (2)$$

where τ_x represents the time constant of the activating or inactivating variable, x ,

$$\tau_x = 1 / (\alpha_x + \beta_x) \quad (3)$$

and α_x and β_x are gating functions shown with each ion's conductance.

Sodium current, I_{Na}

$$I_{\text{Na}} = \bar{g}_{\text{Na}} \times m^3 \times h \times (V - E_{\text{Na}}) \quad (4)$$

$$m_{\infty} = \alpha_m / (\alpha_m + \beta_m) \quad (5)$$

$$\tau_m = \tau_1 / (\alpha_m + \beta_m) \quad (6)$$

$$\alpha_m = 0.32 (13 - v_2) / (\exp((13 - v_2)/4) - 1) \quad (7)$$

$$\beta_m = 0.28 (v_2 - 40) / (\exp((v_2 - 40)/5) - 1) \quad (8)$$

$$h_{\infty} = \alpha_h / (\alpha_h + \beta_h) \quad (9)$$

$$\tau_h = \tau_1 / (\alpha_h + \beta_h) \quad (10)$$

$$\beta_h = 4 / (1 + \exp((40 - v_2)/5)) \quad (11)$$

$$\alpha_h = 0.128 \exp((17 - v_2)/18), \quad (12)$$

with \bar{g} being the conductance of the ion; m and h representing the activation and inactivation, respectively (Fig. 1); and v_2 the voltage relative to spike threshold (for both I_{Na} and I_K):

$$v_2 = V - (v_{\text{threshold}}) \quad (13)$$

Potassium current, I_K

$$I_K = \bar{g}_K \times n^4 \times (V - E_K) \quad (14)$$

$$n_{\infty} = \alpha_n / (\alpha_n + \beta_n) \quad (15)$$

$$\tau_n = \tau_1 / (\alpha_n + \beta_n) \quad (16)$$

$$\beta_n = 0.5 \exp((10 - v_2)/40) \quad (17)$$

$$\alpha_n = 0.032 (15 - v_2) / (\exp((15 - v_2)/5) - 1) \quad (18)$$

with n representing activation.

The temperature control was maintained over the sodium and potassium currents by the following equation:

$$\tau_1 = Q_{10}^{(C-36)/10} \quad (19)$$

where C is temperature ($^{\circ}\text{C}$). Q_{10} was set at 1.5 in all the simulations (Machne and Orozco 1970; McCormick and Huguenard 1992). All simulations were run using 25°C unless noted otherwise.

Calcium current, I_{Ca}

The calcium current is modeled after those recorded in cockroach (*Periplaneta americana*) interneurons (Wicher and Penzlin 1997) and the L-type Ca^{2+} current in cricket (*G. bimaculatus*) myocytes (Mutoh and Yoshino 2004; Numata and Yoshino 2005) (Fig. 1). The current is assumed to be uniform across all areas of the cell and is represented by the Goldman–Hodgkin–Katz current equation (Hille 1992):

$$I_{\text{Ca}} = \bar{P}_{\text{Ca}} \times m_{\text{Ca}}^2 \times \Gamma(V, [\text{Ca}^{2+}]_i, [\text{Ca}^{2+}]_o, 2) \quad (20)$$

where \bar{P}_{Ca} is the permeability of the calcium channel and m_{Ca} is the voltage-activating variable ranging from 0 to 1,

$$m_{\text{Ca}} = 1 / [1 + \exp(-(V+13)/8.6)] \quad (21)$$

Γ uses the typical Goldman–Hodgkin–Katz parameters to describe the driving force as a function of voltage and internal and external Ca^{2+} ($[\text{Ca}^{2+}]_i$ and $[\text{Ca}^{2+}]_o$) (this notation previously used by Engel et al. 1999). The valence of Ca^{2+} equals 2.

With respect to the $[\text{Ca}^{2+}]_i$, the omega neuron has at least two distinguishable compartments with relatively fast and slow change in $[\text{Ca}^{2+}]_i$: whereas the $[\text{Ca}^{2+}]_i$ in the dendritic areas decreases after excitation with a time constant (τ_{removal}) near 237 ms, the $[\text{Ca}^{2+}]_i$ in the spike generation zone exhibits slower changes, $\tau_{\text{removal}} = 1265$ ms (Baden and Hedwig 2007). When taken together with the fact that ON1 spike frequency adaptation exhibits a double exponential decay (Fig. 2), it suggests that these different Ca^{2+} dynamics contribute to different rates of Ca^{2+} -activated adaptation. Thus, based on the differing rates of Ca^{2+} extrusion in different anatomical compartments and the double exponential spike frequency adaptation found in vivo, the model contains two pools of internal Ca^{2+} : $[\text{Ca}^{2+}]_{i,\text{pool1}}$ and $[\text{Ca}^{2+}]_{i,\text{pool2}}$ (the notation ‘i’ is not an index and represents internal Ca^{2+}). The modeled pools are functionally independent with respect to their associated Ca^{2+} -activated after-hyperpolarizing conductance, or I_{ahp} (i.e., a compartmentalization of $[\text{Ca}^{2+}]$ and associated Ca^{2+} -activated currents). This separation allows for the separate contributions of two Ca^{2+} -activated after-hyperpolarizing conductances (Wang 1998; but see Roper et al. 2003). It is important to note, however, that $[\text{Ca}^{2+}]_i$ used to calculate I_{Ca} in the Goldman–Hodgkin–Katz current equation above is the sum of all $[\text{Ca}^{2+}]_i$. That is, $[\text{Ca}^{2+}]_i = [\text{Ca}^{2+}]_{i,\text{pool1}} + [\text{Ca}^{2+}]_{i,\text{pool2}}$. Thus, although there are two pools that affect

an associated $I_{\text{ahp}1}$ or 2, there is a simplification to only one pool that affects the driving force of I_{Ca} .

Ca²⁺ change and removal

The rate of Ca²⁺ entry and exit in each pool is described by a first-order differential equation,

$$\frac{d[\text{Ca}^{2+}]_{\text{internal},j}}{dt} = -\alpha_{\text{Ca},j}I_{\text{Ca}} + \left(\frac{-[\text{Ca}^{2+}]_{\text{internal},j}}{\tau_{\text{removal},j}} \right) \quad (22)$$

where j is an index that designates the pool, $j = 1, 2$; α_{Ca} is a constant chosen to scale $[\text{Ca}^{2+}]_i$ so that it varies across the steep portion of the I_{ahp} calcium activation curve (Fig. 1). The relative size of $\alpha_{\text{Ca},1}$ and $\alpha_{\text{Ca},2}$ reflects the surface area to volume ratio of the two pools: $\alpha_{\text{Ca},2} = \alpha_{\text{Ca},1}/3$. Although the value of this relationship was previously chosen (Wang 1998) to help describe the different Ca²⁺ modes of the soma and dendrites of cortical pyramidal neurons, it is also used here because it satisfactorily modeled that for the proposed location of the two pools in ON1: the spike generation zone and dendrites of ON1 (Baden and Hedwig 2007). $\tau_{\text{removal},j}$ determines the rate at which $[\text{Ca}^{2+}]_{i,j}$ changes in each pool. The removal represents the functional result of all mechanisms of Ca²⁺ removal: binding, sequestration and extrusion.

Ca²⁺-activated after-hyperpolarizing current, $I_{\text{ahp},j}$

The currents are modeled after voltage-independent SK conductances (Engel et al. 1999; Faber and Sah 2007) and control spike frequency adaptation and input resistance by generating hyperpolarizing current in response to $[\text{Ca}^{2+}]_{i,j}$ in one of the two compartments (j):

$$I_{\text{ahp},j} = \bar{g}_{\text{ahp},j} \times ms_j^2 \times (V - E_{\text{K}}) \quad (23)$$

where $\bar{g}_{\text{ahp},j}$ is the maximum conductance and ms is the sensitivity of I_{ahp} activation to $[\text{Ca}^{2+}]_i$:

$$ms_j = \left(\frac{[\text{Ca}^{2+}]_{i,j} / \text{Ca}_{\text{half_activation}}}{1 + \left(\frac{[\text{Ca}^{2+}]_{i,j} / \text{Ca}_{\text{half_activation}}}{1} \right)} \right), \quad j=1, 2 \quad (24)$$

The open–close action of the ahp (c.f. SK) channel is fast compared to the calcium concentration change, so the dynamics of the ahp channel is described by a function instead of a differential equation. Although Ca²⁺-activated K⁺ channels have been identified in insects (McCormack 2003) and other invertebrates (Wei et al. 1996; Burrell and Crisp 2008), most measures of their Ca²⁺ sensitivity are from vertebrate cells. For those, the reported half activation concentrations for SK currents vary (Leinders and Vijverberg 1992; Tucker and Fettiplace 1996; Xia et al. 1998; Hirschberg et al. 1999; Stocker et al. 2004). In our simulations $\text{Ca}_{\text{half_activation}} = 2.5 \mu\text{M}$ (Engel et al. 1999), but its actual numeric value is irrelevant to the model, as $[\text{Ca}^{2+}]_i$ has been scaled to vary across the non-saturating portions of the I_{ahp} activation curve (ms) using α_{Ca} (Fig. 1). This is a requirement, as in vivo recordings of ON1 show that variance in $[\text{Ca}^{2+}]_i$ elicits variance in adaptation (Sobel and Tank 1994; Baden and Hedwig 2007), which would not occur if $[\text{Ca}^{2+}]_i$ saturated I_{ahp} sensitivity (Roper et al. 2003).

