Kinetics and connectivity properties of Parvalbumin- and Somatostatin-positive inhibition in layer 2/3 medial entorhinal cortex

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Parvalbumin (Pvalb⁺)- and somatostatin (Sst⁺)-positive cells are the two largest subgroups of inhibitory interneurons. Studies in visual cortex indicate that synaptic connections between Pvalb⁺ cells are common while connections between Sst⁺ interneurons have not been observed. The inhibitory connectivity and kinetics of these two interneuron subpopulations, however, have not been characterized in medial entorhinal cortex (mEC). Using fluorescence-guided paired recordings in mouse brain slices from interneurons and excitatory cells in layer 2/3 mEC, we found that, unlike neocortical measures, Sst⁺ cells inhibit each other, albeit with a lower probability than Pvalb⁺ cells (18% versus 36% for unidirectional connections). Gap junction connections were also more frequent between Pvalb⁺ cells than between Sst⁺ cells. Pvalb⁺ cells inhibited each other with larger conductances, smaller decay time constants and shorter delays. Similarly, synaptic connections between Pvalb⁺ and excitatory cells were more likely and expressed faster decay times and shorter delays than those between Sst⁺ and excitatory cells. Inhibitory cells exhibited smaller synaptic decay time constants between interneurons than on their excitatory targets. Inhibition between interneurons also depressed faster, and to a greater extent. Finally, inhibition onto layer 2 pyramidal and stellate cells originating from Pvalb⁺ interneurons were very similar, with no significant differences in connection likelihood, inhibitory amplitude, and decay time. A model of short-term depression fitted to the data indicates that recovery time constants for refilling the available pool are in the range of 50-150 ms and that the fraction of the available pool released on each spike is in the range 0.2-0.5.

Significance

Two large and distinct classes of interneurons in medial entorhinal cortex (mEC) include parvalbumin (Pvalb⁺)- and somatostatin (Sst⁺)-positive cells. Previous work has demonstrated unique functions with regards to spatial tuning and network oscillations for these two interneuron populations. Potential differences in kinetics of inhibition and likelihood of connection from these two interneuron groups, however, have not been quantified. Here, using fluorescence to guide intracellular recordings, we quantified the synaptic connections from both types of interneurons. We indicate that Sst⁺ and Pvalb⁺ express different synaptic kinetics that

are target-cell specific. In contrast to neocortical measures, we find substantial connections between Sst⁺ interneurons.

Introduction

Medial entorhinal cortex (mEC) plays a significant role in spatial navigation (Burgalossi and Brecht, 2014; Sasaki et al., 2015). In layer 2/3 mEC, the neurophysiological correlates of this role are partially supported by spatially tuned cells ('grid cells') that generate spikes at the vertices of a hexagonal grid formed during an animal's movement (Fyhn et al., 2004; Hafting et al., 2005). The region also generates theta-nested gamma frequency oscillations (Colgin et al., 2009) that synchronize grid cell spiking to specific phases of a network-wide theta oscillation (Hafting et al., 2008; Reifenshtein et al., 2012).

Both spatial tuning and oscillations in mEC are often accounted for using a canonical circuit composed of excitatory and inhibitory cells connected through recurrent excitation and negative feedback (Shipston-Sharman et al., 2016). In mEC, fast-firing interneurons participate in theta-nested gamma oscillations through a mechanism similar to pyramidal-interneuron gamma oscillations in cortex and hippocampus (Pastoll et al., 2013); by inhibiting stellate cells, these neurons set the phase of spiking and frequency of gamma oscillations during network activation. Fast-firing interneurons also provide the sole synaptic communication path between stellate cells, which lack recurrent excitatory connections (Couey et al., 2013; Pastoll et al., 2013) (but see (Fuchs et al., 2016)). In contrast, a subset of low threshold-spiking interneurons have been shown to suppress network oscillations in mEC (de Filippo et al., 2021). Further, in behaving animals, inhibition from layer 2/3 parvalbumin (Pvalb⁺)- and somatostatin (Sst⁺)-positive interneurons have separate roles in setting the spatial tuning and firing rates of layer 2 grid cells (Miao et al., 2017).

In cortex and hippocampus, Pvalb⁺ and Sst⁺ interneurons comprise two families of interneurons that, to a first approximation, correspond to fast-firing and low threshold-spiking interneurons, respectively (Rudy et al., 2011; Kepecs and Fishell, 2014; Tremblay et al., 2016; Pelkey et al., 2017). Across different regions, fast-acting, perisomatic-targeting inhibitory feedback typically originates from fast-firing, Pvalb⁺ cells (Kawaguchi and Kubota, 1997, 1998; Kubota et al., 2016). These cells are highly interconnected through synapses and gap junctions (Bartos et al., 2002; Hjorth et al., 2009). Conversely, Sst⁺ interneurons are more diverse electrophysiologically (Tremblay et al., 2016; Yavorska and Wehr, 2016), target dendrites (Kawaguchi and Kubota, 1998), can be electrical coupled (Amitai et al., 2002; Fanselow et al., 2008), and provide inhibition onto other, non-Sst⁺ neurons, with little evidence of synaptic connections between Sst⁺ cells (Pfeffer et al., 2013). Specifically, measures in visual cortex indicate a lack of synaptic connections between Sst⁺ interneurons (Pfeffer et al., 2013).

To date, the likelihood and synaptic kinetics of inhibition from Pvalb⁺ and Sst⁺ interneurons in mEC have not been measured. Although some properties are likely shared with other brain regions, significant differences have been observed between regions (Tremblay et al., 2016; Yavorska and Wehr, 2016). Measures in mEC, therefore, can guide specific mechanisms and models of mEC activity with regards to the role of inhibition in spatial tuning and network synchrony. Using Cre-based expression of the tdTomato fluorophore, we targeted Pvalb⁺ and Sst⁺ interneurons and used paired recordings from mouse slices to establish the properties of inhibition, both between interneurons, as well as onto excitatory cells, in layer 2/3 mEC.

Methods

Ethics statement

All experimental protocols were approved by the Boston University Institutional Animal Care and Use Committee.

