

CELL TYPE SPECIFIC CONNECTIONS FROM PRIMARY MOTOR TO PRIMARY SOMATOSENSORY CORTEX

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Anatomical studies have shown that primary somatosensory (S1) and primary motor (M1) cortices are reciprocally connected. The pathway from primary motor cortex (M1) to primary somatosensory cortex (S1) is thought to influence activity in S1 by conveying a general modulatory signal and/or a copy of the motor command. In these studies, we investigated M1 synaptic inputs to S1 by injecting an AAV virus containing channelrhodopsin-2 and a fluorescent tag into M1. Consistent with previous results, we found labeling of M1 axons within S1 that was most robust in the deep layers and in L1. We recorded *in vitro* from excitatory neurons and two classes of inhibitory interneurons, fast-spiking and somatostatin-expressing inhibitory interneurons. All three cell types had a high probability of receiving direct excitatory M1 input, with both excitatory and inhibitory cells in L4 being the least likely to receive input from M1. Disynaptic inhibition was observed frequently, indicating that M1 recruits substantial inhibition within S1.

A subpopulation of pyramidal neurons in layers 5 and 6 received especially strong input from M1, suggesting M1 differentially contacts classes of pyramidal neurons, such as those projecting to different sensorimotor centers at cortical and subcortical levels. We tested this hypothesis by combining optogenetic techniques to specifically label M1 synaptic inputs to S1 and retrograde tracing to identify specific populations of projection neurons in infragranular layers of S1. We determined that both the intrinsic properties and the magnitude of M1 input to an S1 pyramidal neuron is highly dependent on its projection target.

Overall, our results suggest that activation of M1 evokes within S1 a general increase in excitatory and inhibitory synaptic activity that could contribute in a layer-specific manner to state-dependent changes in S1. Our results further indicate that M1 may specifically engage sub-circuits within S1 in order to differentially regulate particular downstream cortical and subcortical processing centers.

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PREFACE

The reason I decided to come to the University of Pittsburgh for graduate school was simple: I wanted to learn how to become a scientist. I am very grateful to the Center for Neuroscience and the Neurobiology department for creating a wonderful environment to help their students to learn and grow into neuroscientists.

The people I would like to thank first are my advisors, Dr. Erika Fanselow and Dr. Dan Simons. I am so grateful that Dr. Fanselow took me on as a graduate student, the dedication of her time and energy to my scientific career has meant so much to me. Her encouragement and support allowed me to continue to grow and pursue new things in the lab. To Dr. Simons, I feel privileged to have the opportunity to learn from such an amazing scientist. Perhaps even more importantly, Dr. Simons is a wonderful teacher and I have learned many things from him that I plan to take with me as I move forward.

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1.0 INTRODUCTION

During exploration, rats and mice use their whiskers to investigate novel objects and environments, in much the same way primates use their fingertips to discern the size, texture, and shape of objects (Carvell and Simons, 1990). This behavior requires the integration of sensory and motor information, as rodents actively sweep their whiskers back and forth in a motor behavior known as ‘whisking’ in order to touch and palpate objects. How this behavior is governed by sensory and motor systems within the central nervous system is a matter of active research.

A detailed understanding of the underlying structure between the sensory and motor pathways is not well understood, making it difficult to determine how these two systems interact dynamically with one another during active touch. In this thesis we focus on the projection from primary motor cortex (M1) to primary somatosensory cortex (S1), examining M1 synapses onto specific cell types in S1. The first part of this introduction will focus primarily on the rodent vibrissal sensory systems and sensory processing within S1, whereas the second part will incorporate information about the motor system and its connections with the sensory pathways to provide an overall picture of how sensorimotor integration is organized within the rodent somatosensory system.

1.1 SOMATOSENSORY SYSTEM IN RODENTS: THE WHISKER-BARREL SYSTEM

Whisker-related sensory information ascends through several stages in the brain, beginning with peripheral innervation of the whisker pad, through the brainstem and thalamus, before finally reaching the cortex. Although the fundamental sensory processing seems to occur as information passes from peripheral to central centers, additional structures are interconnected with these sensory pathways, resulting in a series of converging and diverging pathways that can form direct and indirect ‘loops’ between and among areas. The whisker-barrel system as a whole is comprised of multiple, parallel pathways that have distinct input-output organization and likely serve different roles in processing of whisker-related sensory input. Much progress has been made in deciphering the contribution of each structure to sensory processing, although an overall understanding remains elusive in large part because we have yet to elucidate critically important details of the circuitry.

1.1.1 Lemniscal pathway

The primary whisker-related sensory pathway, known as the lemniscal system (Figure 1.1), begins at the whisker follicle, which is directly innervated via the infraorbital nerve by primary afferent neurons residing in the trigeminal ganglion. The trigeminal ganglion cells carry information to the brainstem, where they synapse onto neurons in the principal nucleus of the trigeminal nuclei (PrV). From PrV, signals are transmitted to the ventral posterior medial (VPM) thalamic nucleus. Finally, VPM axons innervate the primary somatosensory cortex (S1).

Topographic representations of the whisker pad that are anatomically well defined (e.g. cortical ‘barrels’) are found at each stage in the lemniscal pathway.

Because of this topography, neurons throughout the lemniscal pathway respond robustly when a sensory stimulus is applied to individual whiskers. Primary afferent trigeminal neurons respond at a short-latency to deflection of a single whisker and have a strong preference for the direction, amplitude, and velocity of the whisker deflection, indicating the ability of the system to reliably encode the mechanical details of sensory stimuli applied to the animal’s face (Zucker and Welker, 1969; Jones et al., 2004; Kwegyir-Afful et al., 2008). Neurons in PrV that project to VPM also exhibit robust whisker responses and directional tuning, but compared with trigeminal neurons they have larger receptive fields although they still respond most robustly to a principal whisker (Minnery and Simons, 2003). The response properties of VPM neurons are similar to those of PrV neurons, the main source of afferent drive to VPM, except that VPM neurons have lower response magnitude to whisker deflections (Minnery et al., 2003). Sensory responses undergo a substantial transformation from VPM to S1, as layer 4 excitatory neurons in S1 barrels have less robust whisker-evoked sensory responses, smaller receptive fields, and considerably less or no amplitude sensitivity compared to VPM neurons (Simons and Carvell, 1989; Bruno and Simons, 2002; Pinto et al., 2003).

Throughout the lemniscal system, information from whiskers is relayed in a spatially and temporally precise manner, with response transformations occurring at each processing center. Although we have learned much about the anatomy and response properties of each structure, understanding the overall function of the system and its regulation by connections with other pathways (see next section) is still incomplete.

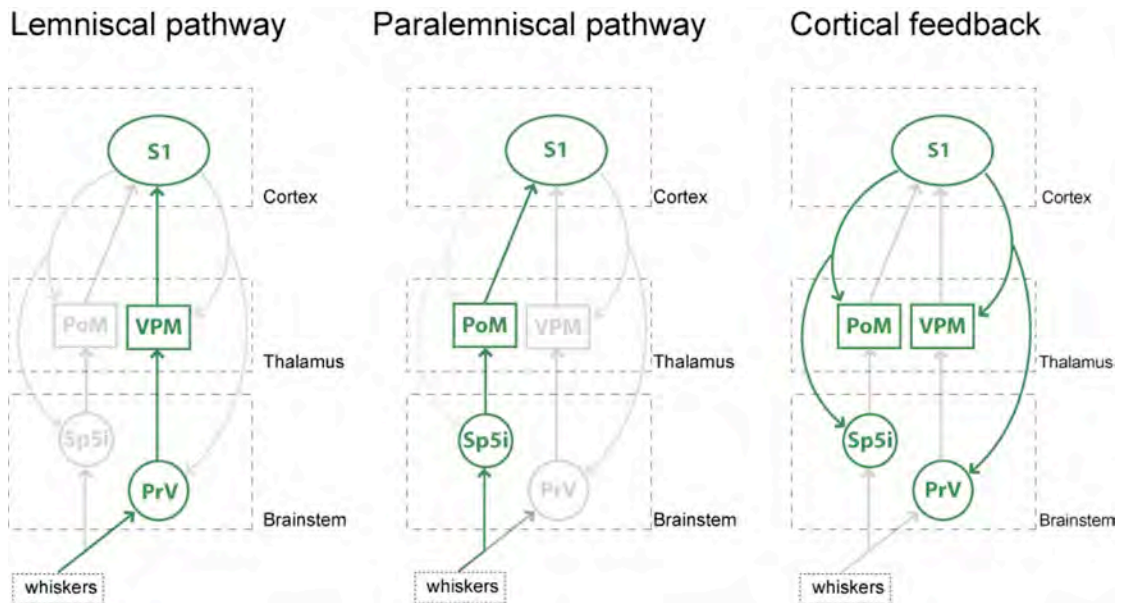


Figure 1.1 Sensory pathways of the whisker-barrel system. Schematic illustrates the two parallel, ascending whisker-related sensory pathways (*left and middle*) and the cortical feedback projections onto subcortical structures (*right*).

1.1.2 Paralemniscal pathway

A second system involved in whisker processing, known as the paralemniscal system, also travels via the brainstem, thalamus, and cortex (Figure 1.1). Axons from trigeminal ganglion neurons synapse in another trigeminal nucleus in addition to PrV, the interpolaris division of the spinal trigeminal nucleus (Sp5i). Sp5i projects to the posteromedial (PoM) thalamic nucleus, and PoM neurons project to S1 as well as to primary motor cortex (M1; see later section).

In contrast to the lemniscal pathway, neurons in the paralemniscal system do not faithfully encode details of sensory stimuli applied to the whisker pad. Neurons in Sp5i that project to the thalamus have large, multi-whisker receptive fields (Jacquin et al., 1986; Furuta et al., 2010). In PoM, despite receiving substantial input from Sp5i, thalamic neurons respond weakly to whisker deflection and at long latencies (Diamond et al., 1992a; Masri et al., 2008). In fact, in contrast to neurons in VPM, which derive most of their sensory properties from ascending brainstem input, sensory responses of PoM neurons instead are thought to depend on cortical input from S1 (Diamond et al., 1992b).

Due to these unusual features, the role of the paralemniscal pathway in sensory processing has remained largely unclear. Activity of neurons in Sp5i and PoM is highly state dependent, through mechanisms that involve cholinergic modulation (Timofeeva et al., 2005; Masri et al., 2006), inhibitory projections via the zona incerta (Trageser and Keller, 2004; Lavallee et al., 2005; Trageser et al., 2006), and cortical feedback (Guillery and Sherman 2011; see next sections). The pathway has also been implicated to play an important role in processing high intensity or pain information in models of central pain syndrome (Masri et al., 2009). The function of the paralemniscal pathway may therefore depend on the behavior state of the animal.

Part of the uncertainty may reflect a more complex anatomical organization than the straightforward whisker-to-barrel pathway of the lemniscal system.

1.1.3 Cortical feedback

In addition to lemniscal and paralemniscal afferent pathways, primary sensory cortices send projections subcortically, innervating the same sensory structures they receive afferent input from (Figure 1.1). In the somatosensory system, S1 sends corticothalamic feedback to both VPM and PoM. S1 also projects to both PrV and Sp5 in the brainstem, as well as to other midbrain and brainstem structures including the superior colliculus, pons, and the spinal cord (White and DeAmicis, 1977; Wise and Jones, 1977; Welker et al., 1988; Killackey et al., 1989). S1 is therefore capable of providing cortical modulation of ascending sensory information at even initial stages of sensory processing.

Corticothalamic feedback from S1 comprises two different pathways, one originating from layer 5 (L5) and the other from layer 6 (L6; see also next section). L5 corticothalamic neurons innervate only PoM, whereas cells in L6 project to VPM or PoM, with a small subset projecting to both thalamic structures (Bourassa et al., 1995). Anatomical and functional data suggest that these two pathways have different roles in modulating thalamic activity. S1 cells in L5 are corticofugal neurons that send a collateral to PoM as they project to the brainstem (Deschenes et al., 1994). These axon collaterals form large ‘giant’ terminals, each containing multiple synapses, onto the proximal dendrites of thalamic neurons in PoM (Hoogland et al., 1991). Stimulation of L5 in S1 evokes large amplitude excitatory currents in PoM neurons, which are highly depressing upon repetitive stimulation (Reichova and Sherman, 2004; Groh et al., 2008). In contrast, L6 corticothalamic neurons synapse onto the distal dendrites of thalamic

neurons in VPM and PoM (Hoogland et al., 1991). These synapses are small, and the excitatory responses in VPM and PoM neurons are weak but facilitating (Reichova and Sherman, 2004; Landisman and Connors, 2007).

These results in the somatosensory system are consistent across primary and higher-order thalamic nuclei in other sensory systems and may reflect two different functions for L5 versus L6 corticothalamic projections. It has been proposed that corticothalamic inputs from L5 act as ‘driver’ and L6 corticothalamic input as ‘modulator’ inputs (reviewed in Sherman 2005). When viewed from a hierarchical perspective, the L5 corticothalamic pathway may be considered as a “feedforward” pathway, and L6 corticothalamic inputs to be a “feedback” system (Hoogland et al., 1991).

Despite growing evidence regarding the anatomy and synaptic organization of corticothalamic feedback, the contribution of each corticothalamic pathway in sensory processing is unclear. When S1 is inactivated the sensory responses of PoM neurons are largely eliminated; in contrast, the fundamental response properties of VPM neurons remain largely unaffected when S1 is inactivated (Diamond et al., 1992b; Ghazanfar et al., 2001). Interestingly, the majority of L6 corticothalamic neurons that project to VPM are silent *in vivo*, exhibiting no spontaneous or whisker-evoked responses (Swadlow, 1989; Swadlow and Hicks, 1996), although they do possess subthreshold whisker-related receptive fields (Kwegyir-Afful and Simons, 2009). Pharmacologically increasing the activity of neurons in L6 of S1 results in enhanced sensory responses in aligned barreloids and suppression of activity in non-aligned barreloids (Temereanca and Simons, 2004). This suggests that when engaged, cortical feedback to VPM does modify ascending sensory inputs in thalamocortical circuits. L6 corticothalamic neurons may be activated by long-range cortical inputs that target the deep layers, such as M1 (Lee et al.,

2008), although the exact circuit mechanisms are still unknown. This is one of the issues we will address in this thesis.

Corticofugal neurons in S1 send projections to both Sp5i and PrV in the brainstem (Wise and Jones, 1977; Welker et al., 1988). Little is known about the anatomy or function of the projection from S1 to PrV. The projection from S1 to Sp5i is topographically organized, and S1 corticotrigeminal neurons specifically contact multi-whisker neurons in the corresponding barrelette within Sp5i (Welker et al., 1988; Furuta et al., 2010). S1 is therefore situated to modulate the earliest stages of afferent sensory processing in both the paralemniscal and lemniscal pathways via its projections to Sp5i and PrV, respectively.