Leak currents, I_{Leak}

The leak current is the sum of each ion's (Na^+ , K^+ , Ca^{2+}) leak, which takes the form of the Goldman–Hodgkin–Katz current equation (Hille 1992; Huguenard and McCormack 1994):

$$I_{\text{Leak}} = \sum I_{\text{Leak,ion}} \quad (25)$$

$$I_{\text{Leak,ion}} = \bar{P}_{\text{ion}} \times \Gamma(V, \text{ion}_i, \text{ion}_o, \text{valence}). \quad (26)$$

The model was simulated using the Runge–Kutta method in XPP software (<http://www.math.pitt.edu/~bard/xpp/xpp.html>).

Calibration of model parameters

Model parameters were calibrated using ON1 responses to a single acoustic pulse in nine female *Gryllus rubens* (Farris et al. 2004). This species was chosen because the male calling song consists of a long-duration series of pulses (trill), making the amplitude modulation structure relatively simple in that there is only one interpulse interval (11 ms duration pulses at 45 pulses/s at 22°C; Walker 1962; Bentley and Hoy 1972). Adaptation of the ON1 spike trains were produced in response to five repetitions (2 s stimulus interval) of a 400 ms duration pulse of 5 kHz (90 dB SPL). In simulating these spike trains, very little change was required for the voltage-sensitive parameters (e.g., parameters of I_{Ca} , I_{Na} and I_{K}) taken from vertebrate neurons, roach neurons and cricket muscles; these parameters were “tuned” by eye and remained the same for all nine simulations of each cell’s spike frequency adaptation. The five parameters that contribute to adaptation, however, were systematically adjusted using a least-squares fit procedure between the real and simulated spike frequencies. The amplitude of I_{stim} , an idealized square pulse, was calibrated first to a value that generated the appropriate initial spike frequency (instantaneous frequency of the first five spikes). Subsequently, $\bar{g}_{\text{ahp},1}$ and $\tau_{\text{removal},1}$ were adjusted to model the spike frequency change of the initial short time constant. Lastly, the same procedure was then used to calibrate parameter values for the mechanisms producing the slower time constant for change in spike frequency ($\bar{g}_{\text{ahp},2}$ and $\tau_{\text{removal},2}$). Whereas starting parameter values in this least-squares method were arbitrarily chosen for $\bar{g}_{\text{ahp},j}$, starting values for $\tau_{\text{removal},1}$ and $\tau_{\text{removal},2}$ were the Ca^{2+} decay times measured in the dendritic (237 ms) and spike generation zone (1265 ms) of ON1 in *G. bimaculatus*, respectively (Baden and Hedwig 2007).

With respect to adaptation, it is important to note that studies in another cricket auditory interneuron (AN1) have shown that the reduced coding of fast amplitude modulation rates during sinusoidal input (but not randomly modulated input) may not be due to the adaptive shift in the input/output (I/O) curve, but instead to a subsequent compression of the I/O curve itself: the spike frequency dynamic range of the shifted curve becomes smaller, reducing the mean spike frequency and thus limiting the modulation rates that spikes can copy (Benda and Hennig 2008). In contrast, however, for ON1 in vivo, neither the adapted nor the maximum spike frequencies explain the shape of the TMTF at high-amplitude modulation rates (Marsat and Pollack 2004); and adapted I/O curves appear to shift to the right, but not compress (Sobel and Tank 1994). This is illustrated by the fact that although the mean adapted firing rate to amplitude-modulated ultrasound stimuli is lower (20–30 Hz less) than that for calling song stimuli, its TMTF extends to higher amplitude modulation rates (Marsat and Pollack 2004). Furthermore, ON1 spike rate often remains greater than the behaviorally relevant amplitude modulation rates (Nabatiyan et al. 2003; Baden and Hedwig 2007). Thus, our model makes a

parsimonious assumption as to the effect of adaptation and does not incorporate any special terms to reduce the maximum adapted firing rate (Benda and Hennig 2008). The model uses only the known mechanisms of selective attention: excitation-induced ahp. This mechanism simply shifted the I/O curve to the right, rather than compressing it (Fig. 3). Such a shift without compression would not be unique to ON1, as it has been found in other auditory interneurons (Dean et al. 2005, 2008).

Stimulus and spike train analysis

The amplitude-modulated stimulus was a periodic square wave at nine rates (1–120 Hz), with amplitude alternating from 0 to I_{stim} in each cycle (3 s total duration). Although synaptic and receptor filtering are implicated in ON1 responses (Marsat and Pollack 2004; Sabourin and Pollack 2010), this was chosen as an idealized input so that our analysis focused solely on filtering due to the post-synaptic mechanisms of the model and not the input itself (Faulkes and Pollack 2001). The stimulus had a 50% duty cycle, meaning response variance depends only on the rate of stimulation, not on overall energy. The amplitude of I_{stim} was calibrated to the mean voltage that best simulated the responses of ON1 to 90 dB stimuli ($n = 9$ cells; see results).

Although several methods have been used to assess response to amplitude-modulated stimuli in crickets (Hennig 2003; Nabatiyan et al. 2003; Farris et al. 2004; Marsat and Pollack 2004), we employed two methods similar to those previously shown to be successful in analyzing ON1 temporal sensitivity in vivo. The first method tests whether the number of peaks in instantaneous spike frequency varied between stimuli with different amplitude modulation rates (Nabatiyan et al. 2003). Thus, the dependent variable for the filter is the number of spike frequency peaks (SFP) per stimulus pulse. The threshold for discriminating an SFP was 200 Hz, the same as that used for ON1 in vivo (Nabatiyan et al. 2003). The limitation of this method, however, is that it fails to record high-pass filtering well, as low-frequency amplitude-modulated stimuli still elicit SFPs even though adaptation causes a failure to code their long-duration pulses. Thus, this method yielded temporal modulation transfer functions that were low pass in shape. This low-pass filter was analyzed using a single-sided rounded exponential function (see below).

The second method of analysis takes a spectral approach (e.g., Tunstall and Pollack 2005). Voltage responses are first converted to instantaneous spike frequency, $r(t)$. Sub-sequently, in lieu of bins, we used a linear interpolation (50 μ s sampling period) variant of the discontinuous fft (Fan and Liu 2004) to transform $r(t)$ into the spectral domain: $r(t)$ is first converted to a continuous function, $R(t)$, and then analyzed for spectral content using the fft transform. This generates the power in the spike frequency waveform at the stimulus modulation frequency. The transfer function is then plotted using the signal-to-noise ratio in the spectrum at each stimulus frequency: $10 \cdot \log(\text{SNR})$, where $\text{SNR} = \text{response power at the stimulus modulation frequency} / \text{spectral density in } \pm 1 \text{ octave}$. This method has no limitation on calculating high and low-pass components and delivered band pass like TMTFs.

The flanks of this band-pass filter (i.e., on either side of the best amplitude modulation frequency) or the single flank of the low-pass filter produced by the peak counting method above were analyzed with the rounded exponential function, $roex(p,r)$:

$$W(g) = (1 - r)(1 + pg) \exp(-pg) + r \quad (27)$$

where g is the normalized deviation of frequency from the center frequency, cf , $g = |(f - cf)|/cf$; p is a dimensionless parameter determining the slope of the flanks of the TMTF, and r is a

constant that sets the range limitation of the roex filter. The TMTFs were assumed to be asymmetrical around the center amplitude modulation frequency, and separate slopes (p) were calculated for the upper and lower frequency flanks; r was limited to be the minimum measure on either side of the cf. When there was no high-pass component, only the low-pass component was calculated. The model was fit using a least-squares method to solve for the slope parameter (p) of each side of the peak frequency. From these parameters, the equivalent rectangular bandwidth (ERB) of the filter can be calculated (Hartmann 1998), facilitating comparisons of filtering properties across taxa (Patterson et al. 1982; Moore et al. 1988; Farris and Hoy 2002; Witte et al. 2005; Farris et al. 2006).

Note that we did not use methods measuring synchrony (Farris et al. 2004; Marsat and Pollack 2004) for the following reason. When I_{stim} elicited a spike, it was often in the same stimulus phase. Thus, even when many stimulus pulses were not coded, those that were coded produced high measures of phase locking, resulting in very little variance in the synchronization coefficient.

All filter parameters were solved using the Minerr function in MathCad (PTC, Needham, MA, USA). All means are reported as \pm standard deviation.