Transgenic mice

To target Pvalb⁺ and Sst⁺ interneurons, we used mice expressing the red fluorescent protein tdTomato in either interneuron population. For Pvalb⁺ neurons, C57BL/6J background, Pvalb-Cre mice (Taniguchi et al., 2011) (Jackson Labs, stock # 017320) were crossed with the lox-stop-lox tdTomato reporter mice (Zariwala et al., 2011)(Jackson Labs, stock # 007914). The same reporter mouse was crossed to Sst-Cre mice (Taniguchi et al., 2011)(Jackson Labs, stock # 013044) to visualize Sst⁺ interneurons.

Slice preparation

Horizontal slices of entorhinal cortex and hippocampus were prepared from 2-8 month-old mice of either sex. After anesthetization with isoflurane and decapitation, brains were removed and immersed in 0 °C sucrose-substituted artificial cerebrospinal fluid (in mM): sucrose (185), KCl (2.5), NaH₂PO₄ (1.25), MgCl₂ (10), NaHCO₃ (25), Glucose (12.5), CaCl₂ (0.5). Slices were cut to a thickness of 400 μm (Leica VT 1200, Leica Microsystems). Slices were then incubated at 35°C for 20 minutes in artificial cerebrospinal fluid (ACSF) consisting of the following (in mM): NaCl (125), NaHCO₃ (25), D-glucose (25), KCl (2), CaCl₂ (2), NaH₂PO₄ (1.25) and MgCl₂ (1). Afterwards, slices were cooled to room temperature (20 °C). After the incubation period, slices were moved to the stage of a two-photon imaging system (Thorlabs) with a mode-locked Ti:Sapphire laser (Chameleon Ultra II; Coherent) set to wavelengths between 915 nm and 950 nm, which was used to excite both the Alexa Fluor 488 and tdTomato using a 20X, NA 1.0 (Olympus) objective lens. Laser scanning was performed using resonant scanners and fluorescence was detected using two photo-multiplier tubes (Hamamatsu) equipped with red and green filters to separate emission from Alexa Fluor 488 and tdTomato. The stage of the microscope contained recirculating ACSF, with all recordings conducted between 34 °C and 36 °C.

Electrophysiology

Electrodes were pulled using a horizontal puller (Sutter Instruments) using filament, thin-wall glass (Sutter Instruments). Intracellular pipette solution consisted of the following (in mM): K-gluconate (120), KCl (20), HEPES (10), diTrisPhCr (7), Na₂ATP (4), MgCl₂ (2), Tris-GTP (0.3), EGTA (0.2) and buffered to pH 7.3 with KOH. To visualize electrodes, the cyan-green fluorescent dye Alexa Fluor 488 hydrazide (Thermo Fisher Scientific) was added to the intracellular electrode solution (0.3 % weight/volume). To patch non-fluorescent, excitatory cells, we used a “shadow” patch technique (Kitamura et al., 2008) in which extracellular green fluorescence contrast with cells that do not fluoresce. Although this does not exclude the possibility of recording from interneurons, the values of spike half-width in probable excitatory cells indicated little overlap with those of Pvalb⁺ and Sst⁺ neurons.

Electrode resistances were between 4 and 7 MΩ, with access resistance values between 15 and 38 MΩ. Seal resistance values were always greater than 2 GΩ. Capacitance was fully

compensated in voltage clamp during the on-cell configuration prior to breaking into the cell. For current-clamp recordings, full bridge balance compensation was used. Series resistance compensation between 45-65 % was used during voltage clamp recordings. Voltage trace signals were amplified and low-pass filtered at 10-20 kHz before being digitized at 20-50 kHz. For current traces, signals were low pass filtered at 4 kHz. All electrophysiology was carried out using a Multiclamp 700B (Molecular Devices) and a Digidata 1550 (Molecular Devices). Liquid junction potentials were not corrected.

Recording protocols

A series of 1-second-long hyperpolarizing and depolarizing current pulses were used to generate spike frequency-current relationships. Spike half-width was taken from the first current pulse that generated spikes. Using the same data, the membrane decay time constant was acquired using an exponential fit to the voltage. To determine the presence of a synaptic connection, the post-synaptic cell was depolarized to -40 mV, while driving the pre-synaptic cell with a brief (2 ms), strong (>0.5 nA) pulse that drove a single spike. For measures of excitatory synaptic connections on interneurons, cells were voltage clamped at -70 mV. For frequency-dependent synaptic depression measures, pulses were delivered at 5, 10, 20, 50, 100, and 200 Hz. Synaptic current responses were averaged across 25-50 trials. For gap junction measures, a square pulse or spike was generated in the pre-synaptic cell and 25-50 trials averaged in the post-synaptic cell. A measured junction potential of ~11 mV was not subtracted from recordings. Recordings were taken from slices between 3.2 mm and 4.3 mm from the dorsal surface (bregma) of the brain.

Data Analyses

For current clamp analyses of spike shape, spike threshold was defined using the peak of the second derivative of the spike waveform. Spike half-width was taken as the width of the spike at voltages corresponding to the half amplitude (mid-point between spike peak amplitude and threshold). For time constant measures, a current pulse was used to depolarized cells to a value slightly below spike threshold. A single exponential decay function was used to fit the membrane voltage time course associated with hyperpolarization and the return to resting voltage resulting from the end of the current pulse.

Averaged individual post-synaptic current decay and rise time courses were fit with single exponential functions. Synaptic delay was measured as the time between spike threshold and the 10 % rise time of the averaged synaptic current response. All peak amplitudes were taken as the peak of the averaged synaptic current response. In a subset of recordings, where individual responses were large, we compared the averaged response to the distribution across trials. In this data set, mean responses sat near the center of the trial-to-trial distribution, suggesting that averaging did not impact our estimates of rise kinetics and delays.

The probability of chemical synaptic connectivity was calculated assuming that the probability of connections in each direction between a pair of neurons were equal and independent.

Statistical Analyses

All values are presented as the mean along with the standard deviation. Normality of data points was established using a Shapiro-Wilk and Lilliefors test. A positive result ($p < 0.05$) from either of these tests was used to determined normality and the use of non-parametric statistical tests noted in the Results section. For non-normally distributed data points, results are presented as the median along with the first (Q_1) and third (Q_3) quartile values.