1.2 LAMINAR AND CELLULAR PROPERTIES OF PRIMARY SOMATOSENSORY CORTEX

Primary sensory cortices, including S1, share a common, ‘canonical’ columnar organization (reviewed in Douglas and Martin 2004). In whisker-barrel cortex, a cortical column is defined morphologically as the ‘barrel’ structure within L4 as well as the supragranular layers above the barrel and the infragranular layers below. Functionally, neurons within a barrel-related column respond maximally or only to the same principal whisker, which corresponds anatomically to the layer 4 barrel (Simons, 1978). Processing of complex information from the whisker array is accomplished by interactions within and among these columnar entities.

1.2.1 Basic laminar organization

As discussed above, S1 receives thalamocortical input via the lemniscal pathway from VPM. Thalamocortical terminals from VPM heavily innervate S1 within layer 4 (L4) where they terminate densely within the topographically aligned barrel (Bernardo and Woolsey, 1987; Jensen and Killackey, 1987; Land et al., 1995). Neurons in L4 project to the superficial laminae, layers 2 and 3 (L2/3), above the barrel. Pyramidal neurons in L2/3 are primarily corticocortical, and have an axon that either projects horizontally to superficial layers of other cortical areas, and/or descends into L5 within the same barrel-related column (Feldmeyer et al., 2006). L5 contains a diversity of pyramidal neurons, the majority of which project out of the cortex to the striatum, thalamus, and/or brainstem, among other targets (Wise and Jones, 1977; Killackey et al., 1989). Finally, L6 receives some excitatory input from L4 and L5 within the same column, but most local excitatory inputs are from other L6 neurons (Zarrinpar and Callaway, 2006; Llano and Sherman, 2009; Tanaka et al., 2011). Approximately half of L6 neurons project to either VPM or PoM, as discussed above, and the other half project to other cortical areas, such as M1 or S2.

In addition to thalamocortical input from VPM, S1 is the recipient of many other extrinsic sources of excitatory input from both subcortical and cortical sources. Input from non-VPM external sources (e.g. PoM) largely avoids barrel centers, and instead targets either superficial and/or deep layers in S1 or the inter-barrel septa in layer 4. Thalamocortical axons from PoM neurons heavily target L5A and L1 (Koralek et al., 1988; Wimmer et al., 2010). S1 is reciprocally connected with other cortical areas, in particular secondary somatosensory cortex (Carvell and Simons, 1987) and primary motor cortex (M1; see next section), that in turn send strong projections to L5 and L6 in S1 (Zhang and Deschenes, 1998).

1.2.2 Cellular organization within S1

Each cortical layer is populated by a diversity of cell types. The intrinsic and network properties of these neurons form the basis for the sensory transformations that occur within each layer. Cortical neurons can be broadly classified as either excitatory or inhibitory neurons, with approximately 80% of cortical cells being excitatory and the remaining 20% inhibitory. Excitatory neurons are the main projection neurons of the cortex and are responsible for transmitting information locally, intracortically, and subcortically. Cortical inhibitory neurons usually project locally, either within or across layers in S1. The role of each specific cell type in cortical processing is still unclear, however a wealth of information exists describing the morphology, physiology, and synaptic properties of the different types of cortical neurons.

Excitatory neurons: Each cortical layer in S1 contains a different composition of excitatory neurons (Lefort et al. 2009; reviewed in Feldmeyer 2012). Excitatory neurons in L4 are unique as they are primarily spiny stellate neurons, having a round soma and radially extending dendritic processes (Lubke et al., 2000). These cells receive primarily thalamocortical input from VPM and recurrent excitatory inputs from other L4 neurons (Petersen and Sakmann, 2000); most L4 spiny stellate cells then project to superficial layers. In contrast, the majority of excitatory neurons in cortical layers 2/3, 5, and 6 exhibit pyramidal neuron morphology, with a triangular-shaped soma and prominent apical dendrite that extends upwards toward superficial layers. Pyramidal neurons in each layer have a rich diversity of features that are often dependent on the projection target of the neuron. The axonal target of a pyramidal neuron is, arguably, its defining characteristic as the morphology, physiology, and circuit behavior all depend on the cell's projection target (reviewed in Brown and Hestrin 2009).

Pyramidal neurons in L2/3 and upper L5, known as L5A, are primarily corticocortical neurons that project either locally within S1 and/or cortico-cortically to contralateral S1, M1, or S2 (Koralek et al., 1990; Schubert et al., 2006). Pyramidal neurons in L2/3 and L5A are classified as thin-tufted pyramidal neurons with an apical dendrite that branches in L1 (Manns et al., 2004; Shepherd and Svoboda, 2005). Recordings from *in vitro* preparations show that both L2/3 and L5A thin-tufted pyramidal neurons typically demonstrate regular-spiking (RS) electrophysiology characteristics (Gottlieb and Keller, 1997; Schubert et al., 2006).

Lower L5, or L5B, contains sub-populations of pyramidal neurons that project to a number of different cortical or subcortical targets. Many pyramidal neurons in L5B are among the largest pyramidal neurons in S1 and are morphologically classified as ‘thick-tufted’ pyramidal neurons due to their thick apical dendrites that arborize within L1. L5B pyramidal neurons can be classified as either RS cells or as cells that intrinsically burst (IB) upon depolarization (Agmon and Connors, 1989; Chagnac-Amitai et al., 1990). L6 pyramidal neurons have small, pyramidal shaped somas with an apical dendrite that extends upwards but terminates in L4, not L1 (Zhang and Deschenes, 1997).

Inhibitory interneurons: Cortical inhibitory interneurons comprise an especially diverse population of neurons. Inhibitory interneurons can be characterized on the basis of their morphology, physiology, biochemical composition, and/or subcellular axonal target (Markram et al., 2004; Ascoli et al., 2008). Despite this heterogeneity, expression of particular protein markers allows for classification of almost all cortical inhibitory interneurons into one of the following three groups: parvalbumin-expressing (PV), somatostatin-expressing (SOM), or 5HT3a-expressing (5HT3aR; Rudy et al. 2011). All three types of inhibitory interneurons are found throughout cortical layers 2-6, although the relative abundance of each type varies by layer

(Lee et al., 2010). Of these three groups, PV and SOM interneurons have been most extensively described; our understanding of 5HTaR neurons is more limited, and they will not be further discussed here.

PV and SOM inhibitory interneurons have many differences, but one of the most functionally relevant may be the differential subcellular location of their inhibitory inputs onto pyramidal neurons. PV interneurons have basket cell morphology, with multipolar dendritic trees and an axon that ramifies extensively within the same layer (Kawaguchi, 1993). Their axon innervates the soma and proximal dendrite of nearby pyramidal neurons (Somogyi et al., 1983; Wang et al., 2002). On the other hand, though SOM interneurons are also multipolar and have locally arborizing dendrites, their axons travel vertically and synapse within L1 on the distal apical dendrites of pyramidal neurons (Kawaguchi and Kubota, 1996; Wang et al., 2004). Therefore, within a cortical column PV and SOM cells constitute primary sources of somatic and dendritic inhibition, respectively, onto pyramidal neurons.

PV and SOM inhibitory interneurons have unique cellular and synaptic properties that are likely related to their function *in vivo*. PV cells (hereafter referred to as ‘fast-spiking’ or FS cells) can be distinguished on the basis of their narrow, or ‘fast-spiking’, action potential waveform. FS cells are less excitable than SOM interneurons based on their intrinsic properties as they have lower input resistance, more hyperpolarized resting membrane potentials, and higher voltage thresholds for action potential initiation (Beierlein et al., 2003; Fanselow et al., 2008). FS cells also receive larger unitary inputs than SOM cells from thalamocortical axons and local corticocortical neurons (Bartley et al., 2008). However, upon repetitive stimulation, excitatory synapses onto FS cells exhibit strong synaptic depression, whereas excitatory synapses onto SOM interneurons show synaptic facilitation (Reyes et al., 1998; Beierlein et al., 2003; Fanselow

et al., 2008). An issue that we address in this thesis is whether these cell types differ similarly with respect to their M1 synaptic inputs.

While extensive work has characterized the local connectivity patterns of FS and SOM interneurons (for review see Thomson and Lamy 2007), much less is known about the extrinsic sources of excitatory input to inhibitory interneurons, with the exception of thalamocortical input from VPM. For example, do inhibitory interneurons receive direct, long-range excitatory corticocortical inputs? If so, do the same synaptic properties apply to these inputs as with local excitatory and thalamocortical inputs? Here, we have addressed these questions with regard to one source of long-range cortical input, that from M1.

1.3 INTEGRATION OF SENSORY AND MOTOR PATHWAYS IN THE RODENT VIBRISSAL SYSTEM

1.3.1 Structure and function of the rodent vibrissal motor system and the control of whisking behavior

The motor cortex of the rodent contains a disproportionately large region representing the face and whiskers, illustrating the importance of whisker movements in behavior. Electrical stimulation of vibrissal M1 can evoke whisker movements, even upon activation of only a single M1 neuron (Brecht et al., 2004). Anatomical tracing has revealed connections between vibrissal M1 and several subcortical structures implicated in the generation and maintenance of motor behavior (Figure 1.2), including the ventral lateral thalamus, superior colliculus, caudate and

putamen, pons, and multiple brainstem nuclei (Porter and White, 1983; Miyashita et al., 1994; Hattox et al., 2002).

Physiological evidence indicates that activity in vibrissal M1 is correlated with whisker movement, as the firing rate of M1 neurons increases during whisking bouts. Activity in M1 actually increases just prior to whisking onset, indicating a possible role of M1 in the generation, or initiation, of whisking (Carvell et al., 1996; Friedman et al., 2006). Activity in vibrissal M1 neurons, however, does not obviously correlate with parameters of whisker motion on a cycle-by-cycle basis (Carvell et al. 1996; Friedman et al. 2012; but see also Hill et al. 2011); this could depend on both the area within M1 (Smith and Alloway, 2013) and the types of pyramidal neurons being recorded (Beloozerova et al., 2003). This raises the question of whether vibrissal M1 directly controls individual whisker movements. Rhythmic whisking persists after inactivation of vibrissal M1, although whisking kinetics are altered (Semba and Komisaruk, 1984; Gao et al., 2003). This suggests that rhythmic whisker movements are not directly under control of M1 and are instead generated subcortically by a central pattern generator (CPG) in the brainstem (Gao et al., 2001; Cramer and Keller, 2006; Cramer et al., 2007).

Muscle contractions producing whisker movement are under the control of motor neurons in the facial motor nucleus. Direct M1 projections to motor neurons in the facial nucleus exist, but are sparse (Grinevich et al., 2005). Most M1 corticofugal axons instead innervate other brainstem nuclei, such as the reticular nucleus, that in turn innervate the facial motor nucleus (Miyashita et al., 1994; Hattox et al., 2002). Recent attempts to identify a brainstem CPG network for whisking has focused on the interactions of whisking and other head and neck motions, such as breathing, sniffing, and licking (reviewed in Deschenes et al, 2012). Neurons in a small region of the brainstem, the intermediate band of the reticular formation, were recently

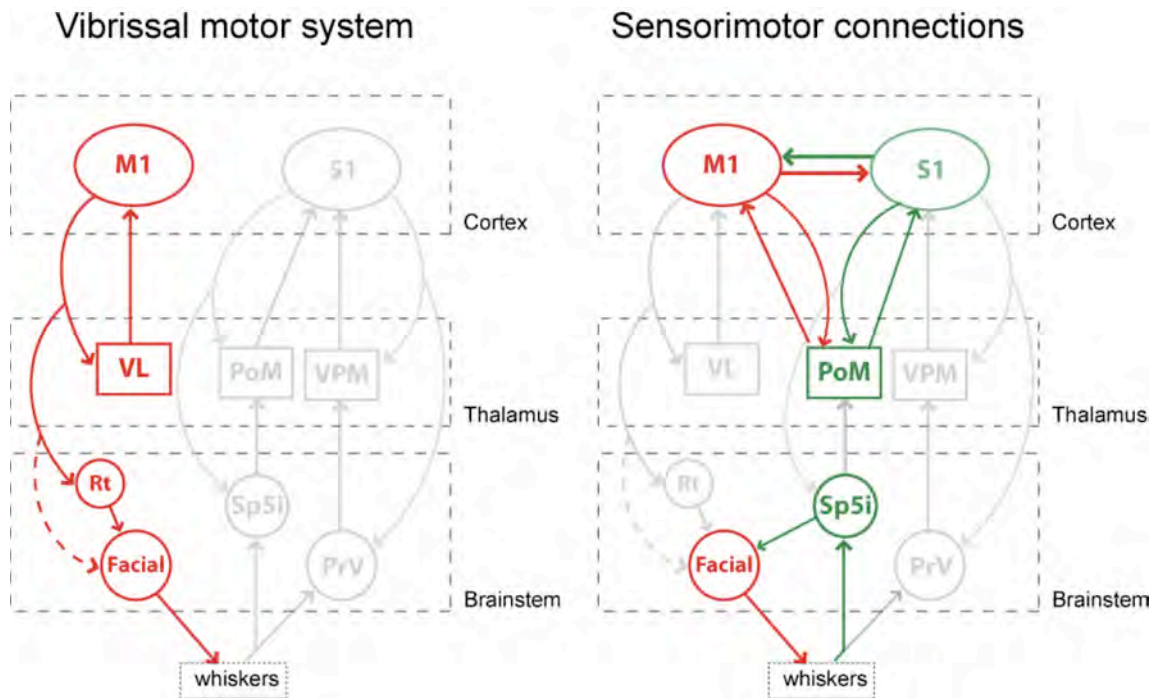


Figure 1.2 Motor and sensorimotor pathways of the whisker-barrel system. Schematic illustrating descending projections of the vibrissal motor system (*left*) and interconnections between the sensory and motor pathways (*right*). Dashed line indicates sparse anatomical connections.

discovered to fire rhythmically during whisking and drive each protraction of the whiskers through their connections with motor neurons in the facial nucleus (Moore et al., 2013). This region receives input from the pre-Botzinger complex, which serves to keep whisking cycles in phase with breathing and sniffing rhythms (Moore et al., 2013). Neurons in the reticular formation also receive input from motor cortex (Hattox et al., 2002), which may allow for cortical modulation of whisking patterns.