Results

Adaptation to a single pulse

For a single acoustic stimulus (400 ms duration; 5 kHz), spike frequency adaptation of ON1 (in vivo) in *G. rubens* exhibited a double exponential decay (Fig. 2): compared to the single exponential fit, the double exponential fit explained more of the change in spike frequency (R^2 single 0.527 ± 0.078 ; R^2 double 0.655 ± 0.085 ; $n = 9$ cells). There was a consistently large difference between the slow ($\tau_{slow} = 1054.3 \pm 470.2$ ms) and fast ($\tau_{fast} = 23.2 \pm 13.2$ ms) time constants for the double exponential fit, as the within cell ratio was $41.3 (\pm 0.01)$. Thus, we assume that multiple processes with different rates contribute to adaptation and therefore the model required a fast and a slow mechanism for its simulation. This was accomplished by having two separate Ca^{2+} removal times ($\tau_{removal,1}$ and $\tau_{removal,2}$) and two associated after-hyperpolarizing Ca^{2+} -activated conductances ($\bar{g}_{ahp,1}$ and $\bar{g}_{ahp,2}$) (Fig. 4).

During single pulse acoustic stimulation in vivo (Fig. 5), the maximum (i.e., initial) spike frequency was different for each cell (356 ± 68 Hz). Thus, simulations required different model stimulus amplitudes ($I_{stim} = 0.074 \pm 0.034$ mA/cm²; used in all multipulse experiments below). This variance in peak spike frequency notwithstanding, the extent of adaptation was relatively constant for all cells ($83 \pm 8\%$), meaning no cells used to calibrate the model exhibited evidence for saturation of the adaptation mechanisms. In simulating the single pulse response (Fig. 5), we found that Ca^{2+} removal times ($\tau_{removal}$) for the two Ca^{2+} pools were similar to the two exponential time constants measured above, generating mean removal times of $\tau_{removal,1} = 34.4 \pm 11$ ms and $\tau_{removal,2} = 1,265$ ms. There was no variance for $\tau_{removal,2}$ because the least-squares fit showed little or no improvement for changes to the initial seed value. Thus, this value was the same for all nine cells.

The size of the \bar{g}_{ahp} conductances (and thus I_{ahp}) differed by an order of magnitude for the two Ca^{2+} pools, with $\bar{g}_{ahp,1} = 0.047 \pm 0.02$ S/cm² and $\bar{g}_{ahp,2} = 0.0053 \pm 0.003$ S/cm². This difference reflects the need to generate greater adaptation at the onset of the stimulus (Figs. 2, 4, 5). The mean parameter values generated from simulating the nine cells ($\tau_{removal,1}$; $\tau_{removal,2}$; $\bar{g}_{ahp,1}$ and $\bar{g}_{ahp,2}$) were then used to test whether parameters based on adaptation to a single acoustic pulse were sufficient to simulate the response to paired and multipulse stimuli.

Paired pulse selective attention

Similar to the response found in vivo to paired pulse acoustic stimuli (Sobel and Tank 1994), a preceding pulse affects the response to a subsequent one in simulations (Fig. 6). Shown to be a mechanism of selective attention (Pollack 1988), responses to the second of a pair of current pulses are reduced when the second pulse is delivered during the after-hyperpolarization elicited by the preceding pulse (Fig. 6). The model captures well the shift of the linear I/O curves of spike number per pulse (Sobel and Tank 1994) and reveals that the mechanisms of adaptation generate a temporal filter, as stimuli occurring close together in time (i.e., faster amplitude modulation) are less likely to elicit full responses.

Multipulse stimuli: temporal modulation transfer function

$[Ca^{2+}]_{i, pool}$ exhibits the fastest dynamics and is considered here to functionally represent the faster Ca^{2+} dynamics of the dendrites (Baden and Hedwig 2007). Thus, the hypothesis that the broader TMTF for ultrasound is caused by less dendritic calcium-induced ahp can be tested by comparing coding of amplitude modulation when $\bar{g}_{ahp,1}$ is reduced relative to the mean parameter value. Whether coding is analyzed using SFP or the spectrum of the spike frequency waveform, the model confirms the hypothesis, as adjusting $\bar{g}_{ahp,1}$ affects amplitude modulation coding. The TMTFs generated from the spike frequency waveforms (e.g., Fig. 7) for three different values of $\bar{g}_{ahp,1}$ are shown in Fig. 8. In vivo, dendrites coding ultrasound show ~47% of the $[Ca^{2+}]_i$ compared to those responding to 5 kHz (Baden and Hedwig 2007). The TMTF of the model dendrite coding ultrasound (47% of the mean $\bar{g}_{ahp,1}$) is a broader filter, as both the high-pass and low-pass portions are affected (low-pass cutoff frequency increased to 46.5 Hz; Fig. 8; red curve). In contrast, when the mean value (or greater) for $\bar{g}_{ahp,1}$ is used, the TMTF is band limited and more narrow with a lower amplitude modulation cutoff frequency of 26.1 Hz (Fig. 8; black and blue curves). As shown in other cells (Benda and Herz 2003; Benda and Hennig 2008), the source of the high-pass filter is spike frequency adaptation (Fig. 3). However, we do not have compression of the I/O curve in ON1. Thus, exploration of the source of the low-pass filter shows that it results from PSP depression (Fortune and Rose 1997, 2001; Rose and Fortune 1999). When I_{ahp} is on there is a change in input resistance, shunting I_{stim} (Fig. 8). This means the membrane's RC response is changed:

$$\frac{V_{out}}{V_{in}} = \frac{1}{\sqrt{1 + \omega^2(RC)^2}}$$

R is the membrane resistance, RC is the time constant and ω is the radian input frequency. When simulations use greater $\bar{g}_{ahp,1}$, there is more I_{ahp} , reducing input resistance and reducing the coupling between I_{stim} and PSP. Thus, R functionally increases (i.e., the sub-threshold passive response to inward current is diminished), increasing the time constant and thus lowering the bandwidth of response (Getting 1974; Horowitz and Hill 1989; Hartmann 1998). With respect to amplitude modulation, longer duration pulses are now required to overcome the slower integration. Furthermore, faster amplitude modulation increases the number of pulses occurring when I_{ahp} is on. Consequently, increases in I_{ahp} ($\bar{g}_{ahp,1}$) or the duration it is activated imposes greater low-pass filtering of amplitude modulation (Fig. 9). Although the other Ca^{2+} -dependent parameters were also capable of shifting sensitivity in both amplitude modulation cutoff frequency and bandwidth, the parameters affecting the fast component ($\bar{g}_{ahp,1}$ and $\tau_{removal,1}$) were slightly more effective, having the largest effect on filter bandwidth (ERB; Fig. 9; Table 2).

Overall, these results show that calibrating the model to the single pulse response in vivo was sufficient to simulate selective attention (i.e., paired pulse) and the TMTF of this species.

Furthermore, the results show that the amplitude modulation sensitivity difference between ultrasound and calling song input can be simulated by adjusting the fast (i.e., dendritic) adaptation parameters; and such adjustments are sufficient to produce continuous changes to the TMTF of the cell.

Temperature and the TMTF

Temperature is only coupled to the model through the I_{Ca} , I_K and I_{Na} activation and inactivation parameters (and not I_{ahp}). Thus, change in temperature affects the TMTF response through effects on spike structure (Fig. 10). Using the mean values for the four adaptation parameters, the model shows that increases in simulation temperature caused a decrease in spike duration in the first spike, as well as those after the first adaptation time constant (i.e., spikes later than $\tau_{removal,1}$). There was no measurement after $\tau_{removal,2}$, as adaptation had eliminated spiking at this time under the mean parameter conditions. The TMTF shows a linear increase in the filter cutoff frequency with increasing temperature (slope = 1.11; intercept = 0.88). However, this change was much less than that found for singing males (Fig. 10).

Discussion

Cells may code aspects of stimulus amplitude, temporal structure and duration using intracellular Ca^{2+} accumulation. This mechanism often leads to spike frequency adaptation and PSP depression, as Ca^{2+} -activated K^+ conductances (e.g., SK) respond to the 'accumulation measure' by providing a form of feedback inhibition, reducing the input resistance of the cell and producing hyperpolarization (Wang 1998; Engel et al. 1999). A primary function of these effects is to reduce responses to background noise or low amplitude stimuli (Fig. 6; see also Pollack 1988; Sobel and Tank 1994; Benda et al. 2005). But, coding of repetitive or amplitude-modulated stimuli is also affected, causing responses to change at both ends of the modulation scale. We have shown in this computational study that variance in Ca^{2+} feedback mechanisms (ahp) can contribute to differences in temporal responses of ON1 (e.g., like those for different carrier frequencies; Pollack 1988; Marsat and Pollack 2004; Sabourin et al. 2008). Furthermore, we also found that these mechanisms can produce sufficient variance in TMTFs to function in the context of the evolution of interspecific differences in sensitivity.