Modeling

The model by Markram and Tsodyks (1996) predicts the amplitude of inhibitory currents from a differential equation for the available fraction of transmitter x that evolves according to:

$\frac{dx}{dt} = \frac{1-x}{\tau_r} - U_{SE}x\delta(t-t_s)$, where t_s is the spike time, τ_r is the recovery time to replenish the available pool of vesicles for release, U_{SE} is the fraction of available pool released by each spike, and the value of x prior to a spike is proportional to the peak current value of the IPSC. The

solution to the differential equation between presynaptic spikes is $x(t) = 1 - (1 - x_i)e^{-\frac{(t-t_s)}{\tau_r}}$. However, there is a discontinuous decrement, x , by an amount $U_{SE}x$ after each spike so a

discrete map from spike to spike is required: $x_i = 1 - (1 - x_{i-1}(1 - U_{SE}))e^{-\frac{(t-t_s)}{\tau_r}}$ where i denotes presynaptic spike number. Normalizing the spike amplitude so that the initial value of x is 1 and substituting one over the frequency of the presynaptic spike train for $t-t_s$, we can write this recursively:

$$x_k = \left(1 - e^{-\frac{1}{f\tau_r}}\right) \frac{1 - \left[(1 - U_{SE})e^{-\frac{1}{f\tau_r}}\right]^k}{1 - (1 - U_{SE})e^{-\frac{1}{f\tau_r}}} + [(1 - U_{SE})e^{-\frac{1}{f\tau_r}}]^k$$

We fit each experimental train of the normalized inhibitory peak amplitudes to the expression above by adjusting the two parameters U_{SE} and τ_r to minimize the least squared error using *curve_fit* from *scipy.optimize*. If the R^2 for the linear regression between each train and the predicted values from the best fit was greater than 0.8, the trace was used in the calculation of the mean of the parameter values. Smaller values indicated a noisy trace, likely from a small synapse. Maximum likelihood estimation of the parameters yielded similar values and the data sets passed a Shapiro-Wilcox test for normality. The Akaike Information Criterion and RMS yielded similar results to R^2 but we chose to use R^2 to accept or reject data sets because the values are constrained (between 0 and 1) whereas the other metrics are not. The expression above sufficed for the connections between interneurons (Sst-Sst and Pvalb-Pvalb). The fit to the data from synapses onto the excitatory cells were improved by adding a correction for

temporal summation from up to three previous spikes $x_{k-j}e^{-\frac{j}{f\tau_r}}$ where j is the index of whether the contribution is from the immediately preceding spike ($j=1$) or farther back in the spike train. The previously measured time constant for synaptic decay at each synapse was used rather than fitting an additional unknown parameter.

Results

High degree of connectivity between mEC Pvalb⁺ interneurons

To address the likelihood of connections and synaptic kinetics of inhibition from layer 2/3 parvalbumin (Pvalb⁺)- and somatostatin (Sst⁺)-positive interneurons, we performed dual intracellular patch recordings in mice expressing the tdTomato fluorophore in Pvalb⁺ or Sst⁺ cells. We targeted both fluorescent and non-fluorescent cells and used intracellular electrophysiological measures to distinguish between subtypes of non-fluorescent cells (pyramidal and stellate cells) in layer 2.

Aside from the fluorescent marker, Pvalb⁺ cells could be differentiated from excitatory cells due a much smaller membrane time constant (6.8 ms [*Q1* 3.72 *Q3* 8.0]; $p < 0.001$, Kruskal-Wallis ANOVA; Fig. 1Aii) and spike half-width (0.29 ms [*Q1* 0.27 *Q3* 0.32], $p < 0.001$; Kruskal-Wallis ANOVA; Fig. 1Aiii), which is consistent with the fast-firing phenotype. In addition, we distinguished between layer 2 pyramidal and stellate cells using the membrane overshoot in response to a negative current pulse associated with the expression of a hyperpolarizing-activated cation current (I_H) and the membrane decay time constant; stellate cells expressed significantly larger membrane overshoots in response to a hyperpolarizing current (3.43 ± 2.13 mV vs. 0.45 ± 0.44 mV, $p < 0.001$, two-way student t-test; Fig. 1Aiv). Combined with a smaller membrane time constant, the two factors could be used to differentiate most stellate cells from pyramidal cells (Fig. 1Aiv). In cases where neither of these two factors was applicable, we used the presence of peri-threshold oscillations to distinguish cells (only used in 5 cells). To detect synapses and gap junctions, we averaged between 25 and 50 sweeps in voltage clamp (held at -40 mV) while driving a spike using a brief current pulse (2 ms) in the other neuron.

At pair distances between 0 and 150 μm , we observed a high probability of connection between mEC Pvalb⁺ interneurons. Over all distances tested, we found that 76/122 pairs expressed a connection either in the form of a chemical synapse (43/122), a gap junction (6/122) or both (27/122) (Fig. 1Bi); therefore, the probability of chemical synaptic connectivity was 35.7% (87/244) in each direction and the probability of bidirectional gap junctional connectivity was 27% (33/122). At distances greater than 150 μm , connection probability (of either type) fell sharply (Fig 1Biii), which is consistent with the extent of axonal arborization of Pvalb⁺ interneurons within layer 2/3 (Martínez et al., 2017; Grosser et al., 2021). We also found that the synaptic amplitude for uni-directionally connected pairs (21.4 pA [*Q1* 12.9 *Q3* 41.0]) did not differ significantly from the size of synapses present in bi-directionally connected pairs (30.9 pA [*Q1* 17.8 *Q3* 60.7], $p = 0.12$; Mann-Whitney test; Fig.1Biv). Using a step depolarization between 10 and 20 mV in one cell, we measured the strength of gap junction connections between Pvalb⁺ cells as the ratio of the two voltage responses. This value was 0.03 [*Q1* 0.02 *Q3* 0.06] across 17 measured pairs.

Next, the prevalence of gap junctions was measured between synaptically connected and unconnected pairs. In our data set, synaptically unconnected pairs occurred at longer pair distances than those of unidirectional or bidirectional connections (Fig. 1Bii). To eliminate this feature from artificially lowering the prevalence of gap junctions in synaptically unconnected pairs, we limited our analysis to pairs at distances between 20 and 125 μm . Under this constraint, the pair distances for all 3 categories of synaptic connections were statistically similar ($p = 0.57$; one-way ANOVA; no connection: 76.4 ± 23.9 μm ; unidirectional: 75.4 ± 26.8 μm ; bidirectional: 68.9 ± 27.3 μm ; Fig. 1Bv, inset). Using this data set, we found that the probability of gap junctions was significantly greater in bi-directionally (0.53 vs. 0.17, $p = 0.01$; two-sided Fisher test), but not uni-directionally, connected pairs when compared to unconnected pairs (Fig. 1Bv).