The vibrissal motor system therefore appears to function such that cortical control from vibrissal M1 can initiate onset of whisking, through its corticofugal projections to ‘pre-motor’ brainstem nuclei. Rhythmic whisking behavior itself seems to be controlled on a whisk-by-whisk basis via a series of brainstem nuclei that enable the rhythmic behavior of whisking to occur as part of a larger ensemble of head and face movements. Whether whisker motion can be dynamically adjusted on the basis of sensory feedback, either at the brainstem or cortical level, is addressed in the following sections.

1.3.2 Anatomical connections between vibrissal motor and sensory systems

Active sensing requires constant adjustment of whisker motion and position based on continually updated sensory information arriving from the periphery. This interaction is thought to be mediated via sensorimotor ‘loops’, which connect the sensory and motor systems at the level of the brainstem, thalamus, and cortex (reviewed in Kleinfeld et al, 2006).

Connections from Sp5i to the facial motor nucleus (Erzurumlu and Killackey, 1979) form a loop within the brainstem (whiskers → Sp5i → facial nucleus → whiskers; Figure 1.2). The function of this circuit is not clear, although sensory input from Sp5i to the facial nucleus is not

required to activate the brainstem CPG network, as severing the trigeminal nerve does not affect the generation or patterning of whisking (Gao et al., 2001).

M1 and S1 are connected directly via reciprocal corticocortical connections and indirectly via their reciprocal connections with the thalamic nucleus PoM (Figure 1.2). Corticothalamic inputs to PoM are branches of axons traveling to subcortical centers, as mentioned earlier, suggesting that the indirect transthalamic pathway via PoM may integrate information from M1 and S1 regarding subcortical commands (Sherman and Guillery, 2011).

Reciprocal corticocortical connections between M1 and S1 follow a pattern seen commonly throughout the cerebral cortex. That is, S1 innervates primarily the middle layers of M1, L2/3 and L5A (Aronoff et al., 2010), whereas M1 input to S1 avoids the middle layers, instead terminating in L5/6 and L1 (Veinante and Deschenes, 2003). It has been suggested that pathways targeting middle layers are the stronger ‘feedforward’ inputs, and those avoiding middle layers represent a ‘feedback’ projection (Felleman and Van Essen, 1991). According to this hierarchical cortical model, information should flow primarily from S1 to M1, with the M1 to S1 projection being weaker and modulatory. Evidence supporting this model in the barrel system shows that sensory activity in the cortex is propagated from S1 to M1 (Ferezou et al., 2007). S1 may therefore help regulate whisking behavior via its projection to M1. The function of the feedback pathway from M1 to S1 is less clear, and will be further discussed below.

Recent advances in optogenetics technology have allowed for the selective activation of specific populations of synaptic input. This approach has yielded important findings regarding the excitatory circuits that connect S1 and M1. A recent study combined optogenetics with retrograde tracers and found that S1 inputs to M1 are most prevalent in the superficial layers of

M1 (Aronoff et al., 2010; Mao et al., 2011), providing especially strong input to S1-projecting neurons in L2/3 and L5A (Mao et al., 2011).

Both callosal and corticofugal neurons in the infragranular layers of M1 project to S1 (Veinante and Deschenes, 2003). Callosal fibers terminate within L5/6 and L1 of S1, whereas corticofugal neurons only terminate within L1. Individual pyramidal neurons in L2/3 and L5 receive input from M1 on both the basal dendrites, near the soma, and the apical dendrite in L1 (Petreanu et al., 2009). Here, using similar optogenetic techniques, we determine the laminar dependence of M1 input to S1 neurons, and also investigate the differential input from M1 onto several types of excitatory and inhibitory interneurons within S1.

1.3.3 Influence of whisker movements on sensory processing

The animal gathers sensory information either passively, when whiskers incidentally contact an object, or actively when whiskers are motorically brought into contact with an object. The first behavior, passive touch, is a purely sensory process whereas the latter, active touch, requires the integration of motor and sensory systems. Investigators have compared activity in sensory structures during passive versus active touch to try to understand differences in sensory processing that occurs in the two states and hence infer the role(s) of internally generated motor commands to perception. Comparisons have yielded information about how sensory processing is transformed when information is received passively versus accompanied by movement. Furthermore, activity in M1 is state-dependent, as M1 neurons increase their firing prior to and during whisking. However, given that sensory and motor systems interact at multiple levels, the contribution of peripheral versus central mechanisms is not always clear. In addition, transitions to whisking states also change the attentiveness of the animal, indicating a role for

neuromodulator systems in state-dependent changes in sensory processing (Castro-Alamancos, 2004).

One of the most common findings across species is that sensory responses are reduced during the course of movement, a phenomenon known as ‘motor-gating’ (Chapin and Woodward, 1982; Chapman, 1994). In the whisker system, sensory-evoked activity in VPM and S1 is reduced during whisking (Fanselow and Nicolelis, 1999; Ferezou et al., 2006; Lee et al., 2008). Recent work has demonstrated a role for Sp5i in gating sensory responses in PrV, as inactivation of Sp5i abolishes the whisking-related reductions in VPM activity (Lee et al., 2008). This suggests that whisking-related sensory activity may be suppressed within the brainstem and this reduction is reflected in VPM and S1. How whisking modulates activity levels in Sp5i, however, is not clear. Sp5i receives input from neuromodulator centers within the brainstem (Timofeeva et al., 2005), as well as top-down cortical input from S1 and S2 (see previous section), either of which could drive state-dependent changes in Sp5i neurons.

Besides reduced activity during whisking, S1 shows other state-dependent changes that appear to originate centrally (Poulet and Petersen, 2008). Recordings from L2/3 pyramidal neurons in S1 *in vivo* demonstrate striking changes in the dynamics of the subthreshold membrane potential. Awake states wherein the animal is alert but not whisking are characterized by low frequency, large amplitude fluctuations that occur synchronously in nearby pyramidal neurons; upon start of whisking membrane potential dynamics de-correlate and transition to higher frequency and lower amplitude fluctuations (Crochet and Petersen, 2006; Poulet and Petersen, 2008). Activity of cortical inhibitory interneurons is also state-dependent, with FS and SOM interneurons exhibiting decreased activity in whisking versus non-whisking states (Gentet

et al., 2010; Gentet et al., 2012). Other inhibitory interneurons, presumably 5HT3aR-expressing interneurons, have increased activity during whisking (Gentet et al., 2010).

The mechanisms governing state-dependent changes in S1 are unknown, but could involve effects of neuromodulator systems (Constantinople and Bruno, 2011), and/or changes in thalamocortical or corticocortical inputs. Of those possibilities, the effects of state-dependent changes in corticocortical activity remain the least understood. This is in part due to a lack of knowledge regarding the detailed microcircuits involved in such connections. Here, we have examined the cell type specific connections of one important cortical input to S1, that coming from M1.

1.4 OVERVIEW

In this thesis, we investigate connections from M1 onto specific cell types in S1. The study described in Chapter 2 tests the hypothesis that inhibitory interneurons are involved in mediating M1 inputs to S1 and, additionally, examines the relative amount of M1 input to different layers in S1. Appendix A contains a previous study by Dr. Fanselow and I on the development of SOM-expressing inhibitory interneurons, which could serve as useful background regarding SOM interneurons. The third chapter builds on findings from Chapter 2, examining whether M1 inputs differentially target S1 pyramidal neurons on the basis of their projection target. Overall, our findings demonstrate M1 inputs synapse broadly with many types of S1 neurons, but that specific sub-circuits exist as a result of greater M1 input to certain types of pyramidal neurons.

2.0 MOTOR CORTEX BROADLY ENGAGES EXCITATORY AND INHIBITORY NEURONS IN SOMATOSENSORY BARREL CORTEX

2.1 ABSTRACT

Anatomical studies have shown that primary somatosensory (S1) and primary motor (M1) cortices are reciprocally connected. The M1 to S1 projection is thought to represent a modulatory signal that conveys motor-related information to S1. Here, we investigated M1 synaptic inputs to S1 by injecting an AAV virus containing channelrhodopsin-2 and a fluorescent tag into M1. Consistent with previous results, we found labeling of M1 axons within S1 that was most robust in the deep layers and in L1. Labeling was sparse in L4 and was concentrated in the inter-barrel septa, largely avoiding barrel centers. In S1, we recorded *in vitro* from regular-spiking excitatory neurons and fast-spiking and somatostatin-expressing inhibitory interneurons. All three cell types had a high probability of receiving direct excitatory M1 input. Both excitatory and inhibitory cells within L4 were the least likely to receive such input from M1. Disynaptic inhibition was observed frequently, indicating that M1 recruits substantial inhibition within S1. Additionally, a subpopulation of L6 regular-spiking excitatory neurons received exceptionally strong M1 input. Overall, our results suggest that activation of M1 evokes within S1 a bombardment of excitatory and inhibitory synaptic activity that could contribute in a layer-specific manner to state-dependent changes in S1.

2.2 INTRODUCTION

During active touch sensory and motor-related signals are thought to interact in a fashion that regulates on-going exploration and stimulus discriminability. In the rodent whisker system the motor and somatosensory systems are intricately linked at the level of brainstem, thalamus, and cortex (Kleinfeld et al., 2006), providing multiple mechanisms for sensorimotor integration. At the cortical level, primary motor (M1) and primary somatosensory (S1) cortices are reciprocally connected (White and DeAmicis, 1977; Porter and White, 1983), with the S1 to M1 connection thought to represent the “forward” pathway and M1 to S1 to represent the “backward” pathway (Felleman and Van Essen, 1991; Cauller et al., 1998). In keeping with this view, recent studies have demonstrated that the synaptic inputs from S1 to M1 are stronger than those from M1 to S1 (Rocco-Donovan et al., 2011). In addition, sensory-evoked activity is first present in S1 and subsequently propagated to M1 (Ferezou et al., 2007).

Functionally, the role of M1 inputs is hypothesized to play a ‘modulatory’ role within S1 by sending a copy of motor-related information that could alter processing of whisking-related sensory information in S1 (Fee et al., 1997; Kleinfeld et al., 2006; Hill et al., 2011; Friedman et al., 2012). Neurons in M1 are active prior to and during whisking (Carvell et al., 1996; Friedman et al., 2006), at which time S1 neurons exhibit smaller responses to whisker deflection (Chapin and Woodward, 1982; Faselow and Nicolelis, 1999; Ferezou et al., 2007). In addition, pairs of S1 neurons exhibit a reduction in membrane potential correlation relative to non-whisking states (Poulet and Petersen, 2008). The peripheral and/or central origin of many of these state-dependent changes in S1 is unclear; however, given the extensive reciprocal connectivity between M1 and S1, whisking-associated changes in S1 firing could reflect direct M1 to S1 projections.

An understanding of M1-S1 interactions has been hindered by the inability to activate selectively motor cortex inputs to S1. In reciprocally connected neural systems, commonly used techniques involving electrical stimulation may be confounded by inadvertent antidromic activation of cells in the target population via their locally recurrent axons. The recent development of optogenetic tools allows for selective activation of neuronal populations that project to a distant location, permitting investigation of the properties of their synaptic connections. In addition, mapping of connectivity between M1 and S1 has so far focused solely on inputs to excitatory neurons (Petreanu et al., 2009; Mao et al., 2011; Rocco-Donovan et al., 2011). However, inhibitory interneurons can regulate the impact of excitatory inputs between brain areas, such as thalamus to cortex, through feedforward inhibition (Simons and Carvell, 1989; Pouille and Scanziani, 2001; Swadlow, 2003; Gabernet et al., 2005). Inhibitory interneurons comprise a diverse group (Markram et al., 2004; Ascoli et al., 2008), and the effect of inhibition on sensory processing is dependent on the types of inhibitory interneurons involved (Porter et al., 2001; Pouille and Scanziani, 2004; Lee et al., 2012; Wilson et al., 2012).

Here, we utilized an optogenetic approach combined with *in vitro* whole-cell recordings to examine synaptic inputs from M1 onto specific classes of excitatory and inhibitory cells in S1. We focused on three identified types of neurons in S1: regular-spiking (RS) pyramidal neurons, fast-spiking (FS) inhibitory interneurons, and somatostatin-expressing (SOM) inhibitory interneurons. We recorded responses *in vitro* from each of these cell types in S1 evoked by photic activation of M1 terminals expressing channelrhodopsin-2 (ChR2). Our findings indicate a high probability of M1 input to all three cell types. The nature of the inputs is laminar and cell-type specific. Results demonstrate that both inhibitory and excitatory neurons are strongly and

widely recruited by M1, providing circuit-level mechanisms for the regulation of S1 activity during movement-associated activity in M1.

2.3 METHODS

All experiments were carried out in compliance with the University of Pittsburgh School of Medicine animal use policies and were approved by the University of Pittsburgh Institutional Animal Care and Use Committee.

Virus injections

Experiments were conducted using a transgenic mouse line that contains GFP in a subset of somatostatin-expressing GABAergic neurons (“GIN” mice; Oliva et al., 2000). Mice were first anesthetized using isoflurane (1-2%/oxygen), a small craniotomy was performed over primary motor cortex (1.0 mm anterior to and 0.8 mm lateral from bregma), and the adeno-associated virus AAV2/5.CamKII α .hChR2(H134R)-mCherry.WPRE.SV40 (University of Pennsylvania Vector Core; permission from Dr. Karl Deisseroth) was unilaterally pressure injected into primary motor cortex (M1) using a picospritzer. In a single penetration we injected the virus separately into both deep and superficial layers of M1 (0.8 mm and 0.4 mm depth, respectively). A volume of 0.1-0.2 μ l was injected at each depth. At the time of injection the mice were 10-15 postnatal days of age (P10-15).

Verification of M1-to-S1 labeling

At the outset we examined the nature of the ChR2 labeling. Injected animals were

perfused transcardially using 0.1M PBS followed by 4% paraformaldehyde. The brain was then placed in 30% sucrose for 48-72 hours. Next, the brain was frozen and sectioned in either the coronal or tangential plane at 40 μ m using a cryostat.

Initial experiments were done using the AAV2/1.CAG.hChR2(H134R)-mCherry.WPRE.SV40 viral vector. When the tissue was examined we consistently observed ChR2-mCherry labeled cells within S1, suggesting a small percentage of neurons were retrogradely transporting the virus from M1, the injection site, back to S1 (data not shown). Because we wished to examine only M1 inputs to S1 and not possible recurrent collaterals of retrogradely labeled S1 cells, we decided to use AAV2/5.CamKII α .hChR2(H134R)-mCherry.WPRE.SV40 for all of our experiments, which we thought may result in little or no retrograde transport while still producing strong ChR2-expression in M1 neurons. To confirm this, in a subset of animals injected with this viral vector we stained tissue for NeuN as a marker for cell bodies. Free-floating coronal sections were rinsed using 0.1M PBS, incubated in blocking solution (containing 10% donkey serum and 0.3% Triton-X in 0.1M PBS), and then incubated in primary antibody for 18-24 hours (1:1000 anti-NeuN; Chemicon). The tissue was rinsed in 0.1M PBS and subsequently incubated for 2-3 hours in secondary antibody (1:500 donkey anti-mouse Alexa 647; Jackson Immunoresearch). The tissue was placed on slides, coverslipped (Vector Laboratories, Vectashield), and examined using a confocal microscope.