Significance of multiple Ca^{2+} pools

Spike frequency adaptation in ON1 is well described by a double exponential decay (Fig. 2; see also Tunstall and Pollack 2005), necessitating the use of two ahp conductances and associated Ca^{2+} pools for adequate simulation. Integration of multiple adaptation rates within cells are not uncommon and often reflect the contribution of different ahp conductances expressing different activation characteristics or kinetics (Storm 1990). These separate pools, and thus adaptation characteristics, could be the result of different Ca^{2+} dynamics in different cellular zones. Baden and Hedwig (2007) showed that the rate of Ca^{2+} decay in ON1 dendrites and the spike generation zone differ by an order of magnitude. Our model suggests that with respect to adaptation, these two Ca^{2+} zones are functional, as integration of two similar Ca^{2+} removal times simulates adaptation well. If Ca^{2+} dynamics were separated in such a way as to limit or bias integration of one Ca^{2+} pool relative to the other (e.g., for different frequency inputs), the characteristics of adaptation would be changed, being carrier frequency specific. This seems to be the case in ON1 as Ca^{2+} imaging shows that the amount of internal Ca^{2+} in certain portions of the dendritic tree depends on stimulus carrier frequency (Baden and Hedwig 2007). When taken together with our model, we propose that such a separation could contribute to the different temporal response of ON1 for different carrier frequency inputs (Marsat and Pollack 2004). Supporting this hypothesis are data showing: (1) anatomical segregation of

afferent central projections from different frequency receptors (Esch et al. 1980) and the contralateral ON1; (2) ultrasound input does not generate large after-hyperpolarizations (Pollack 1988); (3) the input resistances of the two frequency inputs appear different, as integration time and recovery from adaptation suggest higher input resistance (less I_{ahp}) for high frequencies (Sabourin et al. 2008); (4) the synaptic input from receptors to ON1 differs between ultrasound versus low carrier frequency-sensitive receptors (Pollack 1994; Faulkes and Pollack 2001). In contradiction though, direct synaptic contact from low-frequency and high-frequency receptors to ON1 does not appear to be located in very different ON1 dendritic regions (Imaizumi and Pollack 2005). The hypothesis, however, does not require great differences in synaptic locations and could be mediated in microdomains between the post-synaptic zone and the spike generation zone. Indeed, stimulus-dependent separation of Ca^{2+} accumulation has been shown in the visual neuropil of flies (Egelhaaf and Borst 1995). It is important to note that this endogenous filtering hypothesis does not eliminate synaptic or pre-synaptic mechanisms (French 1986; Gollisch and Herz 2004; Marsat and Pollack 2004; Sabourin and Pollack 2010). But, taken together with the empirically derived Ca^{2+} -sensitive adaptation mechanisms (Sobel and Tank 1994; Baden and Hedwig 2007), the model does argue that variance in these Ca^{2+} -dependent parameters alone is sufficient to explain the differences in temporal code, as the model does not need intervening circuits or synaptic filters.

Implications for selective pressure and song coding

In orthopterans, auditory decisions, including species recognition, are based in part on the structure of amplitude modulation in calling songs, such that phonotaxis is elicited by a particular species-specific range of modulation rates (Walker 1957; Thorson et al. 1982; Schildberger 1984; Doherty 1985b; Hennig and Weber 1997; Schul 1998; Shaw and Herlihy 2000). Comparisons across taxa show that song modulation rate can evolve across a wide range, as interspecific variance is broadly distributed (Walker 1962) and exhibits evidence for quantitative underlying genetics (Shaw et al. 2007). Indeed, hybrids of species with different amplitude modulation rates produce intermediate songs: hybrid modulation rate is halfway between those of parental species. Interestingly, song preference in females matches this variation, with female hybrids preferring the intermediate songs (Hoy and Paul 1973; Hoy 1974; Hoy et al. 1977; Shaw 2000). This suggests that, like the motor mechanisms of song production, the mechanisms underlying temporal auditory filtering are under quantitative control and thus capable of being altered to produce shifts in sensitivity along a continuous amplitude modulation scale.

Although other neurons in crickets are more directly implicated in delivering song modulation information to the brain (Hennig 1988; Schildberger et al. 1989), there are several lines of evidence that support ON1's role in amplitude modulation coding as well. First, amplitude modulation tuning to calling song carrier frequencies matches that of phonotaxis behavior and is positively correlated to species-specific amplitude modulation song rates (Nabatiyan et al. 2003; Marsat and Pollack 2004; Tunstall and Pollack 2005). Second, sensitivity to amplitude modulation is carrier frequency dependent (Marsat and Pollack 2004; Sabourin et al. 2008), and there is little evidence for Ca^{2+} -induced ahp for ultrasound stimulation (Pollack 1988). Third, as an intrasegmental neuron, ON1 functions, in part, to increase binaural contrast through inhibition of the contralateral ON1 and ascending units (Selverston et al. 1985; Horseman and Huber 1994). But, the mechanisms of coding amplitude modulation appear to operate independently of the directional mechanisms (Marsat and Pollack 2005), meaning sensitivity to amplitude modulation is not an artifact or emergent property of binaural mechanisms. Fourth, during the Ca^{2+} -mediated selective attention (Sobel and Tank 1994), filtering must be timed correctly to prevent the filtering of subsequent pulses from the attended male (i.e., filter too slow) or coding song pulses from other males falling in the attended male's

interpulse interval (i.e., filter too fast). Fifth, specialized amplitude modulation coding in ON1 increases the binaural contrast in ascending neurons for behaviorally relevant sounds (Marsat and Pollack 2004). Thus, the temporal properties of ON1 appear important to processing song amplitude modulation and phonotaxis.

The model shows that Ca^{2+} -sensitive adaptation and depression in ON1 are sufficient to mediate these temporal properties and capable of shifting sensitivity to different song modulation rates. Thus, this model generates a hypothesis for the underlying mechanism for *interspecific* (and hybrid) differences in temporal sensitivity: differences in the size and kinetics of the biophysical mechanisms of Ca^{2+} -mediated adaptation and PSP depression (e.g., Hirschberg et al. 1999; Pedarzani et al. 2001; Pedarzani et al. 2005; Nolting et al. 2007). As with other systems, our results suggest that small changes would be required to shift sensitivity (Katz and Harris-Warrick 1999).

Temperature coupling between sender and receiver

Temporal sensitivity in orthopterans varies not only between species, hybrids and for different carrier frequencies, but also with the temperature (Pires and Hoy 1992a, b). There is a linear relationship between song modulation rate and temperature (Walker 1962), which female sensitivity tracks (Doherty 1985a). With respect to omega neurons, *in vivo* recordings by Janiszewski and Otto (1989) showed that temporal coding varied with thoracic temperature. Lower temperatures were correlated to: reduced instantaneous spike frequency; reduced coding of faster song modulation rates; and increased spike duration. The latter result suggests, at the very least, that the kinetics of spike production is under temperature control. Thus, in proposing and testing a mechanism for changing temporal sensitivity in these neurons, the model included temperature in calculating I_{Na} activation and inactivation, and I_{K} activation (note that although temperature is used in the Goldman–Hodgkin–Katz equation to calculate I_{Ca} , the change in temperature over the range tested did not affect the activation curve of I_{Ca} ; data not shown). As in those results *in vivo* (Janiszewski and Otto 1989), the model showed that temporal coding was positively correlated to temperature: higher temperatures increase the TMTF cutoff frequency. There are two possible mechanisms for this effect. First, reductions in spike duration at high temperatures (Fig. 10) increase the simulation's ability to respond to faster changes in stimulus envelope by increasing the spike rate. Second, the reduction in spike duration at high temperatures decreases the time spent depolarized, as shown by changes in the integral of the spike train as a function of temperature (Fig. 10). Because this integral (i.e., the depolarization time) determines the activation of the voltage-sensitive Ca^{2+} conductance, it also determines the extent of hyperpolarization or adaptation. Thus, reduced spike duration reduces Ca^{2+} feedback, which broadens the TMTF.

The changes in temperature did not produce the quantitative shift in TMTF like that found in singing males (Walker 1962; Fig. 10), which is predicted for the temperature coupled cricket system (Doherty 1985a). Thus, based on our model, temperature-sensitive mechanisms other than those voltage-sensitive mechanisms that generate spikes must also contribute to the temporal response of omega neurons. Hypothetically, one such mechanism could be to couple temperature to the rate of Ca^{2+} removal, in which buffering, sequestering or extrusion would slow down at lower temperatures (Kang et al. 2008), producing longer activation of Ca^{2+} -activated ahp currents, increasing PSP depression (Lee et al. 2005).

In conclusion, this study adds to the other predictive models for amplitude modulation filtering in insects (Benda et al. 2001; Hennig 2003; Bush and Schul 2006). It is distinguished, however, in that it offers computational evidence that particular biophysical mechanisms, the Ca^{2+} -mediated mechanisms of selective attention, are sufficient to explain the temporal sensitivity of omega neurons. Furthermore, the model shows that small changes (<1 SD) in the parameters

of these biophysical mechanisms can shift temporal sensitivity, an important character in the contexts of speciation and predator detection. From a neuroethological point of view, empirical validation of this prediction would require, at the very least, an extension of the comparative analysis by Marsat and Pollack (2004) to include measurements of ahp rates across taxa (or hybrids) with different calling song modulation structure.