Connectivity and kinetics between mEC Pvalb⁺ interneurons and excitatory neurons

We carried out a similar set of analyses for connections between Pvalb⁺ and excitatory cells ($I_{\text{Pvalb}} - E$; Fig. 2A). Unlike interneuron pairs, we never observed gap junctions in $I_{\text{Pvalb}} - E$ cell pairs. Connection likelihood was also generally lower for $I_{\text{Pvalb}} - E$ pairs compared to $I_{\text{Pvalb}} - I_{\text{Pvalb}}$ pairs. Including both excitatory cell types, 38 of 85 $I_{\text{Pvalb}} - E$ pairs were connected in comparison to 76 of 122 $I_{\text{Pvalb}} - I_{\text{Pvalb}}$ pairs ($p = 0.02$; two-sided Fisher test). There were 26 unidirectional connections and 12 bidirectional connections; therefore, the probability of chemical synaptic connectivity was 29.4% (50/170) in each direction. Both pyramidal and stellate cells expressed statistically similar ratios of unconnected and connected pairs (33/26 vs. 14/12, $= 0.99$; two-sided Fisher test; Fig. 2 Bi, Ci) For both cell types, connection probability also dropped off at

distances greater than 100 μm (Fig. 2Biii, Ciii). Like $I_{\text{Pvalb}} - I_{\text{Pvalb}}$ pairs, we found no significant difference in the size of inhibitory synapses between uni-directionally and bi-directionally connected pairs of Pvalb⁺ and pyramidal ($I_{\text{Pvalb}} - E_{\text{Pyr}}$) cells (31.1 pA [$Q1$ 18.4 $Q3$ 100.2] vs. 14.2 pA [$Q1$ 7.6 $Q3$ 94.4], $p = 0.23$; Mann-Whitney test; Fig. 2Biv); we did not have enough bi-directionally connected Pvalb⁺-stellate ($I_{\text{Pvalb}} - E_{\text{Stel}}$) cell pairs to test potential differences in this measure (Fig. 2Civ).

The kinetics of inhibition indicated clear differences among the 3 cell types. We focused on amplitude, decay time, delay and rise time as these properties are crucial in many models of inhibitory activity with regards to synchrony and oscillations (Bartos et al., 2007; Economo and White, 2012; Keeley et al., 2017). The largest difference amongst these parameters was the inhibitory decay time constant (single exponential fit; Kruskal-Wallis ANOVA, $p < 0.001$; Fig. 3Aii). Synaptic inhibition decayed much faster in Pvalb⁺ cells than in either pyramidal or stellate cells (2.0 ms [$Q1$ 1.7 $Q3$ 2.6] vs. 5.5 ms [$Q1$ 4.9 $Q3$ 6.6] and 12.1 ms [$Q1$ 6.0 $Q3$ 14.5], $p < 0.001$; Dunn's test; Fig. 3Aii). Although generally larger, the difference in decay time constants between stellate and pyramidal cells did not reach the significance threshold ($p = 0.75$; Dunn's test, Fig. 3Aii).

We also found differences in the rise time constant between $I_{\text{Pvalb}} - I_{\text{Pvalb}}$ and stellate cell pairs ($p = 0.002$; Kruskal-Wallis ANOVA), with stellate cells showing larger rise time constants (0.34 ms [$Q1$ 0.28 $Q3$ 0.44] vs. 0.58 ms [$Q1$ 0.40 $Q3$ 0.81], $p = 0.002$; Fig. 3Aiii). In contrast, neither delay time ($p = 0.12$; one-way ANOVA) nor peak amplitude ($p = 0.06$; Kruskal-Wallis ANOVA) differed amongst the 3 pair categories (Fig. 3Ai, iv). Consistent with previous measures of excitatory (Song et al., 2005) and inhibitory (Bartos et al., 2001) synapses in other regions, the amplitude of inhibitory synapses expressed strong, non-Gaussian distributions (Shapiro-Wilk and Lilliefors test, $p < 0.001$). Distributions of $I_{\text{Pvalb}} - I_{\text{Pvalb}}$ and $I_{\text{Pvalb}} - E$ cell pair synaptic amplitudes fit an exponential decay and were dominated by small synapses accompanied by fewer, larger synapses (Fig. 3Bi, ii).

Connectivity and kinetics between mEC Sst⁺ interneurons

Next, we focused on inhibition from Sst⁺ interneurons. As with Pvalb⁺ neurons, Sst⁺ neurons could be differentiated from pyramidal cells through their membrane decay time constant and spike half-width (Fig. 4A). Although not as small as Pvalb⁺ neurons, both membrane time constant (13.1 ms [$Q1$ 7.9 $Q3$ 25.3] vs 39.2 ms [$Q1$ 29.0 $Q3$ 51.0]) and spike half-width (0.57 ms [$Q1$ 0.41 $Q3$ 0.68] vs 1.0 ms [$Q1$ 0.98 $Q3$ 1.2]) were significantly smaller than those of pyramidal cells ($p < 0.001$; Dunn's test, Fig. 4Aii).

Over all distances tested, we found that 16/59 Sst⁺ pairs expressed a connection either in the form of a chemical synapse (14/59) or a gap junction (2/59) (Fig. 4Bi). Compared with $I_{\text{Pvalb}} - I_{\text{Pvalb}}$ pairs, synaptic and gap connections between $I_{\text{Sst}} - I_{\text{Sst}}$ neurons were far less likely ($p < 0.001$, Fisher test). Crucially, due to the lack of connections at distances greater than 100 μm (Fig. 4iv), our average $I_{\text{Sst}} - I_{\text{Sst}}$ pair distance was shorter than $I_{\text{Pvalb}} - I_{\text{Pvalb}}$ pairs (46.7 μm [$Q1$ 28.9 $Q3$ 65.0] vs 81.3 μm [$Q1$ 61.9 $Q3$ 116]). Thus, despite shorter pair distances, connections probabilities were still far lower than those of $I_{\text{Pvalb}} - I_{\text{Pvalb}}$ pairs. We also found that the amplitude of synapses between $I_{\text{Sst}} - I_{\text{Sst}}$ pairs was significantly different depending on whether cells were unidirectionally or bidirectionally connected (6.0 pA [$Q1$ 5.2 $Q3$ 8.1] vs. 15.1 pA [$Q1$ 8.6 $Q3$ 26.4], $p = 0.02$; Mann-Whitney test; Fig. 4Biii). Unlike $I_{\text{Pvalb}} - I_{\text{Pvalb}}$ pairs, gap junctions in $I_{\text{Sst}} - I_{\text{Sst}}$ pairs were observed in only two pairs, both of which also had bidirectional synaptic connections.