For sections within M1, we determined that about 71% of NeuN+ neurons near the injection site (range 52-85%; n=3 animals) were also positive for ChR2, indicating that they expressed the virus. This included neurons in layers 2-6. This percent decreased with distance from the center of injection site. The promoter we used to drive viral expression, CaMKII α , is largely specific for excitatory neurons (Dittgen et al., 2004; Nathanson et al., 2009). It is not

known whether CaMKII α is preferentially expressed in subpopulations of pyramidal neurons. However, ChR2 was present in the majority of neurons near the injection site, consistent with the assumption that the virus is being expressed non-selectively in M1 pyramidal neurons.

We examined S1 labeling in 8 slices from 3 animals for co-expression of NeuN and ChR2-mCherry. In a count of 4007 NeuN-positive cells distributed across cortical layers, we found 4 that also labeled with mCherry. Interestingly, the 4 retrogradely-labeled neurons were found in layers 2/3. Due to the sparse nature of the retrograde labeling ($\sim 1/1000$ neurons), we concluded that any such labeling with this viral construct would be unlikely to affect our results. All of the data presented in this manuscript were therefore collected from animals injected with the AAV2/5.CamKII α .hChR2(H134R)-mCherry.WPRE.SV40 virus construct.

Preparation of *in vitro* S1 slices

Electrophysiological experiments began a minimum of 3 weeks following virus injection to allow for transport and full expression of the virus. At the time of experiments the animals were 32-51 days of age. Mice were anesthetized with isoflurane, then the brain was removed and placed in ice-cold artificial cerebrospinal fluid (ACSF) containing (in mM): 126 NaCl, 3 KCl, 1.25 NaH₂PO₄, 2 MgSO₄, 26 NaHCO₃, 10 dextrose, and 2 CaCl₂, saturated with 95% O₂-5% CO₂. The tissue was then sliced at 400 μ m in the coronal plane using a vibratome. Slices were incubated at 32° C for 30-45 minutes and then maintained at room temperature until used for recording. Slices containing S1 barrel cortex were identified by the presence of layer IV barrels and a patchy barrel-related pattern of mCherry fluorescence (Figure 2.1B&2A). We recorded primarily from the larger, more medially situated barrels (rows D-E), as this is where fluorescence was typically strongest. We recorded from one to three adjacent barrels per slice.

Recording procedures

Whole cell recordings were performed using glass micropipettes (4–10 M Ω) filled with internal solution containing (in mM): 135 K-gluconate, 4 KCl, 2 NaCl, 10 HEPES, 0.2 EGTA, 4 ATP-Mg, 0.3 GTP-Tris, and 14 phosphocreatine-Tris (pH 7.25, 280-290 mOsm). Biocytin (0.5%) was added to the internal solution in a subset of experiments. Membrane potentials reported here were not corrected for the liquid junction potential. Recordings were conducted at 32°C. When patching, cell-attached seal resistances were 1 G Ω or greater and series resistance after achieving whole cell configuration was 5-20 M Ω . After establishing whole cell configuration, a series of current steps was presented in current-clamp for use in characterization of cell type (see below). Current steps were presented in 20 pA steps, ranging from -100 to 300 pA. Steps were presented 5 seconds apart. In voltage-clamp experiments, series resistances ranged from 10-40 M Ω and were compensated for up to 80%. For EPSC measurements all cells were held at a potential of -80 mV. Data were collected using a Multiclamp 700B amplifier and pClamp10 software (Molecular Devices). Data were collected at a sampling rate of 20 kHz.

Cell type identification

Cells were viewed under infrared-differential interference contrast illumination using a Nikon FN-1 microscope and a Dage IR-1000 CCD camera. In “GIN” mice all of the GFP-expressing cells are inhibitory interneurons that express the neuropeptide somatostatin (SOM), although not all SOM⁺ interneurons are labeled with GFP (Oliva et al., 2000; Ma et al., 2006). Most or all of the GFP-expressing cells in this mouse line are morphologically consistent with Martinotti interneurons (Ma et al., 2006). GFP-expressing neurons are found in layers 2-5 but

not in layer 6 (Ma et al., 2006), preventing us from positively identifying L6 SOM interneurons. All GFP-expressing interneurons were identified by visualization of GFP under epifluorescence illumination (Nikon Intensilight). All neurons identified in this study as “SOM” were labeled with GFP.

Fast-spiking (FS) inhibitory interneurons do not express GFP in this mouse line. Therefore cells with an oval-shaped soma were targeted as putative FS cells and their identity was confirmed based on electrophysiological characteristics (Beierlein et al., 2003; Fanselow et al., 2008). FS cells can exhibit different axonal projection patterns (Helmstaedter et al., 2009); therefore our results may encompass FS cells with different morphologies. Nevertheless, a wealth of both in vitro and in vivo literature exists regarding the role of inhibitory interneurons identified as “fast-spiking” in cortical circuits. Our goal was to place our results regarding the role of FS inhibitory neurons in mediating M1-S1 interactions within the context of these findings.

Regular-spiking (RS) pyramidal cells were targeted for recording based on a triangular-shaped soma. During recording, RS pyramidal cells were similarly characterized based on established electrophysiological criteria (Porter et al., 2001; Beierlein et al., 2003; Hattox and Nelson, 2007). We also recorded from a few pyramidal neurons in L5 that were not regular-spiking, but instead displayed ‘intrinsically bursting’ properties (n=3). These cells showed similar results as the L5 RS cells, and were therefore pooled with the regular-spiking pyramidal neurons.

Laminar definitions

Layer 1 was identified by a low density of cell bodies, and the top of layer 2/3 was defined by the abrupt increase in cell density. The boundary between the bottom of L3 and the top of L4 was identified by the presence of barrels in L4, and the bottom of L4 was indicated by the loss of barrel structure as well as a more diffuse labeling pattern of fluorescent M1 axons. The boundary between layers 5 and 6 was approximated as being roughly half way between the top of L5 and the white matter. In a subset of our data we measured the distance from the pia to the cell body of the recorded cell (see Figure 2.6A). Based on these measurements, our laminar boundaries were very similar to previously published laminar definitions for S1 in the mouse (Hooks et al., 2011).

Optical stimulation procedures

To test for M1 input using optical stimulation, the recorded cell was centered in the field of view. Full-field blue light was delivered through a 40X objective using a 470 nm LED (OptoLED; Cairn Research). Light intensity at the surface of the slice was $\sim 20\text{mW/mm}^2$; light intensity was held constant across all experiments to minimize variability. This intensity is higher than the threshold for channelrhodopsin activation, which in our experiments as well as in previous studies is approximately $8\text{-}12\text{mW/mm}^2$ (Boyden et al., 2005). Keeping the light intensity stronger than the threshold for ChR2 activation allowed for reliable detection of synaptic connections within our experiments. Light intensity was measured using an optical power meter to measure the overall power and then dividing by the surface area of the light spot (to give mW/mm^2). We measured the spatial diameter of the light spot to be $\sim 250\ \mu\text{m}$ with some scattering of light beyond that. Consistent with this measurement, light evoked responses were about 50% of the maximal response when stimulated $250\ \mu\text{m}$ from the neuron's somata. This

indicates the activation range probably extends approximately one barrel-related column in width, or perhaps slightly more. Light stimulation was delivered via TTL pulses using the pCLAMP software. A single stimulation trial consisted of trains of 8-10 pulses (each pulse was 1.0 ms duration) at 1, 10, 20, and 40Hz, with trains separated by >8 seconds to minimize adaptation effects. For each cell 10-30 trials were recorded.

We took multiple steps to limit experimental variability arising from variation in the amount of virus taken up and/or expressed across animals. First, we only recorded from slices exhibiting strong Chr2-mCherry expression that was easily visible under 4X magnification (e.g. Figure 2.2A). Second, we limited recordings to the area in barrel cortex having the highest Chr2-mCherry expression; this was typically one to three medially situated adjacent barrels. Third, we included data only from animals in which at least one significant excitatory response was identified. Fourth, we attempted to sample multiple cells within each slice such that the sample contained a variety of cell types from different layers.

Data analysis

Data were analyzed using in-house programs written in Matlab (The MathWorks, Natick, MA; A. Kinnischtzke). Trials were averaged together, and analyses were performed on averaged voltage traces. Response onset and peak were calculated within a 15ms window following the offset of the light pulse. Response onset was taken as the first of 20 consecutive data points that exceeded a threshold of 1 s.d. above the resting membrane potential. A cell was considered to have received an input if the peak response was greater than 5 times the standard deviation of the resting membrane potential. Excitatory post-synaptic potential (EPSP) and current (EPSC) amplitudes were calculated as the difference between the response onset and the peak response.

In cases where a cell spiked in current-clamp mode, the peak response was taken to be the action potential voltage threshold. For analysis of synaptic dynamics, response amplitudes were calculated using EPSCs, and we used only cells that had a significant response to the first pulse in the train. Response amplitudes were calculated for each pulse then normalized to the first value. To calculate the change in spike probability across a train, we used only cells that spiked at least once across all trials to any pulse in the train.

For comparison of input probabilities between cell types or layers we used a chi-squared test. For all other statistical comparisons, an ANOVA was performed for each data set and *t*-tests were used for post hoc pairwise comparisons. Results are reported as mean \pm SEM.

2.4 RESULTS

Anatomy of the M1-S1 projection

To study synaptic inputs from M1 to identified S1 neurons, we injected an AAV virus containing the channelrhodopsin-2 (ChR2) gene into primary motor cortex (see Methods; Figure 2.1A, left). Experiments were performed a minimum of three weeks following the injection, at which time pyramidal neurons in M1 showed strong expression of ChR2 (Figure 2.1A, right).

Viral

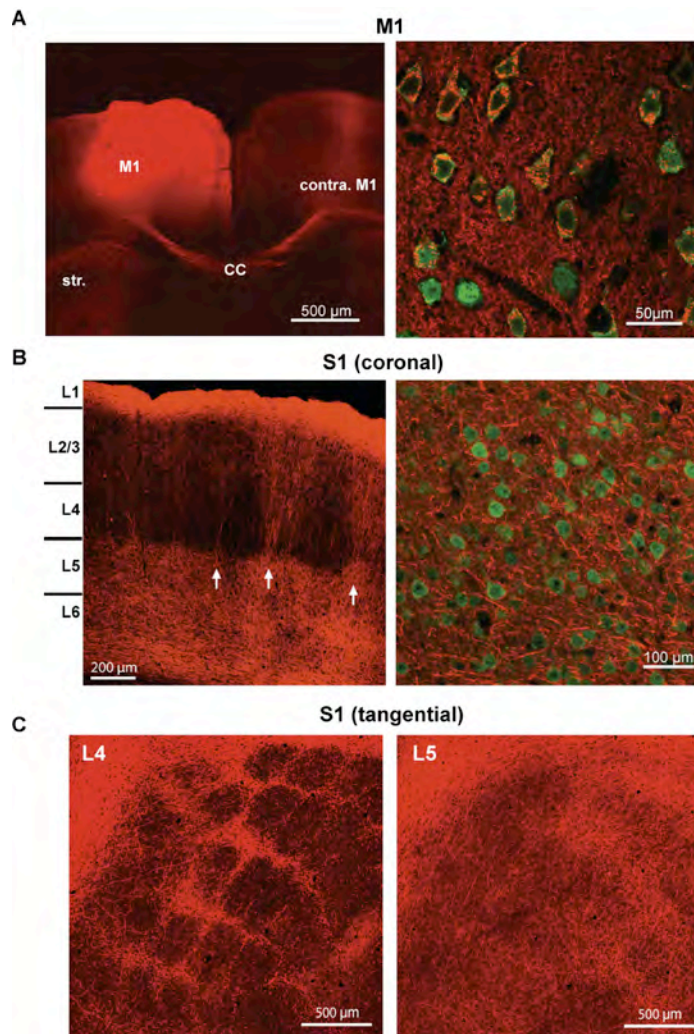


Figure 2.1 Injection of ChR2-mCherry viral vector produces extensive labeling of M1 axons within S1. (A) *Left:* Injection of the ChR2-mCherry viral vector produced robust fluorescent labeling around the injection site, shown here in the coronal plane. Projections from the injection site can be seen in contralateral M1 as well as the striatum. Str=striatum; CC=corpus callosum. *Right:* High power magnification (60x) shows individual neurons, labeled with NeuN (green), co-expressing the ChR2-mCherry virus (red). (B) *Left:* ChR2-mCherry labeling of M1 axons within S1. Labeling pattern is typical of the termination pattern of M1 axons within S1. Arrows indicate septal columns of M1 fibers. *Right:* High power magnification (60x) demonstrates that the ChR2-mCherry virus is only expressed in axons and axon terminals in S1, as no neurons in S1 (labeled with NeuN; green) are co-labeled with ChR2-mCherry (red). (C) S1 section cut in a plane tangential to the pial surface at depths corresponding to L4 (*left*) and L5 (*right*).

expression was primarily located in M1 neurons within the agranular medial field (vibrissal motor cortex) with some in the agranular lateral cortex (Brecht et al., 2006). This produced a stereotypical pattern of axonal labeling within S1 (Figure 2.1B) that was consistent with known patterning of M1 axons (Veinante and Deschenes, 2003; Petreanu et al., 2009). This pattern of labeling was consistent across animals; however we observed some variability between animals in the amount of ChR2 labeling that was present (see Methods). Layers 5 and 6 (L5&6) were characterized by diffuse labeling. At the L4/L5 boundary labeling become concentrated within vertical bands that coursed through L4 and L2/3. Labeling broadened somewhat before becoming extensive and widespread in L1 (Figure 2.1B). The vertical bands of M1 fibers appeared to be concentrated between barrel centers, which were largely devoid of M1 axons. To explore this further we sliced through the S1 barrel field in the tangential plane to highlight barrel versus septal areas. We found that in L4, M1 labeling was concentrated primarily between the barrels, within the septa (Figure 2.1C, left). In deeper layers, as was observed in the coronal slice, M1 axons became more diffuse, however they remained more concentrated under the septa all the way through L5 and L6 (Figure 2.1C, right).