Acknowledgments

Two anonymous reviewers as well as C. Canavier, S. Selvakumar, S. Achuthan and W. Gordon provided useful comments on the project and/or manuscript. N. Bazan generously provided equipment and software. No animals were used for this computational project. The study was funded in part by the NIH Grant # P20RR016816 to N. Bazan.

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Abbreviations

Ahp	After-hyperpolarizing
ERB	Equivalent rectangular bandwidth
I/O	Input/output
ON1	Omega neuron 1
PSP	Postsynaptic potential
SFP	Spike frequency peak
SNR	Signal-to-noise ratio
τ_{fast}	Fast adaptation time constant
TMTF	Temporal modulation transfer function
τ_{removal}	Time constant for Ca ²⁺ removal
τ_{slow}	Slow adaptation time constant

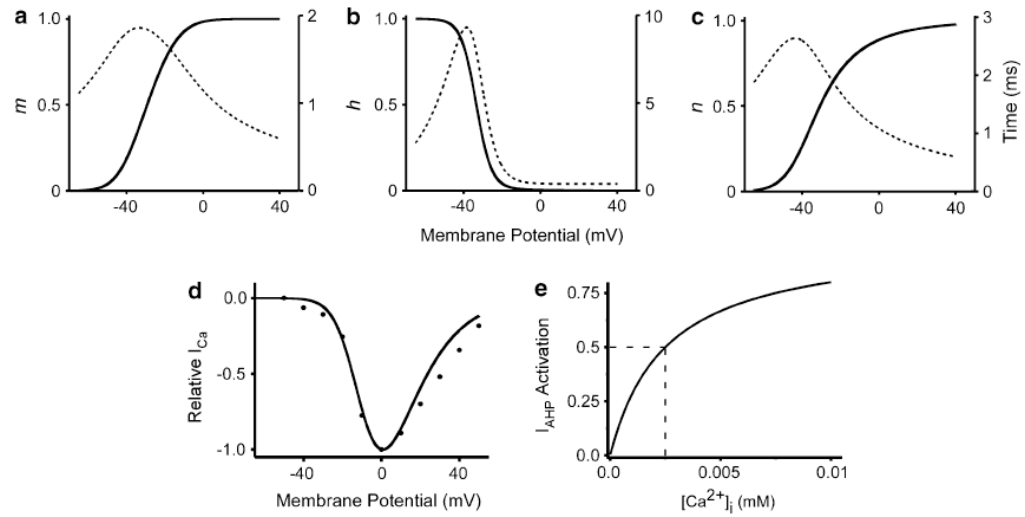


Fig. 1. Activation, inactivation (*solid lines*) and time constants (*dotted lines*) for model conductances. **a** I_{Na} activation. **b** I_{Na} inactivation. **c** I_K activation. *Solid lines* correspond to left axes, *dotted lines* correspond to right axes. **d** I/V relationship for I_{Ca} . Curve is the simulation. Points are data from cockroach (Wicher and Penzlin 1997) **e** Ca^{2+} sensitivity of I_{ahp} ; *dashes* note half activation point

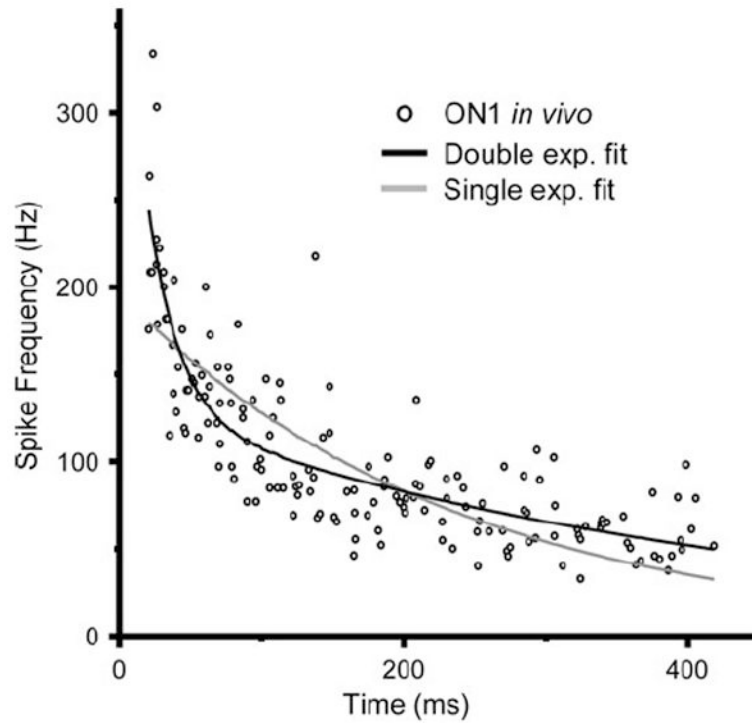


Fig. 2.

Points show instantaneous spike frequency during adaptation for a real ON1 in response to 400 ms duration pulse of 5 kHz (5 repetitions)(Farris et al. 2004). *Black* and *gray* curves are the double ($a \cdot e^{-t/\tau_{\text{fast}}} + b \cdot e^{-t/\tau_{\text{slow}}}$) and single ($d \cdot e^{-t/\tau}$) exponential fits to the data, respectively. Double exponential parameters ($n = 9$): $\tau_{\text{slow}} = 1054.3 \pm 470.2$ ms; $\tau_{\text{fast}} = 23.2 \pm 13.2$ ms; $a = 607 \pm 581$; $b = 188 \pm 44$; single exponential parameters: $\tau = 520 \pm 171$ ms; $d = 235 \pm 53$

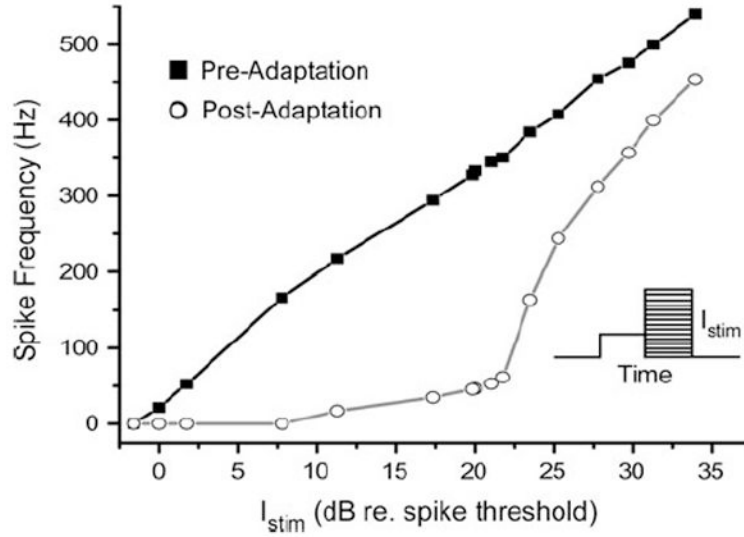


Fig. 3.

Adaptation of the model shifts the rate–intensity function to the right, but does not compress the curve by limiting maximum spike frequency. Squares are the instantaneous spike frequency (first two spikes) when stimulating at rest; circles are that following 1.5 s stimulation of $I_{stim} = 21$ dB (relative to spike threshold). The linear-like response is similar to that found in vivo (Baden and Hedwig 2007). *Inset* shows schematic of stimulus for adapted I/O curve. A 1.5 s pulse at mean I_{stim} is followed by a second 1.5 s pulse at variable amplitude