Connectivity and kinetics between mEC Sst⁺ interneurons and pyramidal cells

For $I_{\text{Sst}} - E$ pairs, we limited our measures to pyramidal cells as we only observed 3 pairs containing stellate cells, with only 1 of these pairs being connected. This is likely due to the

deeper location of Sst⁺ neurons within layer 2/3 (Tremblay et al., 2016), as well as lower overall density across cortical layers (Amitai et al., 2002). This is also consistent with optogenetic experiments indicating greater connectivity between Sst⁺ and pyramidal cells (Kecskés et al., 2020). In contrast to I_{Sst}-I_{Sst} pairs, connection probabilities between I_{Sst}-E_{Py} were not significantly lower than those measured in I_{Pvalb}-E pairs (12/41 vs. 38/85, $p = 0.12$; two-sided Fisher test). If, however, we limited our connection measures in I_{Pvalb}-E pairs to a range similar to those of I_{Sst}-E_{Py} pairs (20 - 125 μ m), connection probability was indeed lower (12/41 vs. 37/73, $p = 0.03$; two-sided Fisher test). The amplitude of synapses between I_{Sst}-E_{Py} pairs also seemed to depend on whether cells were unidirectionally or bidirectionally connected. The difference, however, was at the margin of statistical significance (10.7 pA [$Q1$ 3.7 $Q3$ 20.9] vs. 39.4 pA [$Q1$ 23.5 $Q3$ 81.0], $p = 0.05$; Mann-Whitney test; Fig. 4Ciii).

As with Pvalb⁺ cells, inhibition from Sst⁺ cells onto pyramidal cells was significantly slower to decay than that between Sst⁺ cells (2.8 ms [$Q1$ 2.2 $Q3$ 5.2] vs 10.3 ms [$Q1$ 6.9 $Q3$ 14.9], $p < 0.001$; Mann-Whitney test; Fig. 5Aii). We also noted significantly longer delays in pyramidal cells (1.1 ± 0.14 ms vs. 1.4 ± 0.08 , $p = 0.04$; Student t-test). Both peak amplitude and rise time constants, however, were not significantly different ($p = 0.18$ and $p = 0.89$; Mann-Whitney test; Fig. 5Ai, Aiii). Again, distributions of peak inhibitory amplitude in both Sst⁺ and pyramidal cells expressed non-Gaussian distributions ($p < 0.001$; Shapiro-Wilk and Lilliefors test) and could be fit with exponential functions (Fig. 5Bi, Bii).

Pvalb⁺ and Sst⁺ interneuron-based inhibition express different decay and delay times

Comparison of inhibition in I_{Sst}-I_{Sst} and I_{Pvalb}-I_{Pvalb} pairs also indicated numerous differences. Inhibitory peak amplitude in I_{Sst}-I_{Sst} pairs was significantly smaller ($p = 0.002$; Mann-Whitney test), with larger decay time constants ($p < 0.001$; Mann-Whitney test) and longer delays ($p = 0.003$; Mann-Whitney test) than those measured in I_{Pvalb}-I_{Pvalb} pairs (Fig. 6Ai-iv). Similarly, we also noted differences between I_{Sst}-E_{Py} and I_{Pvalb}-E_{Py} pairs. These included larger decay time constants ($p = 0.005$; Mann-Whitney test; Fig. 6Bii) and delay times ($p = 0.002$; Mann-Whitney test; Fig. 6Biii) in I_{Sst}-E_{Py} pairs.

I-I synapses express greater synaptic depression than I-E synapses

Using 200 ms-long pulse trains between 5 – 200 Hz, we proceeded to measure the degree of synaptic depression at different stimulus frequencies. In both I_{Pvalb}-I_{Pvalb} and I_{Pvalb}-E synapses, depression was often observed at frequencies as low as 5 – 10 Hz (Fig 7A-C). In all 3 cell types, synaptic depression increased as a function of stimulus frequency (Fig. 7Di). To compare differences in the depression, we measured the depression decay time constant and steady-state value at 50 Hz, a frequency in which depression was observed reliably in all 3 cells and which reached a steady-state value within our stimulus time frame. As shown, the depression decay time constant and the steady-state value indicated faster and greater depression in I_{Pvalb}-I_{Pvalb} synapses than in either type of I_{Pvalb}-E connection ($n = 61, 20, 12$, $p < 0.001$, one-way ANOVA; Fig. 7Dii, iii), with I_{Pvalb}-I_{Pvalb} pairs expressing a time constant of 20.6 ± 7.2 ms (versus 34.9 ± 11.7 ms and 40.5 ± 13.6 ms, $p = 0.004$; Bonferroni test; Fig. 7Dii) and a steady-state value at 50 Hz of 0.44 ± 0.11 (vs. 0.55 ± 0.11 and 0.68 ± 0.11 , $p = 0.01$ and $p < 0.001$; Bonferroni test; Fig. 7Diii).

Using an identical stimulus range, we also measured the depression at I_{Sst}-I_{Sst} and I_{Sst}-E_{Py} synapses. Again, synaptic depression was observed at frequencies as low as 5 – 10 Hz. Like Pvalb⁺ cells, I_{Sst}-I_{Sst} synapses depressed more quickly and to a greater extent than I_{Sst}-E_{Py} synapses (Fig. 8A-C). The depression decay time constant in response to 50 Hz stimulus trains for I_{Sst}-I_{Sst} pairs was 20.1 ms [$Q1$ 15.7 $Q3$ 26.9], which was significantly smaller than the 51.9 ms [$Q1$ 38.8 $Q3$ 76.9] measured in I_{Sst}-E_{Py} synapses ($n = 20, 13$, $p = 0.002$; Mann-Whitney test; Fig.