M1 provides input to excitatory and inhibitory neurons in S1

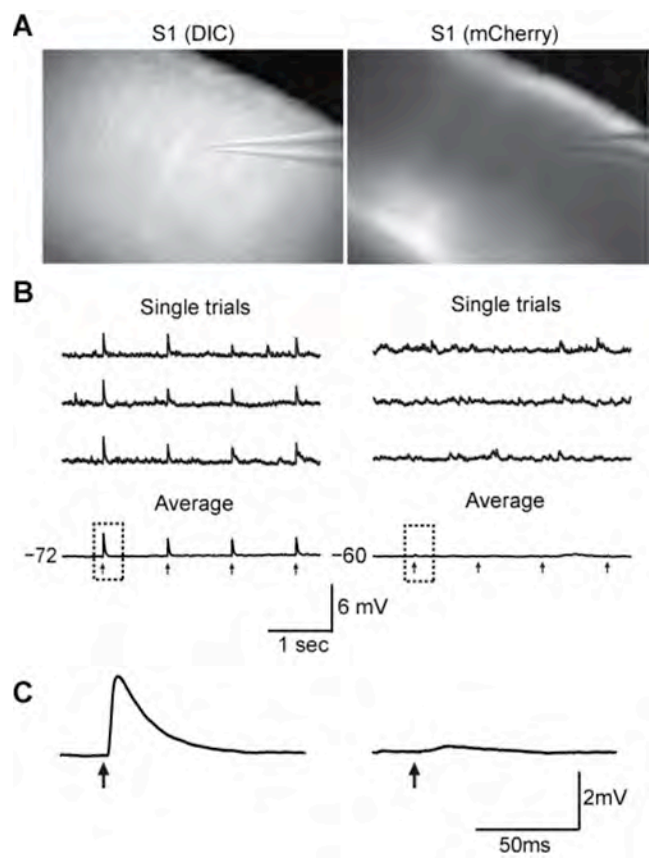


Figure 2.2 Optical stimulation of ChR2-expressing M1 axon terminals (ChR2-M1) results in time-locked excitatory responses in S1 neurons. (A) Left: In vitro brain slice through S1 under DIC at 4X magnification. Right: Same brain slice under mCherry fluorescence shows robust ChR2-M1 labeling in S1. (B) Recording of single S1 pyramidal neurons during optical stimulation with 470 nm light (black arrows) evokes excitatory post-synaptic potentials (EPSPs) in some neurons (left) but not others (right). (C) Higher resolution traces of single optically-evoked response shown in (B).

In order to test for M1 input to individual neurons in S1 we optically stimulated the ChR2-expressing M1 axon terminals while recording from a neuron in S1 (Figure 2.2A). We determined whether the recorded cell received M1 input by stimulating with trains of light pulse (Figure 2.2B). Cells deemed to have an evoked response showed time-locked excitatory postsynaptic potentials (EPSPs) that were typically present following every individual pulse; that is, they displayed little synaptic failure. Excitatory responses were readily observed when averaged across trials (Figure 2.2B&C, left). Occasionally, an inhibitory response was observed as well (see below). In a subset of cells, we computed the latency to EPSP onset on each trial (10-30 trials per cell) and averaged the values for each cell (3.21 ± 0.41 ms; $n=5$ cells). To determine the trial-to-trial ‘jitter’ in the EPSP response onset, for each cell we calculated the coefficient of variation (CV) of the EPSP onset latency across trials. The mean CV was 0.09 ± 0.02 ($n=5$ cells). Because variability between trials was low, we calculated an average EPSP for each cell and performed subsequent analyses on such trial-averaged responses. The trial-averaged EPSP latency for all cells ($n=95$) was 2.99 ± 0.09 ms.

In order to verify that the short-latency responses were monosynaptic we performed a series of experiments where we included 1 μ M tetrodotoxin (TTX) and 1 mM 4-aminopyridine (4-AP) in the bath to block sodium and potassium channels, respectively (Petreanu et al., 2009; Cruikshank et al., 2010). In the presence of TTX only, M1-evoked responses were always abolished (data not shown; $n=11/11$ cells). When we additionally added 4-AP, to enhance indirectly the depolarizing effects of ChR2, evoked responses were present in most cells (data not shown; $n=11/13$). When responses remained in the presence of TTX and 4-AP, for all cells ($n=11/11$) the M1-evoked response latency was significantly longer than under control conditions (control: 3.13 ± 0.04 ms; TTX + 4-AP: 6.46 ± 0.09 ms; $p < 0.005$). Also, almost all

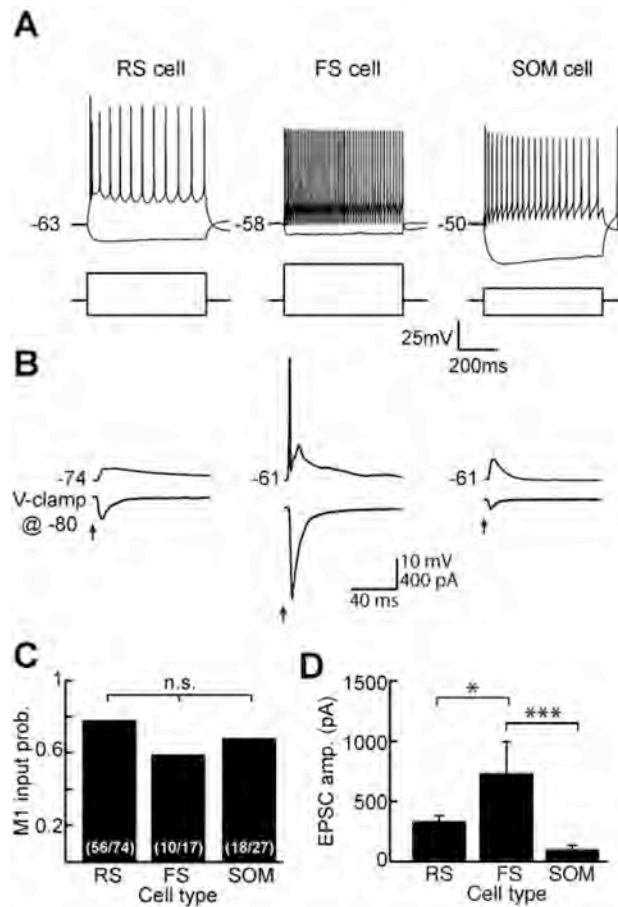


Figure 2.3 M1 contacts three major cell types in S1. (A) Example traces identifying a regular-spiking (RS) pyramidal neuron (*left*), a fast-spiking (FS) inhibitory interneuron (*middle*), and a somatostatin-expressing (SOM) inhibitory interneuron (*right*). Each cell type exhibits characteristic voltage responses (top) to hyperpolarizing and depolarizing current step injections (bottom). (B) Example traces for an RS cell (*left*), FS cell (*middle*), and SOM cell (*right*) demonstrating excitatory responses to optical stimulation of ChR2-M1 fibers. Responses were measured in current-clamp (top) and voltage-clamp (bottom). Holding potential for voltage-clamp is -80 mV for all recordings. Note that the FS cell spiked. (C) Quantification of the probability of receiving an input from M1 by cell type demonstrates equivalence among them ($p = 0.32$). (D) Amplitudes of excitatory currents evoked by ChR2-M1 stimulation are significantly greater in FS than RS ($p < 0.05$) and SOM cells ($p < 0.005$). Amplitudes do not differ between RS and SOM cells ($p > 0.05$). Panels C and D contain cells recorded from layers 2 through 6.

neurons (n=10/11) exhibited reduced M1-evoked response amplitudes in the presence of TTX and 4-AP, although on average this decrease was not significant (control: 6.87 ± 0.33 mV; TTX + 4-AP: 3.80 ± 0.33 mV; $p=0.08$). The cells tested with TTX and 4-AP included all three cell types examined in this study (see below) and the effects of TTX and 4-AP did not depend on cell type. These results are consistent with previous findings (Cruikshank et al., 2010) and illustrate that the light-evoked excitatory responses we observed are directly post-synaptic to ChR2-expressing M1 axon terminals.

Previous work demonstrated that M1 inputs directly contact pyramidal neurons within L2/3 and L5 of somatosensory cortex (Petreanu et al., 2009; Mao et al., 2011; Rocco-Donovan et al., 2011). We wanted to determine here whether M1 also provides synaptic input onto inhibitory interneurons. We therefore recorded from three cell types within S1: regular-spiking pyramidal (RS) neurons, fast-spiking (FS) inhibitory interneurons, and somatostatin-expressing (SOM) inhibitory interneurons (Figure 2.3A). RS and FS neurons were identified using established electrophysiological criteria (see Methods) and SOM interneurons expressed GFP. In addition, we recorded from a population of neurons that could not be readily categorized into one of these three groups yet were too heterogeneous to comprise a meaningful fourth cell type (“non-classified” cells).

We found that in addition to contacting pyramidal neurons with a high probability (n=56/74), M1 also provides input to FS interneurons (n=10/17) and SOM inhibitory interneurons (n=18/27). Connections probabilities were equivalent among the cell types ($p=0.32$; Figure 2.3C). The latency of the evoked responses also did not differ between the three cell types (RS: 2.97 ± 0.02 ms; FS: 2.61 ± 0.07 ms; GIN: 3.08 ± 0.04 ms; $p=0.42$). The “non-classified” cells also received input from M1 (n=9/16; data not shown). The characteristics of the M1 inputs

to the non-classified neurons were not distinct and fell within the range observed for our three identified cell types; therefore we opted to not include these cells in further analyses. Overall, these results suggest that M1 afferents do not discriminate by cell type, but instead provide direct input to most types of cells within S1.

We compared the strength of excitatory responses among cell types. We found that although all three cell types had a similar probability of receiving M1 input, the strength of the inputs differed among them (ANOVA $p=0.0012$; Figure 2.3D). The average amplitude of EPSCs onto FS cells was the largest (730.29 ± 261.1 pA), and this was significantly greater than that of the M1 input onto RS cells (321.83 ± 52.8 pA; $p < 0.05$) or SOM cells (84.43 ± 28.8 pA; $p < 0.005$). This demonstrates that M1 strongly contacts FS cells, which could evoke robust disynaptic inhibition within S1 (see below).

Short-term dynamics of M1 inputs are dependent on post-synaptic cell type

Both thalamocortical and local cortical inputs onto RS, FS, and SOM cells exhibit distinct short-term dynamics that are dependent on the identity of the post-synaptic cell. To determine if M1 inputs show short-term synaptic depression or facilitation, we stimulated the ChR2-expressing M1 terminals with trains of light pulses at 1 Hz and 10 Hz (Figure 2.4A). The dynamics of the M1 input depended on the post-synaptic cell type. With 10 Hz trains, the magnitude of the excitatory post-synaptic currents (EPSCs) displayed short-term depression in the RS cell and FS cell populations and short-term facilitation in the SOM population (Figure 2.4A&B). The time course of short-term depression in the RS and FS cell populations was similar at 1 and 10 Hz (Figure 2.4B).

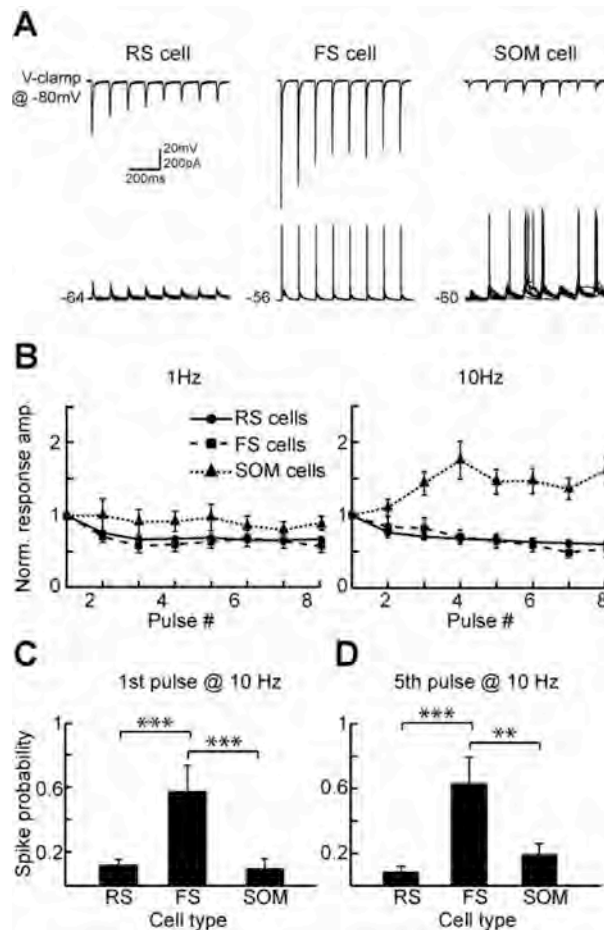


Figure 2.4 Responses to Chr2-M1 optical stimulation exhibit cell type dependent short-term synaptic dynamics. (A) Example traces in voltage-clamp (top) and current-clamp (bottom) demonstrate changes in excitatory response amplitudes and spike probability for RS cells (left), FS cells (middle), and SOM cells (right) across a train of optical stimuli at 10 Hz. Voltage-clamp traces show averaged responses whereas current-clamp traces show multiple single trials overlaid. (B) EPSC responses are normalized to the response amplitude evoked by the first pulse for RS cells (solid line; circles), FS cells (dashed line; squares), and SOM cells (dotted line; triangles) at both 1 Hz (left) and 10 Hz (right). (C) For the first pulse in a 10 Hz train, stimulation of Chr2-expressing M1 terminals resulted in significantly greater probability of spiking in FS than RS cells ($p < 0.005$) and SOM cells ($p < 0.005$). (D) Spiking probability was also greater in FS than RS cells ($p < 0.005$) and SOM cells ($p < 0.01$) for the fifth pulse in a 10 Hz stimulation train. Data in (C) and (D) consist of all cells receiving M1 input.

We also recorded in current-clamp and examined the change in the probability of eliciting spikes across the stimulus train (Figure 2.4A, C&D). Out of the cells in each cell type that spiked at least once (16/56 RS cells; 6/10 FS cells; 10/18 SOM cells), the probability of eliciting a spike across the train tended to increase in SOM cells and decrease in the RS cells, although the change was variable in both populations (data not shown). The incidence of spiking is consistent with SOM cells displaying short-term facilitation and RS cells exhibiting short-term depression (Figure 2.4B). In FS cells, although EPSC responses displayed synaptic depression, we observed spiking in response to each pulse across the 10 Hz train. This is likely due to the large amplitude EPSCs evoked in FS cells (Figure 2.3D) under these conditions.