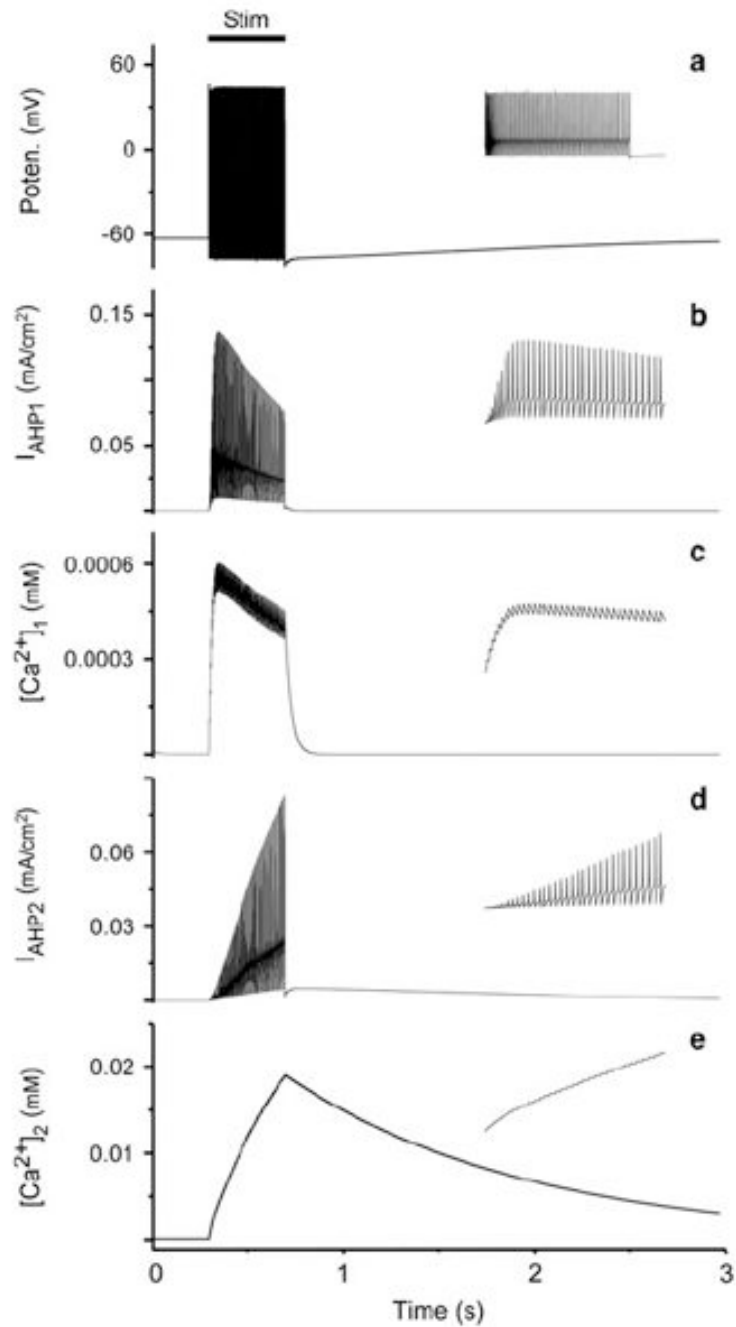


Fig. 4. **a** Voltage response using mean adaptation parameters generated from simulation of the nine cells. **b–e** I_{ahp} and corresponding $[Ca^{2+}]_i$ for the simulation in **a** (see y axis label for parameter). Insets show expansion of response over 500 ms in **a** and over 200 ms in **b–e**. Stimulus duration is 400 ms

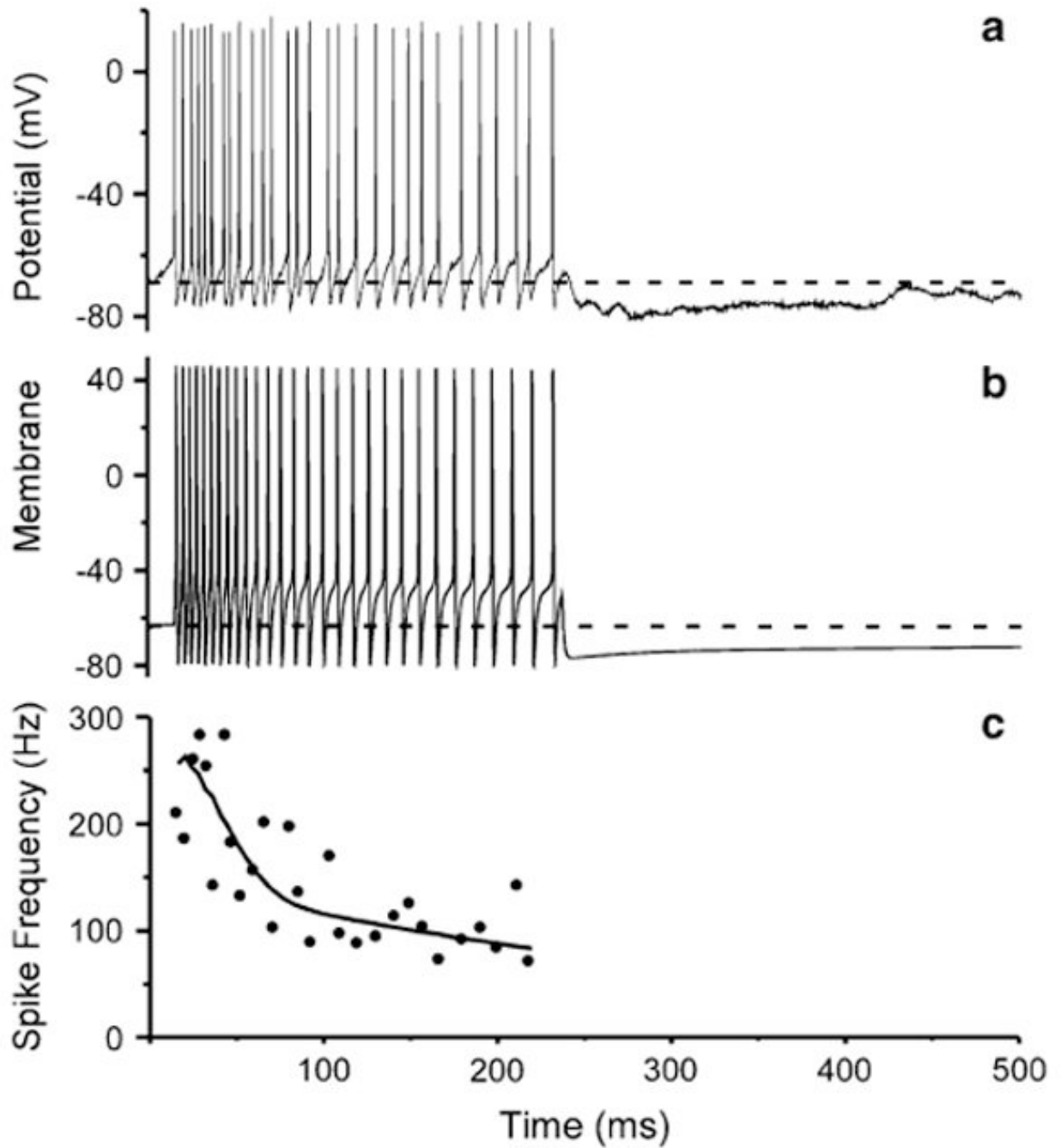


Fig. 5. Simulation of ON1 response. **a** Response of ON1 in *G. rubens* female to a 200 ms duration pulse of 5 kHz (1 ms ramps). **b** Simulation of response in **a** following adaptive least-squares fit of adaptation parameters. *Gray dashed line* is resting potential. **c** Comparison of instantaneous spike frequency in **a** (*filled circles*) and **b** (*black curve*)

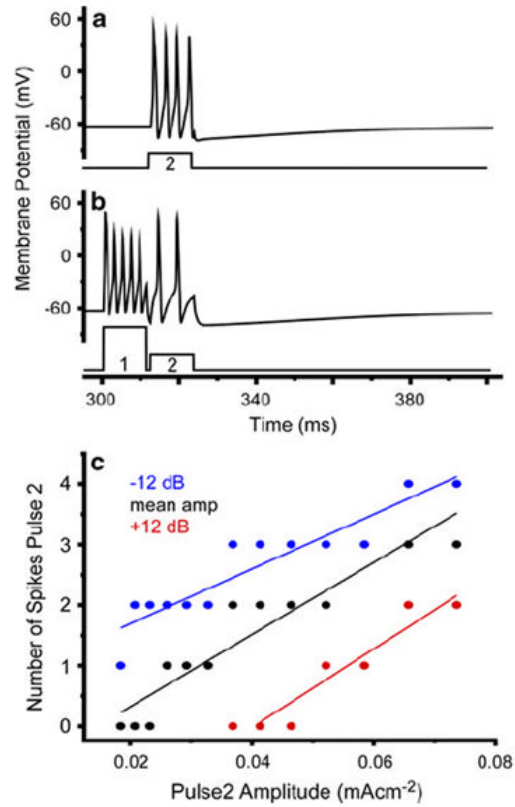


Fig. 6. Simulation exhibits selective attention in the form of greater sensitivity to the preceding pulse. **a** Voltage response to current pulse 2 alone ($I_{stim} = 0.0657 \text{ mA/cm}^2$). **b** Addition of current pulse 1 (0.185 mA/cm^2 ; 9 dB relative to pulse 2) reduces response to pulse 2. **c** Number of spikes (for pulse 2) as a function of pulse 2 amplitude for three different pulse 1 amplitudes (mA/cm^2): 0.0184 (*blue*); 0.074 (*black*); 0.293 (*red*). Plots are linear regression fits for the three pulse 1 amplitudes (*blue* slope = 45.3; intercept = 0.78; *black*: slope = 59.7, intercept = -0.88; *red*: slope = 64.7, intercept = -2.61). Increases in pulse 1 amplitude shifts the I/O response to the right