8Ci). Steady-state levels at 50 Hz also differed, with $I_{Sst-I_{Sst}}$ pairs depressing to 0.44 [$Q1$ 0.42 $Q3$ 0.62] compared with 0.65 [$Q1$ 0.55 $Q3$ 0.74] in $I_{Sst-E_{Pyr}}$ ($p = 0.007$; Mann-Whitney test; Fig. 8Ciii). Finally, there was no difference in the synaptic depression time constants ($p = 0.55$; Mann-Whitney test) and steady-state levels ($p = 0.55$; Mann-Whitney test) of synaptic depression in response to 50 Hz between $I_{Pvalb-I_{Pvalb}}$ and $I_{Sst-I_{Sst}}$ pairs.

Modeling synaptic kinetics and short depression

To examine whether our results could be framed using previous mechanism of synaptic depression, we used a model of short-term depression developed by Markram and Tsodyks (1996) to fit our measures of post-synaptic current decay and depression (Markram and Tsodyks, 1996; Stimberg et al., 2019). In Figure 9, we show the best fitting models of synaptic depression as described in the Methods section. Because the synaptic decay time constants were much smaller for synapses between interneurons (Fig. 9A and D; $I_{Pvalb-I_{Pvalb}}$ and $I_{Sst-I_{Sst}}$, respectively) we were able to ignore temporal summation between successive post synaptic currents and still obtain a good fit. In contrast, synapses onto excitatory cells were generally slow enough that including their previously measured synaptic decay constants in the model as described in the Methods greatly improved the fit of the model (Fig. 9B, C and E for $I_{Pvalb-E_{Pyr}}$, $I_{Sst-E_{Pyr}}$ and $I_{Pvalb-E_{Stel}}$, respectively). This is consistent with the observation that inhibitory currents decay more slowly on excitatory compared to inhibitory neurons (Ma et al., 2012). The summary panel in Figure 9F shows that the time constants for recovery from depression ranges from about 50 to 150 ms, with no systematic differences in their values, as confirmed by the statistics in Table 1 ($p = 0.89$; one-way ANOVA). Most values of U_{SE} fall in the range 0.2 to 0.5, with some lower values (corresponding to less depression) associated with synapses onto pyramidal neurons (green and yellow symbols). Values, however, were not significantly different ($p > 0.05$; Dunn's test).

Discussion

In summary, we find that inhibition in $I_{Pvalb-I_{Pvalb}}$ and $I_{Pvalb-E}$ pairs in layer 2/3 mEC are stronger, faster, and more interconnected, both in term of synapses and gap junctions, than those measured in $I_{Sst-I_{Sst}}$ and $I_{Sst-E_{Pyr}}$ pairs. Further, for both $Pvalb^+$ and Sst^+ cells, inhibitory synaptic currents between interneuron pairs were faster to decay than in the excitatory cells; the inhibitory current decay time constant and delay time, however, were smaller and shorter, respectively, in $Pvalb^+$ cells. Although we noted some minor differences, $Pvalb^+$ inhibition onto pyramidal and stellate cells was generally similar. In contrast to previous measures in visual cortex (Pfeffer et al., 2013), we found significant evidence for inhibitory connections between Sst^+ interneurons. Finally, inhibition between interneurons expressed faster and greater synaptic depression than those onto excitatory cells.

Comparison with past measures of inhibitory synaptic currents

Like measures in neocortex (Beierlein et al., 2003) and hippocampus (Bartos et al., 2002), $I_{Pvalb-I_{Pvalb}}$ pairs expressed fast decaying synapses that were strongly depressing. These cell pairs also contained a high prevalence of gap junction connections, albeit with a lower likelihood than those measured in other neocortical regions (Gibson et al., 1999; Hjorth et al., 2009). Similarly, we also noted gap junction connections in $I_{Sst-I_{Sst}}$ pairs, but these were far less frequent than in $I_{Pvalb-I_{Pvalb}}$ pairs. This contrasts with measures in neocortex in which $I_{Sst-I_{Sst}}$ pairs are highly interconnected through gap junctions ($> 65\%$ of pairs)(Amitai et al., 2002; Fanselow et al., 2008). Also, in contrast to neocortex (Pfeffer et al., 2013), we measured substantial synaptic connections between Sst^+ neurons. These stark difference might arise, in part, from the large

diversity within the Sst⁺ population, which likely includes at least 4 different subtypes of interneurons across different cortical regions and layers (Urban-Ciecko and Barth, 2016; Yavorska and Wehr, 2016). Our analysis of spike half-width shows that SST⁺ cells share a similar range of variance as those in measured Pvalb⁺ cells. This suggests a narrowly defined population in layer 2/3 of the mEC.

Similar to CA1 (Bartos et al., 2002), we found that the inhibition decay time constants from Pvalb⁺ is target cell specific, with smaller decay time constants in Pvalb⁺ cells; this was also true for Sst⁺ interneurons. A key factor in determining the kinetics of GABA_A receptor-mediated inhibition is the subunit composition of the receptor (Fritschy and Brünig, 2003; Barberis et al., 2007). Faster decay times are associated with the presence of the alpha-1 and alpha-2 subunits (Barberis et al., 2007). Immunohistochemical work in layer 2 mEC indicates a greater prevalence of the alpha-1 subunit in Pvalb⁺ cells than in either reelin (stellate)- or calbindin (pyramidal)-positive excitatory cells (Berggaard et al., 2018). As a result, the faster decay time of inhibition can be explained by different subunit compositions that support a faster inactivation of the receptor in Pvalb⁺ cells.

Interneuron role in synchrony and gamma oscillations

Fast-firing interneurons are generally assumed to underlie fast negative feedback that is critical to balancing network excitation and inhibition (Wehr and Zador, 2003; Okun and Lampl, 2008; Atallah and Scanziani, 2009; Isaacson and Scanziani, 2011). In addition, for cells in close proximity (< 100 μm), nearly half of fast-firing interneurons are connected through gap junctions (Deans et al., 2001). These features support network synchrony, as well as the generation of fast gamma frequency oscillations (Beierlein et al., 2000; Deans et al., 2001; Bartos et al., 2002; Mann and Paulsen, 2007; Atallah and Scanziani, 2009; Pastoll et al., 2013). Consistent with this function, we observed that Pvalb⁺ cells inhibition onto pyramidal cells was faster to decay and arrived with a smaller delay time than Sst⁺ cells. Pvalb⁺ cells also expressed a higher prevalence of gap junction connections, and a much higher degree of interconnectivity between Pvalb⁺ neurons.