When we compared all cells receiving M1 input for each cell type under equivalent recording conditions (i.e. same light intensity for stimulation), FS cells (n=10) were significantly more likely to spike to both the first pulse (0.64 ± 0.16 ; Figure 2.4C) and fifth pulse (0.63 ± 0.16 ; Figure 2.4D) in a 10 Hz train than RS (1st pulse: 0.12 ± 0.04 , 5th pulse: 0.09 ± 0.03 ; $p < 0.005$) or SOM (1st pulse: 0.10 ± 0.06 , $p < 0.01$; 5th pulse: 0.19 ± 0.07 , $p < 0.005$) cells. This is consistent with our previous finding that FS cells receive stronger excitatory drive from M1 than RS or SOM cells and likely produce feedforward inhibition onto S1 neurons.

Feedforward inhibition recruited by M1 stimulation

We observed a high probability of spiking in the FS cell population, indicating that M1 activation could recruit widespread disynaptic inhibition within S1. To test this possibility, we recorded in voltage-clamp and held the cells at a potential of -20 mV to determine the presence or absence of an inhibitory post-synaptic current (IPSC; Figure 2.5). We observed IPSCs in RS

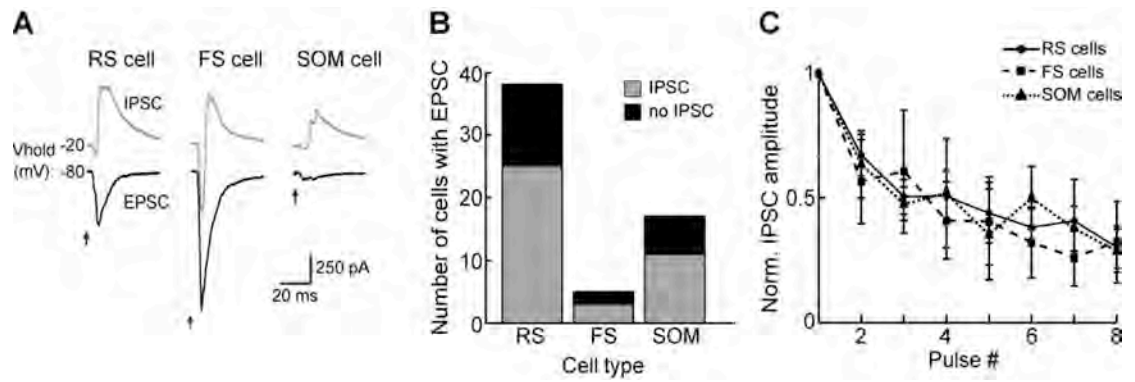


Figure 2.5 Optical stimulation of Chr2-M1 terminals evokes widespread disinaptic inhibition in RS cells, FS cells, and SOM cells within S1. (A) Examples of excitatory and inhibitory post-synaptic currents (EPSCs and IPSCs) in RS cells (left), FS cells (middle), and SOM cells (right) in response to Chr2-M1 stimulation. Cells were held at -80 mV for EPSC recordings and -20 mV for IPSC recordings. (B) Summary of number of neurons with both E&IPSC responses by cell type. (C) Normalized IPSC amplitudes for RS cells (solid line; circles), FS cells (dashed line; squares), and SOM cells (dotted line; triangle) for a 10 Hz optical pulse train.

cells, FS cells, and SOM cells that appeared delayed relative to the initial EPSC (Figure 2.5A). We found that for each cell type a high proportion of the cells that received excitatory input from M1 also displayed a disynaptic inhibitory response (66% of RS cells, 60% of FS cells, 65% of SOM cells; Figure 2.5B). The peak of the IPSC followed the peak of the EPSC in nearly all cells ($n=36/38$; average delay was 4.89 ± 0.57 ms). The IPSC-EPSC peak delay was slightly less in FS cells (3.76 ± 1.07 ms) than RS (4.99 ± 0.77 ms) or SOM cells (4.97 ± 1.02 ms), however there was no significant difference among cell types ($p=0.85$). In contrast to the short-term dynamics of the M1 excitatory inputs, the disynaptic inhibitory responses showed short-term depression in all three cell types (Figure 2.5C). This suggests that M1 activation generates extensive inhibition within S1 that depresses at frequencies of 10 Hz and greater. That FS cells are the most likely to be driven suprathreshold by M1 stimulation strongly suggests that this inhibition is mediated by FS interneurons.

Laminar dependence of M1 input

The pattern of ChR2-M1 axons was distinctly non-uniform across S1 cortical layers (see above; Figure 2.1). We therefore examined in a subset of cells whether particular layers within S1 receive more M1 input than others and whether this depends on cell type (Figure 2.6A). The probability of a cell receiving an M1 input was lowest in L4 (~20%), and this was the case for all three cell types (RS: $p=0.0008$, FS: $p=0.016$, SOM: $p=0.037$; Figure 2.6B). In each of the other layers, input probability was ~80% for each cell type. We similarly examined response amplitudes (EPSPs). For all S1 neurons we recorded, there was a significant difference in response magnitude across layers (ANOVA $p < 0.005$; data not shown). When we separated the data by cell type, RS amplitudes differed significantly by laminar location (ANOVA $p=0.0011$).

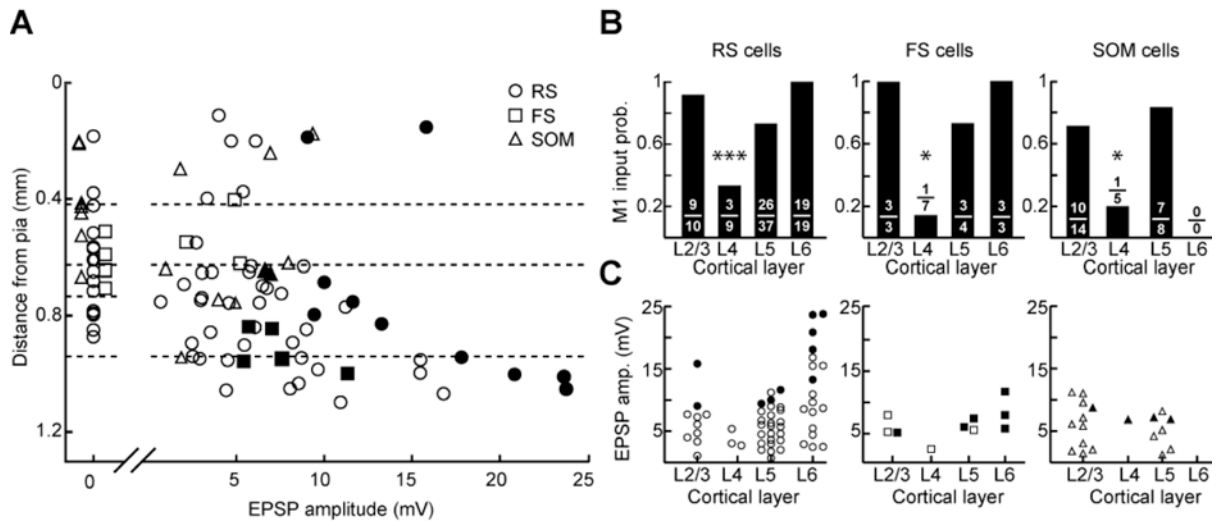


Figure 2.6 M1 inputs to excitatory and inhibitory neurons in S1 exhibit similar laminar distributions. (A) Amplitude of M1 input as a function of depth from pia. Each point represents EPSP amplitude in response to ChR2-M1 stimulation for individual cells (circles: RS cells; squares: FS cells; triangles: SOM cells). Dashed lines indicate laminar boundaries and were placed as follows (in μm from pia): L2/3-L4: 419; L4-L5: 626; L5-L6: 940. Data plotted are a subset of data shown in (B) and (C). Filled circles represent cells that spiked in response to ChR2-M1 on at least 2 trials; open circles are cells that did not spike. (B) Probability of receiving an M1 input by layer for RS cells (left), FS cells (middle), and SOM cells (right). Numbers represent the number of cells that received input out of total number of cells recorded for that layer and cell type. (C) EPSP amplitude of evoked response to ChR2-M1 optical stimulation as a function of layer for RS cells (left; circles), FS cells (middle; squares), and SOM cells (right; triangles). Conventions as in (A).

Laminar dependency was at best at trend level for FS (ANOVA $p=0.18$), and SOM cells did not display laminar differences (ANOVA $p=0.82$). In both RS and FS cells, excitatory responses were largest in L6 as was the proportion of cells that fired action potentials. In RS cells, L6 response amplitudes were on average significantly larger than in the other layers ($p < 0.05$). We were unable to determine whether SOM interneurons also receive their strongest M1 inputs in L6 because there are no GFP+ cells in L6 of GIN mice; SOM cells do exist there, however (Markram et al., 2004; Ma et al., 2006; Lee et al., 2010).

We found unexpectedly that M1 inputs were especially large in a subset of L6 cells. To verify that these results were not an artifact of between-animal variability in ChR2 expression or activation (see Methods), we recorded sequentially from RS neurons in L5 and L6 within the same slice (Figure 2.7). Pair-wise comparison of L5 and L6 RS cells showed that L6 cells indeed receive, on average, significantly larger M1 inputs ($p=0.02$; Figure 2.7B, left) though the difference was not necessarily observed in every pair. When we normalized the paired data to the amplitude of the L5 cell, it was clear that relative to nearby L5 neurons a subset of neurons in L6 receive especially strong M1 input (mean normalized L6: 2.47 ± 0.57 , $p=0.02$; Figure 2.7B, right). Interestingly, it appears that the L6 cells having the largest responses are concentrated in upper L6, near the L5/L6 border (Figure 2.6A). The cells receiving large M1 inputs exhibited regular-spiking characteristics, as did all L6 pyramidal neurons; however they did tend to show an initial spike ‘doublet’ in response to current step injection whereas L6 pyramidal neurons receiving less M1 input did not. These results suggest they could represent a specific class of projection neurons (Otsuka and Kawaguchi, 2011) that remain to be identified in future studies.

2.5 DISCUSSION

Here we took advantage of optogenetic tools to investigate motor cortex to somatosensory cortex synaptic circuitry. By injecting an AAV vector carrying the ChR2 gene into M1 we were able specifically to investigate properties of anterograde connections from M1 to S1. Provided that ChR2 is transported only anterogradely, our experimental approach avoided confounds associated with inadvertent synaptic activation by local recurrent axons in S1. We verified that labeling was indeed virtually entirely anterograde in nature (see Methods).

Our experiments yield four important findings. First, M1 inputs engage three main types of S1 neurons, including two classes of inhibitory interneurons as well as excitatory pyramidal neurons. As a consequence, M1 activation produces both direct excitation as well as substantial disynaptic inhibition within S1. Second, motor cortex inputs to pyramidal neurons, FS and SOM inhibitory interneurons possess cell-type specific synaptic dynamics. The dynamics are similar to those of other extrinsic and intrinsic inputs, including thalamocortical and local corticocortical connections. Third, M1 input to excitatory and inhibitory neurons display similar laminar distributions. Regardless of cell type, neurons in L4 were the least likely to receive input from motor cortex, whereas all three cell types in other layers showed a similar, high probability of M1 input. Fourth, M1 labeling is distributed into vertically-oriented columns that are complementary to those described for thalamocortical inputs from the ventral posteromedial (VPM) thalamic nucleus, the major subcortical station in the whisker-to-barrel pathway. Taken together, our findings suggest that M1 exerts on S1 modulatory excitatory and inhibitory effects that may differentially influence corticocortical vs. thalamocortical processing streams.

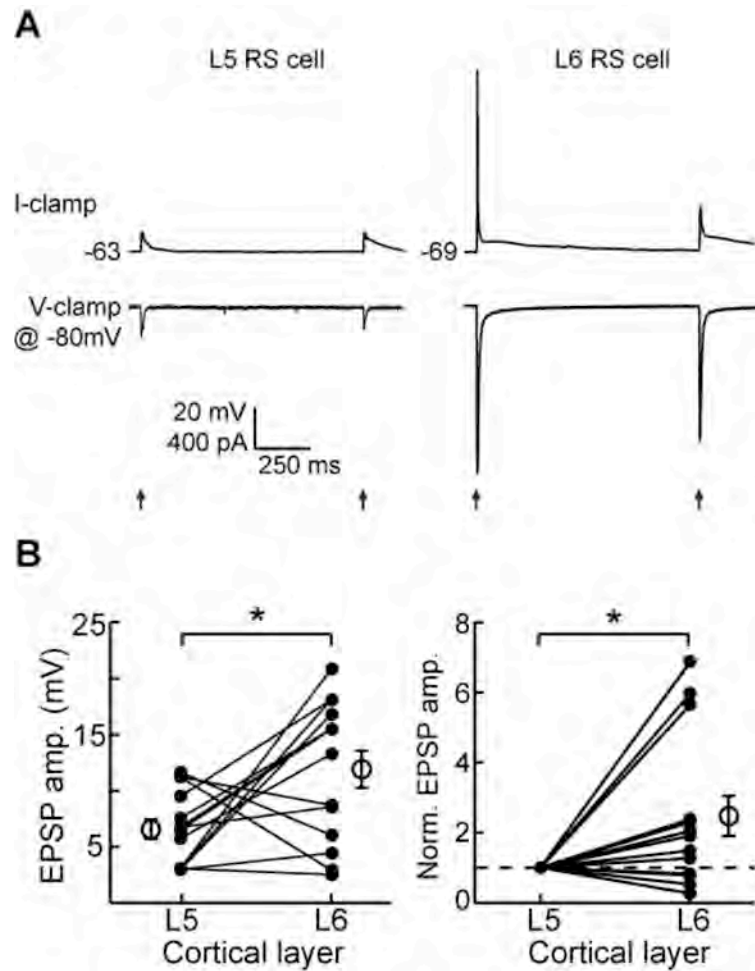


Figure 2.7 Optical stimulation of ChR2-expressing M1 terminals within S1 evokes stronger responses in L6 cells than in L5 cells. (A) Example traces in current clamp (top) and voltage clamp (bottom) of a L5 RS neuron (left) and a L6 RS neuron (right) recorded from the same slice. (B) Average M1 response amplitudes for paired L5 and L6 RS cells within the same slice (left; $p < 0.05$, paired t-test). Same data normalized to the amplitude of the L5 cell (right; $p < 0.05$, t-test). Filled circles represent individual cells and open circles and bars represent the mean and SEM for each population.

Selective stimulation of M1 axons with channelrhodopsin

Use of optogenetic techniques allowed for sole activation within S1 of axons originating from primary motor cortex. Such specificity has been difficult to achieve with traditional electrical stimulation techniques, as unintentional antidromic activation of fibers in reciprocally connected brain regions, such as S1 and M1, is difficult to avoid. In addition, we were able to activate ChR2-expressing fibers directly within S1 as opposed to the neuronal somata in M1. This allows greater confidence for detecting the presence of synaptic connections, as the entire axon tract need not be preserved within the slice. Under these conditions, the latency measure likely reflects primarily the synaptic delay of neurotransmitter release from the presynaptic ChR2-expressing terminal to the post-synaptic neuron.