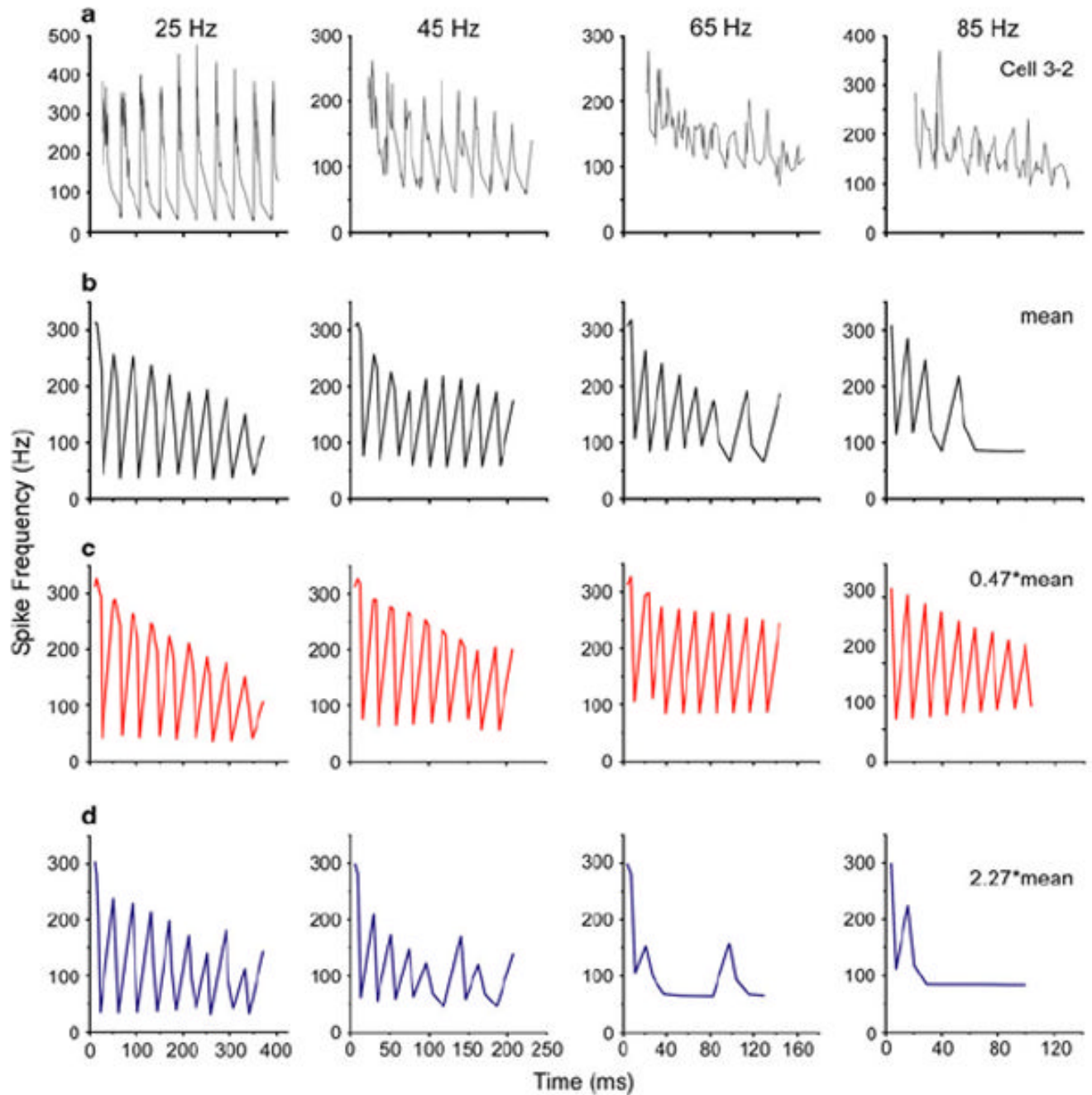


Fig. 7. Examples of spike frequency waveform in response to ten cycles of modulated stimuli. **a** Response of a real ON1 to an amplitude-modulated tone (5 kHz) at four different rates near that for the calling song. The parameters used to simulate this cell's response to a single acoustic pulse were near the mean. **b** Response of the model using the mean parameters. This response is altered when $\bar{g}_{ahp,1}$ is decreased (**c**) or increased (**d**). Modulation rates are noted above each column

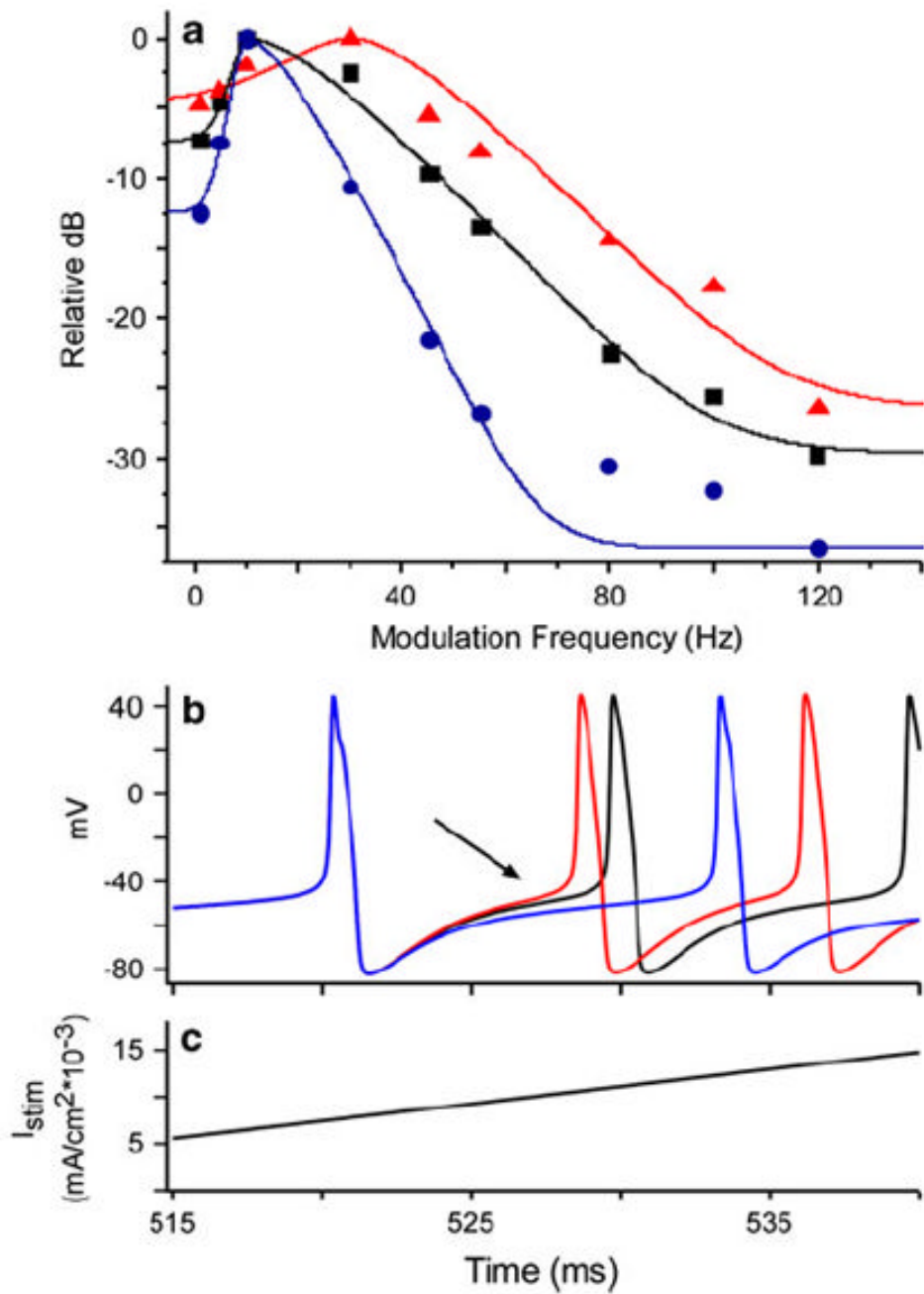


Fig. 8.
a Temporal modulation transfer functions (TMTF) for three values of $\bar{g}_{ahp,1}$. Each point is the relative power in the spike frequency waveform at the stimulus modulation frequency. Curves are the roex model fit for each TMTF. Whereas the black data are for the mean parameter value ($\bar{g}_{ahp,1} = 0.047 \text{ S/cm}^2$), blue are for an increase ($\bar{g}_{ahp,1} = 0.107 \text{ S/cm}^2$) and red a decrease ($\bar{g}_{ahp,1} = 0.022 \text{ S/cm}^2$). These changes had the opposite effect on TMTF bandwidth (cutoff frequency: red 46.5 Hz; black 26.1 Hz; blue 18.7 Hz). The value for the parameter decrease (red curve) was chosen to match the calcium signal observed in ON1 dendrites in response to

ultrasound relative to that for 5 kHz (47%; Baden and Hedwig 2007). **b** Membrane response to a ramped stimulus (**c**) is used to show difference in input resistance (Chen and Toney 2009). *Colors* represent the same $\bar{g}_{\text{ahp},1}$ values as in **a**. *Arrow* shows that after the depolarization of the first spike when I_{ahp} is activated, the same I_{stim} does not elicit the same depolarization. Higher values of $\bar{g}_{\text{ahp},1}$ require longer stimulation to elicit the next spike. Note that ramp stimulus starts at 500 ms