In models, the unique kinetics and connectivity properties of fast-firing interneurons can also generate gamma oscillations using solely an inhibitory network (Bartos et al., 2002, 2007; Tiesinga and Sejnowski, 2009; Tikidji-Hamburyan et al., 2015; Keeley et al., 2017; Tikidji-Hamburyan and Canavier, 2020), with experimental support for this form oscillatory activity noted in the hippocampus and entorhinal cortex (Butler et al., 2016, 2018). In contrast, neocortical Sst⁺ interneurons in vivo do not correlate with oscillatory network activity (Kwan and Dan, 2012). Although differences in inputs are likely involved (Pfeffer et al., 2013; Yavorska and Wehr, 2016), the lower synaptic and gap junction connectivity that we observed may also contribute to lower synchrony levels.

Like Pvalb⁺ cells, inhibition in Sst⁺ cells was faster to decay than that in excitatory neurons. The faster decay times of inhibition between interneurons, therefore, may serve roles independent of network synchrony and oscillations. For example, a longer inhibitory decay time onto excitatory targets may arise as a compensation mechanism for the larger membrane time constants of these cells relative to both Pvalb⁺ and Sst⁺ interneurons. The average firing rates of excitatory cells in layer 2 mEC are also typically lower than those in inhibitory cells (Frank et al., 2001), such that a proportional impact on spike rate requires longer lasting inhibition. As a result, differences in the inhibition kinetics may serve to compensate for differences in the neurophysiology of the post-synaptic target.

Kinetics of inhibition in pyramidal and stellate cells

Inhibition originating from Pvalb⁺ cell was generally very similar in pyramidal and stellate cells. The only small and significant difference was the greater degree of steady-state depression in pyramidal cells when synapses were driven at 50 Hz. This would appear to be consistent with the central role of Pvalb⁺ interneurons in shaping spatial selectivity of grid cells (Miao et al., 2017), which include both pyramidal and stellate cells (Domnisoru et al., 2013; Tang et al., 2014). Our results are also consistent with a recent comparison of Pvalb⁺ inhibition onto stellate and pyramidal cells along the dorsal-ventral axis (Grosser et al., 2021). The authors reported no difference in the amplitude, paired-pulse ratio, and connectivity likelihood between inhibition in stellate and pyramidal cells (Grosser et al., 2021). Nevertheless, past work has indicated stronger labeling for the slower α -3 GABA_A receptor subunit in stellate cells (Berggaard et al., 2018). Although we noted a tendency for inhibition to be slower in stellate cells, this difference did not reach significance.

Synaptic depression in fast firing interneurons

Parvalbumin is a Ca²⁺ buffer with slow kinetics that has been shown to mediate paired-pulse depression (Caillard et al., 2000). Although synapses between fast-firing neurons have not been as well studied as inhibition in excitatory cells, synapses between fast firing neurons in hippocampal area CA3 (Kohus et al., 2016) and the dentate gyrus both exhibit short-term depression (Bartos et al., 2001). Similarly, synapses from Pvalb⁺ neurons in the medial septum diagonal band of Broca (MS-DBB) onto CA1 stratum oriens interneurons and from local hippocampal Pvalb⁺ interneurons onto CA1 pyramidal neurons both exhibit short-term depression (Yi et al., 2021). Previously, different subtypes of Pvalb⁺ CA1 pyramidal neurons were found to have different paired pulse ratios (Maccaferri et al., 2000) in synapses onto principal cells, with the axo-axonic cell exhibiting more paired-pulse depression (PPD) than in basket cells and bistratified cells. In area CA3, the synapses of axo-axonic cells and Pvalb⁺ basket cells onto the principal cells both exhibit short term depression; carbachol decreases the inhibitory synaptic current magnitude but eliminates or greatly reduces short-term depression (Szabó et al., 2010). Pvalb⁺ basket cells synapses onto their principal cell targets in the dentate gyrus also exhibit paired pulse depression (Kraushaar and Jonas, 2000). In the striatum, the synapses of fast spiking interneurons onto medium spiny striatal neurons also exhibit short-term depression (Gittis et al., 2010). Synapses from fast firing interneurons onto pyramidal cells in neocortex also exhibit short-term depression, albeit not as much as synapses between pyramidal cells (Galarreta and Hestrin, 1998). This study extends the evidence for short term depression as a characteristic feature of Pvalb⁺ interneurons. However, the short-term depression observed here for Sst⁺ interneurons may not be as general; for example, in mouse layer 4 somatosensory cortex, synapses made by Sst⁺ cells depressed much less than those made by Pvalb⁺ cells, and have a late component of facilitation (Ma et al., 2012). Short term synaptic plasticity may be an important mechanism that allows neurons to detect complex temporal structures by functioning as a memory of events in the past few hundred milliseconds (Motanis et al., 2018).

Figure 1

Synaptic and gap junction connections between Pvalb⁺ interneurons. **A**, Example traces from Pvalb⁺ cells in response to current steps, along with examples of gap junction and synaptic connections (i). Membrane time constant (ii) and spike half-width (iii) measures in Pvalb⁺,

pyramidal and stellate cells, along with a plot of membrane voltage sag and time constant in pyramidal and stellate cells (*iv*). **B**, Distribution and probability of connection types between Pvalb⁺ interneurons. Pie chart indicating the distribution of connection types between Pvalb⁺ interneurons. Plots of pair distance as a function of synaptic connection type (none, one-way, two-way; *ii*) and connection probability of pairs between 20 and 250 μ m (*iii*). Peak inhibitory synaptic amplitude in one-way and two-way connected Pvalb⁺ interneurons (*iv*). Plot of gap junction probability in unconnected, one-, and two-way connected Pvalb⁺ interneuron pairs at distance between 20 and 125 μ m.