Wide-field optical stimulation, as we used in this study, likely produces synchronous or near-synchronous activation of all or most of the ChR2-expressing M1 terminals within the activated region, a condition that could correspond to the upper-bound of S1 engagement by M1. While our stimulation protocol may not be analogous to physiological M1 activity, it enabled us to identify connections that may not have been apparent with weaker stimuli. Importantly, our approach provided the opportunity to identify similarities, such as feedforward inhibition, and differences, i.e., direct engagement of SOM cells, in afferent cortical circuitry, notably that associated with thalamic input (see below).

Further investigation into the strength of unitary connections, as opposed to population connections, will provide additional insight into whether the large inputs to some S1 neurons result from stronger synaptic connections and/or a convergence of more synapses onto single post-synaptic neurons. Traditional *in vitro* experimental paradigms, such as minimal stimulation

techniques, could be adapted for use with optical stimulation to address these questions in future studies. Nevertheless, our findings are consistent with other results demonstrating that FS cells receive large thalamocortical and corticocortical inputs. Previous studies have demonstrated for these inputs that FS cells receive larger unitary inputs than RS cells (Cruikshank et al., 2007) or SOM cells (Beierlein et al., 2003); it seems likely that the same could be true for M1 inputs onto FS interneurons.

Motor cortex input to inhibitory and excitatory neurons in S1

At a population level, motor cortex neurons contact three major types of S1 neurons with high probability. Indeed, up to 80% of recorded S1 cells displayed short-latency excitatory responses. The strongest M1 inputs are to FS inhibitory neurons. This was evident in both the size of the excitatory currents and the likelihood that these inputs evoked FS spikes. Our finding that FS interneurons overall receive the strongest M1 input could be a reflection of the greater recruitment of FS interneurons located in deeper layers (see below). That FS cells receive large excitatory input is reminiscent of the synaptic properties of thalamocortical inputs to FS cells in layer 4 barrels (Porter et al., 2001; Beierlein et al., 2003; Cruikshank et al., 2007). SOM inhibitory interneurons also have a high probability of receiving M1 input, but the amplitude of the responses and the probability of spiking were substantially smaller. SOM inhibitory interneurons have high input resistances and depolarized resting potentials (Fanselow et al., 2008; Kinnischtzke et al., 2012). However, despite their greater intrinsic excitability relative to FS interneurons the probability of eliciting action potentials with M1-ChR2 activation was much lower at the stimulation frequency used here. Together, these M1-S1 findings parallel those showing strong input from local S1 excitatory neurons onto FS interneurons and weak input onto

SOM interneurons (Markram et al., 1998; Xiang et al., 2002; Beierlein et al., 2003; Thomson and Lamy, 2007). Thus, the distinctive synaptic profiles of FS and SOM interneurons extend to at least one long-range, inter-areal corticocortical system.

Given the high convergence of FS cells onto local excitatory neurons in S1 (Thomson and Lamy, 2007; Oswald et al., 2009; Packer and Yuste, 2011), strong activation of FS cells by motor cortex is likely to result in widespread disynaptic inhibition within S1. We determined the presence or absence of disynaptic IPSCs following M1 stimulation, and indeed found that when an S1 neuron of any cell type received an excitatory M1 input it was usually followed by an IPSC. Based on our findings with FS and SOM inhibitory cells, we hypothesize that this disynaptic inhibition originates from FS cells. We also recorded from “non-classified” neurons. This group likely contains inhibitory neurons that were neither GFP+ (identifying them as SOM-expressing) nor conclusively fast-spiking interneurons, such as VIP-expressing, CCK-expressing, or neurogliaform cells (Markram et al., 2004; Lee et al., 2010). Because the “non-classified” cells also received input from M1 (data not shown), it is possible that these additional inhibitory interneuron classes may also be involved in mediating the IPSCs observed here. This could also vary by depth, as the relative abundance of these inhibitory interneuron groups varies across S1 layers (Lee et al., 2010). However, given the high spiking probability that we observed in FS cells, M1-evoked inhibition in our preparation most likely originated from the FS inhibitory neurons. Direct, efficacious thalamocortical (TC) inputs to FS cells also provide strong, fast disynaptic inhibition for processing sensory signals (Cruikshank et al., 2007). Thus, FS interneurons generate fast, reliable inhibition in response to thalamic, local corticocortical, and M1 signals.

Pyramidal neurons in S1 show considerable diversity in terms of their morphology, intrinsic biophysical properties, projection targets, sources of excitatory input and response properties (e.g. Swadlow, 1989; Elhanany and White, 1990; Agmon and Connors, 1992; Hattox and Nelson, 2007). Given this diversity, we were somewhat surprised by the lack of discernible groups of pyramidal neurons in terms of their M1 inputs. With the exception of some L6 cells (see below), responses tended to be subthreshold (usually <10 mV). These inputs appear to be smaller than those of the reciprocal S1 to M1 projection (Mao et al., 2011; Rocco-Donovan et al., 2011). These findings suggest that M1 provides weak excitation to L2/3 and L5 pyramidal neurons in S1 regardless of their projection target, as well as to many L6 pyramidal neurons. This conclusion is consistent with recent findings that L2/3 and L5A M1-projecting pyramidal neurons receive similar magnitudes of M1 input as nearby pyramidal neurons projecting to other targets (Mao et al., 2011). M1 may therefore provide a general increase in excitation to the majority of pyramidal neurons in S1, with a subset of L6 pyramidal cells being the exception.

Synaptic dynamics of M1 inputs onto neurons in S1

A striking feature of SOM interneurons is how the efficacy of their excitatory synaptic inputs changes with repeated stimulation. Local corticocortical synapses to SOM cells facilitate, whereas excitatory inputs to RS and FS cells depress. We observed qualitatively similar effects here. In response to periodic M1-ChR2 stimulation, RS and FS cells exhibited short-term depression similar to that observed for thalamocortical (Beierlein and Connors, 2002; Chung et al., 2002; Cruikshank et al., 2007) and local intra-cortical synapses (Markram et al., 1998; Reyes et al., 1998; Oswald and Reyes, 2008, 2011). These findings indicate that the distinct synaptic dynamics that apply to FS versus SOM inhibitory interneurons extend to long-range cortical

inputs, such as from primary motor cortex. Because SOM cells exhibit short-term facilitation it has been suggested they provide a delayed source of inhibition in the presence of sustained high frequency inputs (Pouille and Scanziani, 2004; Kapfer et al., 2007; Silberberg and Markram, 2007). We did not measure responses at frequencies above 10 Hz, as we observed at frequencies ≥ 20 Hz a larger probability of response failure as well as a summation of post-synaptic responses. This could be a result of using ChR2 as opposed to electrical stimulation, as ChR2 has relatively slow recovery from inactivation and can continue to conduct for a few milliseconds following removal of the light stimulus (Nagel et al., 2003; Cruikshank et al., 2010; Gunaydin et al., 2010). M1 inputs to SOM cells may continue to facilitate at higher frequencies, as occurs at local RS to SOM synapses (Fanselow et al., 2008), which could lead to greater recruitment of SOM interneurons.

Synaptic depression in RS and FS cells at 10 Hz was modest, particularly considering that channelrhodopsin tends to slightly exaggerate the degree of synaptic depression compared to electrical stimulation (Cruikshank et al., 2010). Nevertheless, our data are consistent with the idea that M1-S1 synapses depress less than thalamocortical synapses. Using a similar optogenetics approach, Cruikshank et al reported greater synaptic depression at TC-FS and TC-RS synapses than we find at M1-FS and M1-RS synapses. M1 inputs may therefore be more similar to local corticocortical synapses, which are generally thought to exhibit less synaptic depression than thalamocortical synapses (Gil et al., 1997; Beierlein et al., 2003). Together, findings raise the interesting possibility that the intrinsic nature of synaptic depression reflects the spiking statistics of the pre-synaptic neurons. That is, populations associated with high firing rates may have more depressing synapses than those associated with lower firing rates. In this regard, *in vivo* firing of VPM neurons is likely higher than that of cells in S1. Correspondingly,

for equivalent stimulation frequencies TC synapses display more depression than S1 synapses. In addition, overall *in vivo* firing frequencies of M1 neurons projecting to S1 may be lower than those of VPM cells, and this may be reflected in the tendency for M1-S1 synapses to exhibit less short-term synaptic depression.

Laminar and columnar organization of M1 input to S1

Our anatomical data demonstrate that L4 is largely devoid of M1 axons, particularly within barrel centers. In accordance with this, neurons in L4 were the least likely to receive input from M1 (Figure 2.6). This was true for all three cell types: RS excitatory cells, FS inhibitory neurons and SOM inhibitory interneurons. L4 is the primary input layer for thalamocortical axon terminals to S1 from VPM (Killackey, 1973). Because we observed relatively few M1 contacts to L4 neurons, we conclude that M1 is unlikely to directly modulate VPM inputs within L4. Furthermore, S1 neurons that project back to M1 are similarly scarce in L4 (Alloway et al., 2004). Therefore, during active whisking M1 may modulate S1 in a layer-specific manner (Krupa et al., 2004), with signal processing in L4 occurring largely independently of direct M1 influences.

Motor cortex axons terminate in a distinct pattern within S1, wherein the axon from a single M1 neuron terminates extensively within L5 and L6 before traveling vertically and ramifying largely within L1 (Veinante and Deschenes, 2003). We observed the same pattern of ChR2 labeling (Figures 2.1&2.2). M1 projections were vertically concentrated in columns interposed between barrel-related columns, particularly in L4. In rats, M1-projecting pyramidal neurons are also concentrated in vertical columns within inter-barrel septa (Alloway et al., 2004). VPM-related circuits in L4 and L6 are more barrel-focused than inputs from the thalamic

posteromedial (POm) nucleus, which target inter-barrel septa (Chmielowska et al., 1989; Wimmer et al., 2010). Present findings thus provide further evidence for a vertical organization within the whisker area of S1 in which barrel- and septal-related columns represent different information streams and/or different modes of sensorimotor integration. As suggested by others (Alloway, 2008), septal-related columns may be more tightly coupled to the motor system.

Pyramidal cells in L6 exhibited more heterogeneity in their M1 input amplitudes than pyramidal neurons in any of the other layers (Figure 2.6). On average, L6 pyramidal neurons had significantly larger M1 inputs. This was due to a subset of cells that received exceptionally strong inputs; unlike other pyramidal cells these often spiked. Pyramidal neurons in L6 are heterogeneous in their projections (Zhang and Deschenes, 1997). Perhaps M1 strongly contacts a particular class of L6 neurons, such as those projecting to a specific target (e.g. VPM, POm, M1, etc.).

M1 input magnitudes to pyramidal neurons are progressively larger in deeper layers, particularly at the border of L5 and L6 (Figure 2.6A). Interestingly, the inputs to FS cells scaled similarly. FS cells in L5 and L6 received strong M1 inputs, and most of them reliably fired action potentials in response to ChR2 stimulation. Although SOM cells constitute a major portion of inhibitory interneurons within L6 (Markram et al., 2004; Lee et al., 2010), in our mouse line SOM cells in L6 are not labeled with GFP (Ma et al., 2006) and we were therefore unable to identify and record from SOM-expressing interneurons in L6. However, our findings with the FS cells provide evidence that M1 recruits strong inhibition within the deeper layers that parallels strong, direct excitatory input from M1.

Functional Significance

Our findings demonstrate that M1 broadly engages S1 circuits and, further, that M1 inputs onto pyramidal neurons are widespread but usually subthreshold. Thalamocortical recipient zones seem to be the exception, with L4 barrel centers receiving few inputs and upper layer 6 receiving especially strong ones. Our findings thus suggest that M1 targets multiple pyramidal cell populations that participate in a variety of local circuits and that collectively project to a broad range of cortical and subcortical targets. M1 neurons also directly contact at least two major types of inhibitory neurons and possibly other types.

M1 neurons abruptly increase their firing prior to the onset of whisking and many continue to fire tonically throughout the whisking bout (Carvell et al., 1996; Friedman et al., 2006; Friedman et al., 2012). Present results show that M1 activates fast feedforward inhibition, likely via FS inhibitory interneurons, in a manner similar to that of thalamocortical inputs to layer 4 of barrel cortex. There, feedforward inhibition creates a ‘window of opportunity’ for L4 neurons to spike, producing a robust and brief signature of stimulus onset (Gabernet et al., 2005). Feedforward inhibition may serve a similar role in M1-S1 interactions. The pre-whisking burst of M1 activity could transiently engage S1 neurons, providing a strong signal of whisking onset; this could set S1 circuits in a state to be further modified by the continued tonic firing of some M1 neurons.

On longer timescales, studies have demonstrated that over the course of a whisking bout the responsiveness of cells in S1 is diminished relative to non-whisking states (Fanselow and Nicolelis, 1999; Hentschke et al., 2006; Ferezou et al., 2007). One possibility is that sustained increases in motor cortex firing during whisking suppress activity in S1. We have demonstrated that such a neural substrate exists in the direct projection from M1 onto S1 inhibitory interneurons. However, given that M1 inputs broadly engage populations of both inhibitory and

pyramidal neurons, it seems unlikely that M1 exerts a predominantly inhibitory or excitatory effect on S1. Furthermore, the activity of S1 FS and SOM inhibitory interneurons also decreases during whisking (Gentet et al., 2010; Gentet et al., 2012). In the whisker-barrel system, as well as in other somatosensory systems, motor-related gating occurs at subcortical levels (Furuta et al., 2008), resulting in reduced stimulus-evoked thalamic firing (Lee et al., 2008). Therefore, the reduced activity of S1 excitatory and inhibitory neurons during whisking may be a reflection of decreased thalamic input that is not fully compensated for by corticocortical input from M1.

Elevations in motor cortex firing that precede and accompany whisking may produce a barrage of excitatory and inhibitory synaptic activity within S1 that contributes to state-dependent changes observed in cortical neurons during active touch. Transformations in input-output functions of individual pyramidal neurons as well as the correlated activity between neurons can be regulated dynamically by the timing and rate of excitatory and inhibitory inputs (Chance et al., 2002; Prescott and De Koninck, 2003; de la Rocha et al., 2007; Renart et al., 2010). Moreover, M1 inputs are located more distally along S1 apical dendrites than VPM inputs (Petreanu et al., 2009). Changes in the balance of proximal and distal synaptic inputs can also affect neuronal responsiveness (Larkum et al., 2004; Oviedo and Reyes, 2005). Whisking-associated reductions in sensory-evoked thalamic activity, accompanied by increases in motor cortex activity, could therefore modify or even sharpen stimulus-encoding properties of S1 neurons. M1-S1 inputs may thus contribute to sensorimotor integration by modulating S1 circuits through broad recruitment of both excitatory and inhibitory cell populations.