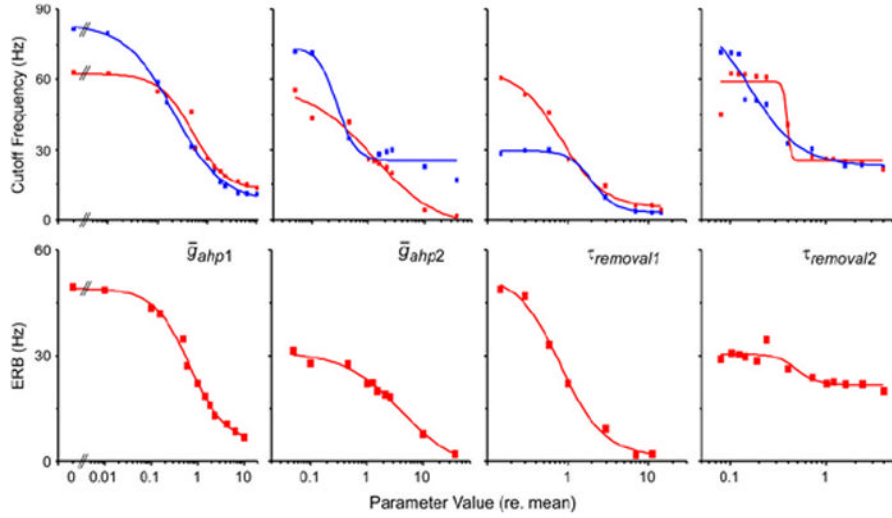


Fig. 9. Change in temporal modulation transfer function associated with variance in the four adaptation parameters (noted in each column). The *top row* is the low-pass cutoff frequency as a function of parameter value (relative to mean parameter value from the simulation of nine *G. rubens* ON1s using response to a single pulse). *Red symbols* are for spectral analysis of the spike frequency waveform. *Blue symbols* are from analysis using spike frequency peaks/stimulus pulse, in which peak discrimination threshold is 200 Hz. The *bottom row* is the equivalent rectangular bandwidth of the spectral filter versus parameter value. Steepness of the ERB curves demonstrates the extent to which the change in cutoff frequency in the top row is due to a change in filter bandwidth as opposed to shifting center frequency. Only one parameter is varied at a time, with other parameter values equaling the mean. All curves are single Boltzman fits to the data (Table 2)

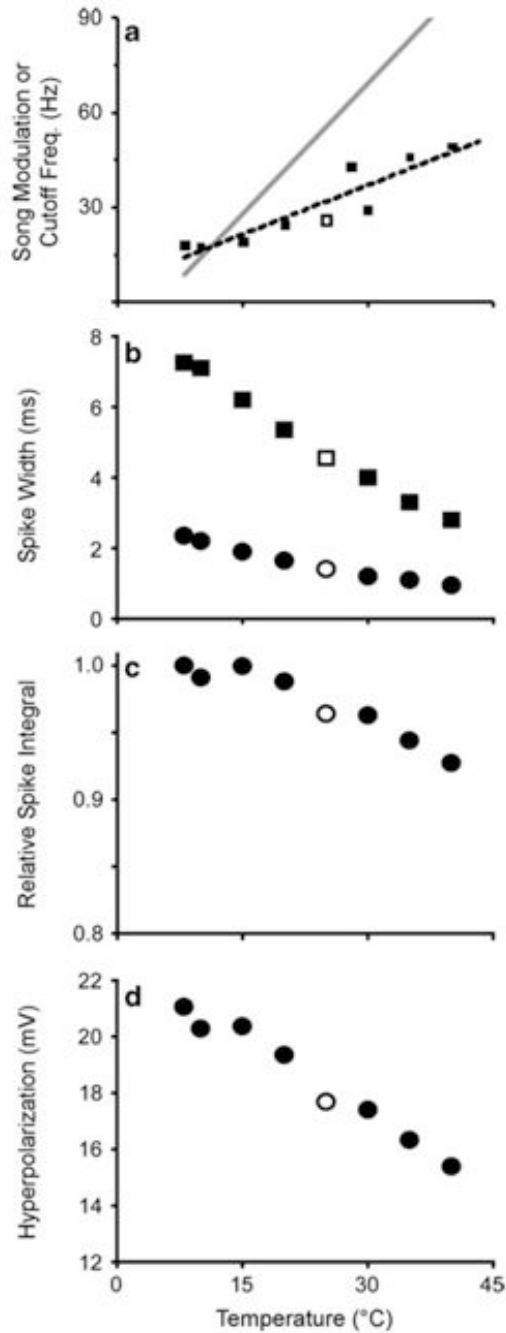


Fig. 10. Effect of temperature on spike generation and adaptation. **a** TMTF of amplitude modulation coding. *Dotted line* is linear fit to data (slope = 1.04; intercept = 5.9). This slope is shallow relative to the change in song modulation frequency with temperature exhibited by males (*solid gray line* slope = 2.756; intercept = -13.12; Walker 1962). **b** Speed of spike generation increases with temperature. *Plot* shows the spike width (duration) during depolarization above -55 mV. *Squares* and *circles* are the widths of the first spike and the first spike after $\tau_{\text{removal},1}$ (34.4 ms), respectively. **c** Reduced spike width caused by increasing temperature

reduces the integral of depolarization by ~ 10%. This reduction leads to reduced activation of I_{Ca} and I_{ahp} , reducing hyperpolarization as shown in **d**. *Open symbols* are simulations run at the standard 25°C

Table 1

Values for constants and initial values (at time = 0) for particular variables

Constants	Value
Sodium equilibrium potential (E_{Na})	55 mV
Potassium equilibrium potential (E_K)	-85 mV
Calcium equilibrium potential (E_{Ca})	100 mV
Faraday constant (F)	96480 C
Action potential threshold ($v_{threshold}$)	-55 mV
Ideal gas constant (R)	$8.314 \text{ V-C}(\text{°C})^{-1}$
Membrane capacitance (C)	$1 \mu\text{F cm}^{-2}$
Sodium conductance (\bar{g}_{Na})	0.11 S cm^{-2}
Potassium conductance (\bar{g}_K)	0.1 S cm^{-2}
Calcium permeability (\bar{P}_{Ca})	$0.0005 \text{ cm sec}^{-1}$
Sodium permeability (\bar{P}_{Na})	$0.00000207 \text{ cm sec}^{-1}$
Potassium permeability (\bar{P}_K)	$0.00000345 \text{ cm sec}^{-1}$
Variables	Initial value
Membrane potential (V)	-65 mV
Internal calcium concentration pool 1 ($[Ca^{2+}]_{i,1}$)	0.00001 mM
Internal calcium concentration pool 2 ($[Ca^{2+}]_{i,2}$)	0.00001 mM
External calcium concentration ($[Ca^{2+}]_o$)	2 mM
Sodium activating (m)	0
Sodium inactivating (h)	1
Potassium activating (n)	0

Table 2

Parameters for fitted curves shown in Fig. 9, which show the change in the cutoff frequency of the temporal modulation transfer functions (for two methods of analysis) and equivalent rectangular bandwidth (ERB; Hartmann 1998) as a function of biophysical parameter value

Spike train analysis method	Biophysical parameter	Inflection point (relative to mean value)	Function range (Hz)	Slope	R ² fit
Spike frequency peaks	$\bar{g}_{ahp,1}$	0.216	75.93	0.88	0.996
	$\bar{g}_{ahp,2}$	0.276	48.40	2.58	0.966
	$\tau_{removal,1}$	2.011	26.48	2.97	0.998
	$\tau_{removal,2}$	0.160	67.14	1.65	0.965
Spike frequency spectrum	$\bar{g}_{ahp,1}$	0.515	49.57	1.29	0.978
	$\bar{g}_{ahp,2}$	1.247	60.61	0.75	0.965
	$\tau_{removal,1}$	0.8177	58.99	1.62	0.989
	$\tau_{removal,2}$	0.392	33.96	22.47	0.931
ERB of spike frequency spectrum	$\bar{g}_{ahp,1}$	0.683	44.05	1.13	0.994
	$\bar{g}_{ahp,2}$	4.169	33.67	0.84	0.987
	$\tau_{removal,1}$	0.808	52.34	1.57	0.995
	$\tau_{removal,2}$	0.497	8.87	4.10	0.875

Columns are the inflection point of the Boltzman curve normalized to the mean biophysical parameter value; the minimum to maximum range of the function; the slope and the R² value of the curve fit. The biophysical parameters are the ahp conductances (\bar{g}_{ahp}) and the Ca²⁺ removal times ($\tau_{removal}$) for pools 1 and 2