Figure 2

Synaptic connections between Pvalb⁺ interneurons and excitatory cells. **A**, Distribution of connections between Pvalb⁺ and pyramidal cells. Pie chart indicating the distribution of connection types between Pvalb⁺ and pyramidal cells. Plots of pair distance as a function of synaptic connection type (none, one-way, two-way; *ii*) and connection probability between 20 and 250 μ m (*iii*). Peak inhibitory synaptic amplitude in one-way and two-way connected Pvalb⁺ interneurons (*iv*). **B**, Distribution of connections between Pvalb⁺ and pyramidal cells. Pie chart indicating the distribution of connection types between Pvalb⁺ and pyramidal cells. Plots of pair distance as a function of synaptic connection type (none, one-way, two-way; *ii*) and connection probability between 20 and 250 μ m (*iii*). Peak inhibitory synaptic amplitude in one-way and two-way connected Pvalb⁺ interneurons (*iv*).

Figure 3

Synaptic inhibitory properties from Pvalb⁺ interneurons. **A**, Comparison of peak amplitude (*i*), inhibitory decay time constant (*ii*), inhibitory rise time constant (*iii*), and delay time (*iv*). **B**, Distributions of inhibitory synaptic amplitude in Pvalb⁺ interneurons and excitatory cells (stellate and pyramidal cells were pooled together). Histograms were fit with a single exponential function.

Figure 4

Synaptic and gap junction connections from Sst⁺ interneurons. **A**, Membrane time constant (*i*) and spike half-width (*ii*) measures in Sst⁺ and pyramidal cells. **B**, Distribution of connection types and probability between Sst⁺ interneurons. Pie chart indicating the distribution of connection types between Sst⁺ and pyramidal cells. Plots of pair distance as a function of synaptic connection type (none, one-way, two-way; *ii*) and connection probability between 20 and 150 μ m (*iii*). Peak inhibitory synaptic amplitude in one-way and two-way connected Sst⁺ interneurons (*iv*). **C**, Distribution of connections between Sst⁺ and pyramidal cells. Pie chart indicating the distribution of connection types between Sst⁺ and pyramidal cells. Plots of pair distance as a function of synaptic connection type (none, one-way, two-way; *ii*) and connection probability between 20 and 150 μ m (*iii*). Peak inhibitory synaptic amplitude in one-way and two-way connected Sst⁺ interneurons (*iv*).

Figure 5

Synaptic inhibitory properties from Sst⁺ interneurons. **A**, Comparison of peak amplitude (*i*), inhibitory decay time constant (*ii*), inhibitory rise time constant (*iii*), and delay time (*iv*). **B**,

Distributions of inhibitory synaptic amplitude in Sst⁺ interneurons and pyramidal cells. Histograms were fit with a single exponential function.

Figure 6

Synaptic inhibitory properties in interneurons (**A**) and excitatory cells (**B**). **A, B**, Comparison of peak amplitude (i), inhibitory decay time constant (ii), inhibitory rise time constant (iii), and delay time (iv) in synaptic connection in interneurons (**A**) and excitatory cells (**B**).

Figure 7

Synaptic inhibitory depression from Pvalb⁺ cells in Pvalb⁺, pyramidal, and stellate cells. **A-C**, Measures of synaptic inhibitory depression in response to 200 ms pre-synaptic pulse trains between 5 and 200 Hz in Pvalb⁺ (**A**), pyramidal (**B**), and stellate (**C**) cells. For each cell, synaptic inhibitory amplitude was normalized by the peak amplitude of the first response during the train (ii). **D**, Plot of average normalized synaptic inhibitory amplitude at the end of different pulse trains (i). Decay time constant associated with synaptic depression in response to 50 Hz pulse trains in Pvalb⁺, pyramidal, and stellate cells (ii). Relative steady-state value of peak inhibitory synaptic amplitude in response to 50 Hz pulse trains in Pvalb⁺, pyramidal, and stellate cells.

Figure 8

Synaptic inhibitory depression from Sst⁺ cells in Sst⁺ and pyramidal cells. **A, B**, Measures of synaptic inhibitory depression in response to 200 ms pre-synaptic pulse trains between 5 and 200 Hz in Sst⁺ (**A**) and pyramidal (**B**). For each cell, synaptic inhibitory amplitude was normalized by the peak amplitude of the first response during the train (ii). **C**, Plot of average normalized synaptic inhibitory amplitude at the end of different pulse trains (i). Decay time constant associated with synaptic depression in response to 50 Hz pulse trains in Pvalb⁺, pyramidal, and stellate cells (ii). Relative steady-state value of peak inhibitory synaptic amplitude in response to 50 Hz pulse trains in Sst⁺ and pyramidal cells.

Figure 9

Modeling synaptic depression. The fits of inhibitory amplitude depression in response to 50 Hz trains of presynaptic stimulation are shown for selected examples for each synapse type. Data points are shown as filled circles. The values predicted by the best fit models are given by other shapes and colors. **A**, Pvalb-Pvalb only includes depression. **B**, Pvalb-Pyr includes both temporal summation and depression. **C**, Sst-Pyr included both temporal summation and depression. **D**, Sst-Sst includes only depression. **E**, Pvalb-Stel included both temporal summation and depression. This was the smallest data set and only the two traces shown had a good enough fit according to the criterion given in the Methods. **F**, Parameter summary. Parameter values are coded by synapse type: Pvalb-Pvalb (blue squares), Pvalb-Pyr (red triangles), Sst-Pyr (upside down orange triangles), SST-SST (green plus signs) and PV-STL (purple X's). Left. Time constant for recovery from synaptic depression. Right. Fraction of available transmitter released by each presynaptic spike.

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Table 1

	PV-PV (n=36)	PV-PYR (n=15)	PV-STL (n=2)	SST-PYR (n=7)	SST-SST (n=11)
U_{SE}	0.345±0.019 (0.166,0.702)	0.261±0.023 (0.087,0.433)	0.317±0.024 (0.284,0.350)	0.183±0.040 (0.061,0.415)	0.378±0.37 (0.197,0.636)
τ_r	105.4±5.5 (49.7,202.9)	104.4±6.6 (74.9,158.8)	96.3±17.0 (72.2,120.4)	112.2±15.9 (55.2,163.8)	96.4±9.4 (47.3,155.5)

Synaptic Depression Model Parameter Statistics. The mean value and the standard error of the mean are given for each parameter value, with n indicating the number of data sets that fit the R^2 criterion given in the methods and were included in Figure 9F and this table.