3.0 DISTINCT INTRINSIC PROPERTIES AND M1 INPUT TO INFRAGRANULAR S1 PYRAMIDAL NEURONS PROJECTING TO DIFFERENT TARGETS

3.1 ABSTRACT

The pathway from primary motor cortex (M1) to primary somatosensory cortex (S1) is thought to influence activity in S1 by conveying a general modulatory signal and/or a copy of the motor command. The nature of the circuits in S1 that receive such signals are still unknown; functionally, the impact of M1 inputs may depend on cell-type specific microcircuits within S1. Recently, we discovered that a subset of pyramidal neurons in the infragranular layers of S1 receive especially strong input from M1 (Kinnischtzke et al., 2013), suggesting that M1 may differentially contact classes of pyramidal neurons, such as those projecting to different sensorimotor centers at cortical and subcortical levels. To test this hypothesis, we combined optogenetic techniques for specifically labeling M1 synaptic inputs to S1 and retrograde tracing identifying specific populations of projection neurons in infragranular layers of S1. In agreement with previous findings, we observed that different projection neurons have distinguishing intrinsic electrophysiological properties. Here we report that the magnitude of M1 inputs to an S1 neuron is highly dependent on its projection target. These results indicate that M1 may engage S1 by circuit-specific mechanisms related to how S1 influences subcortical centers from which it receives afferent input.

3.2 INTRODUCTION

During active exploration, sensory and motor systems function in concert so that sensory feedback can inform ongoing and future movements. Although different regions within the central nervous system process sensory and motor information, the structures are highly interconnected to form large-scale sensorimotor integration systems (Kleinfeld et al., 2006; Petersen, 2007). At the level of the telencephalon, primary somatosensory (S1) and primary motor (M1) cortices are reciprocally connected (White and DeAmicis, 1977; Porter and White, 1983; Miyashita et al., 1994; Cauller et al., 1998). Both anatomical and physiology studies demonstrate that S1 sends a strong projection to superficial layers in M1, in particular to neurons that project back to S1 (Aronoff et al., 2010; Mao et al., 2011). Conversely, M1 densely innervates and provides the strongest input to the infragranular layers in S1 (Zhang and Deschenes, 1998; Veinante and Deschenes, 2003; Kinnischtzke et al., 2013).

Layers 5 and 6 of somatosensory cortex contain a diversity of pyramidal neuron populations that project to a number of cortical and subcortical sites. Our previous study (Kinnischtzke et al., 2013) demonstrated that pyramidal neurons in S1 are broadly contacted by M1, consistent with the view that M1 sends a general ‘modulatory’ signal to S1. On the other hand, we also observed significant variability in the amount of M1 input, with a subset of neurons in the top of L6 receiving especially large inputs. M1 may therefore differentially engage specific circuits within S1, possibly those projecting to particular cortical or subcortical targets.

Here, we investigate synaptic input from M1 onto several classes of S1 pyramidal neurons: neurons projecting back to M1 (M1-projecting), corticothalamic neurons projecting to either the posteromedial thalamus (PoM-projecting) or ventral posterior medial nucleus (VPM-

projecting), and corticotrigeminal neurons (Sp5-projecting). Our results demonstrate that the projection target of the neuron is strongly related to both its intrinsic electrophysiological properties and the strength and effectiveness of the M1 inputs it receives.

3.3 METHODS

All experiments were carried out in compliance with the University of Pittsburgh School of Medicine animal use policies and were approved by the University of Pittsburgh Institutional Animal Care and Use Committee.

Channelrhodopsin and retrograde labeling procedures

Viral constructs carrying channelrhodopsin-2 (ChR2) and mCherry genes were injected into primary motor cortex (M1) as previously described (Kinnischtzke et al., 2013). Briefly, mice (aged 10-16 postnatal days of age) were anesthetized using isoflurane (1–2%/oxygen) and a craniotomy was performed over M1 (1.0 mm anterior to and 0.8 mm lateral from bregma). The virus was pressure-injected at depths of 0.9 and 0.4 mm (corresponding to deep and superficial layers in M1) within a single penetration. A volume of 0.1-0.2 μ l was injected at each depth. We used the adeno-associated virus construct AAV2/5.CamKII α .hChR2(H134R)-mCherry.WPRE.SV40 (University of Pennsylvania Vector Core; permission from Dr. Karl Deisseroth) for all experiments, as this construct results in strong anterograde expression but virtually no retrograde transport (Kinnischtzke et al., 2013).

During the same surgery, mice were also injected with red and green fluorescent

retrograde tracers (RetroBeads, Lumafluor Inc.). All mice had green retrobeads injected in M1; the beads were typically co-injected simultaneously with the ChR2 virus (in approximately 1:4 ratio with the virus). Mice were then injected with red fluorescent retrobeads in one of the following locations: ventral posterior medial (VPM) thalamus, posteromedial (PoM) thalamus, or the spinal trigeminal nucleus (Sp5). For injections in the thalamus, a small craniotomy was performed dorsal to VPM (1.4 mm posterior and 1.7 mm lateral to bregma) or PoM (1.6 mm posterior and 1.4 mm lateral to bregma). Beads were pressure-injected (volume of 0.1-0.2 μ l) using a picospritzer (depth of 3.20 mm for VPM and 3.10 mm for PoM). The pipette was withdrawn slowly 5-10 minutes after injection to limit inadvertent leakage in the cortex. For Sp5 injections, the brainstem was exposed directly posterior to the cerebellum and the pipette was advanced at a 40° angle. Retrobeads were pressure injected at depths of 0.8 and 0.6 mm (volume of 0.1 μ l at each depth). Experiments were conducted using the ‘GIN’ transgenic mouse line (Oliva et al., 2000), in order to be consistent with our previous study.

Slice preparation

Electrophysiological experiments were performed 4-6 weeks following the injection surgery to allow for transport and full expression of the virus. Mice were between postnatal days 44-74 at the time of experiments. Mice were anesthetized with isoflurane, and the brain was removed and placed in ice-cold artificial cerebrospinal fluid (ACSF) containing (in mM): 126 NaCl, 3 KCl, 1.25 NaH₂PO₄, 2 MgSO₄, 26 NaHCO₃, 10 dextrose, and 2 CaCl₂, saturated with 95% O₂–5% CO₂. The tissue was sliced at 400 μ m in the coronal plane using a vibratome. Slices were incubated at 32 °C for 30–45 min and then maintained at room temperature until used for recording. Slices containing S1 barrel cortex were identified by the presence of layer IV

barrels and a patchy barrel-related pattern of ChR2-mCherry fluorescence (Figure 3.4A). We recorded primarily from the larger, more medially situated barrels (rows D–E), as this is where the ChR2-mCherry labeling was typically strongest. We recorded from 1 to 3 adjacent barrel-related columns per slice.

Recording procedures

All recorded cells were positive for either red or green retrobeads, which allowed for unambiguous classification of the projection target. Whole-cell recordings were performed using glass micropipettes (4–10 M Ω) filled with internal solution containing (in mM): 135 K-gluconate, 4 KCl, 2 NaCl, 10 HEPES, 0.2 EGTA, 4 ATP-Mg, 0.3 GTP-Tris, and 14 phosphocreatine-Tris (pH 7.25, 280–290 mOsm). Biocytin (0.5%) was added to the internal solution in the majority of experiments. Membrane potentials reported here were not corrected for liquid junction potential. Recordings were conducted at 32 °C. When patching, cell-attached seal resistances were ≥ 1 G Ω , and series resistance after achieving whole-cell configuration was 5–20 M Ω . After establishing whole-cell configuration, a series of current steps was presented in current-clamp for use in analysis of electrophysiological characteristics. Current steps were 600 ms in duration and presented in 20 pA steps, ranging from –100 to 300 pA. Steps were presented 5 s apart. In voltage-clamp experiments, series resistances ranged from 10 to 40 M Ω and were compensated for up to 70%. For excitatory postsynaptic current (EPSC) measurements, all cells were held at a potential of –80 mV. Data were collected using a Multiclamp 700B amplifier and pClamp10 software (Molecular Devices) at a sampling rate of 20 kHz.

Laminar definitions

Laminar boundaries were measured using Nissl stained coronal sections in S1 (Figure 3.1B). Measurements of laminar position are reported as normalized to the top of L5A (value of 0.0) and the bottom of L6 (value of 1.0) to account for slight variability in cortical depth due, for example, to section plane. The actual, non-normalized, laminar boundary measurements were similar to previously reported values (Hooks et al., 2011). Border boundaries were based on changes in cell densities indicated by the Nissl stain. The L4/L5A boundary showed a clear border, indicated by an abrupt loss of barrel structure and a decrease in the number of cells. The bottom of L5A was estimated by changes in the density of cells and was typically $\sim 100 \mu\text{m}$ below the L4/L5A border. The boundary between L5B and L6 was visible as a notable increase in cell density near the top of L6; it was typically located approximately halfway between the top of L5A and the subcortical white matter.

Optical stimulation procedures

All experimental procedures were carried out as previously described (Kinnischtzke et al., 2013). Full-field blue light (470 nm; OptoLED, Cairn Research) was delivered through a 40x objective; light intensity at the surface of the slice was $\sim 20 \text{ mW/mm}^2$. A single pulse (duration 1.0 ms) was delivered per trial and individual trials were separated by 8 seconds.

Data analysis

Data analyses were performed using custom-written MATLAB codes (The MathWorks, Natick, MA, USA; A. Kinnischtzke). Intrinsic properties were derived from a series of DC current steps that were presented to the cell (see above). Input resistance was calculated as the slope of the voltage-current relationship for subthreshold voltage deflections that were within

± 30 mV of the resting membrane potential. Using hyperpolarizing current steps, sag current magnitude (mV) was measured as the difference in the minimum voltage during the initial 100 ms and the mean voltage during the last 50 ms in the step. Because sag currents exhibit a voltage-dependence, the first current step where the minimum voltage value was greater than -90 mV was used.

Action potential (AP) features were calculated from the first action potential in the first current step that elicited at least two APs at 10 Hz or greater. AP threshold was measured as the maximum of the second derivative in the voltage 10 ms preceding the AP peak. AP half-width was calculated as the width halfway between the AP threshold and the AP peak. To measure afterhyperpolarization (AHP) depth, the minimum voltage (V_{\min}) within the 5 ms following the AP threshold was subtracted from the AP threshold (see Figure 3.3D). To determine the presence or absence of a depolarizing after-potential (DAP), the local maximum (V_{\max}) was found between V_{\min} and the voltage during the following 10 ms. The voltage 17 ms following V_{\min} was also calculated (V_{end}). If $V_{\max} > V_{\text{end}}$, the DAP magnitude was calculated as the difference between V_{\max} and V_{\min} (see Figure 3.3D). If $V_{\max} < V_{\text{end}}$, the DAP magnitude was 0.

Rheobase current was the first current step amplitude that elicited at least one action potential. All spike train properties were calculated from the first current step that resulted in an average of 10 Hz firing or greater. This was typically ~ 30 pA (range 0-80 pA) above the rheobase current. A 'burst' was defined as two or more APs with an inter-spike interval (ISI) of 12 ms or less. An initial adaptation ratio was defined as the ISI between the first and second action potentials divided by the ISI between the second and third action potentials. A steady-state adaptation ratio was calculated as the ISI between the third and fourth spikes divided by the ISI of the last two spikes in the train.

For analysis of M1 input, individual trials were averaged together, because we observed little trial-to-trial variability in the responses to optical stimulation. For analysis of excitatory post-synaptic currents (EPSCs), response onset was taken as the first of 20 consecutive data points that exceeded a threshold of 1 SD below the baseline current. A cell was considered to have received input if the peak of the EPSC exceeded 15 times the standard deviation of the baseline preceding the response. This detected EPSC responses of ~5 pA and larger. EPSC amplitude was calculated as the EPSC peak minus the value at EPSC onset. In current clamp, spike probability was measured as the number of trials that elicited a spike divided by the total number of trials.

Statistical comparisons of multiple groups were performed using an ANOVA, followed by multiple pairwise comparisons tests that were corrected using the Tukey-Kramer method. Statistical comparison of proportions or probabilities between groups was tested using a Fisher's Exact Test. Comparison of EPSC amplitudes for within slice recordings was conducted using a paired t-test.

3.4 RESULTS

Anatomical distribution of L5 and L6 pyramidal neurons

To test for M1 input to populations of pyramidal neurons projecting to a specific cortical or subcortical target, we injected fluorescent retrograde tracers in combination with an AAV virus containing the gene for channelrhodopsin-2 (ChR2; see Methods). In all animals a retrograde tracer was co-injected with the ChR2 virus into primary motor cortex (M1), and a

second retrograde tracer was injected into either the ventral posterior medial (VPM) thalamic nucleus, the posteromedial (PoM) nucleus, or the spinal trigeminal nucleus (Sp5) in the brainstem (Figure 3.1). For each injection target, the resulting laminar pattern of labeled neurons in S1 was consistent with known labeling patterns (Wise and Jones, 1977; Chmielowska et al., 1989; Killackey et al., 1989; Deschenes et al., 1994; Alloway et al., 2004; Hattox and Nelson, 2007; Mao et al., 2011). M1-projecting neurons were found in L2/3, L5A, at the boundary between L5 and L6 (L5/6), as well as deep in L6 (Figure 3.1A). Neurons projecting to Sp5, on the other hand, were located exclusively in L5B. Corticothalamic neurons projecting to PoM were found in L5B, in a similar position as the Sp5-projecting neurons, as well as in deep layer 6. Finally, corticothalamic neurons projecting to VPM were only found in L6 and tended to be concentrated in the upper half of L6 as has been previously described (Zhang and Deschenes, 1997).

The retrogradely labeled neurons recorded in this study were grouped into the following five populations: M1-projecting cells in layer 5A (L5A M1-projecting), M1-projecting neurons near the border of layers 5 and 6 (L5/6 M1-projecting), Sp5-projecting neurons, L5B PoM-projecting neurons (PoM-projecting), and VPM-projecting neurons. Only two L6 PoM-projecting neurons were recorded and therefore were not included as an additional population (see Discussion). Depth measurements, normalized to the layer 4/5A border, revealed that each population of pyramidal neurons was consistently positioned in specific locations in L5A, L5B, and/or L6 (Figure 3.1C). Interestingly, each population also tended to be largely segregated within a layer, with only a small degree of overlap between populations (Figure 3.1C). The exception was that PoM-projecting neurons and Sp5-projecting neurons in L5B were completely overlapping in their laminar location (Figure 3.1C).

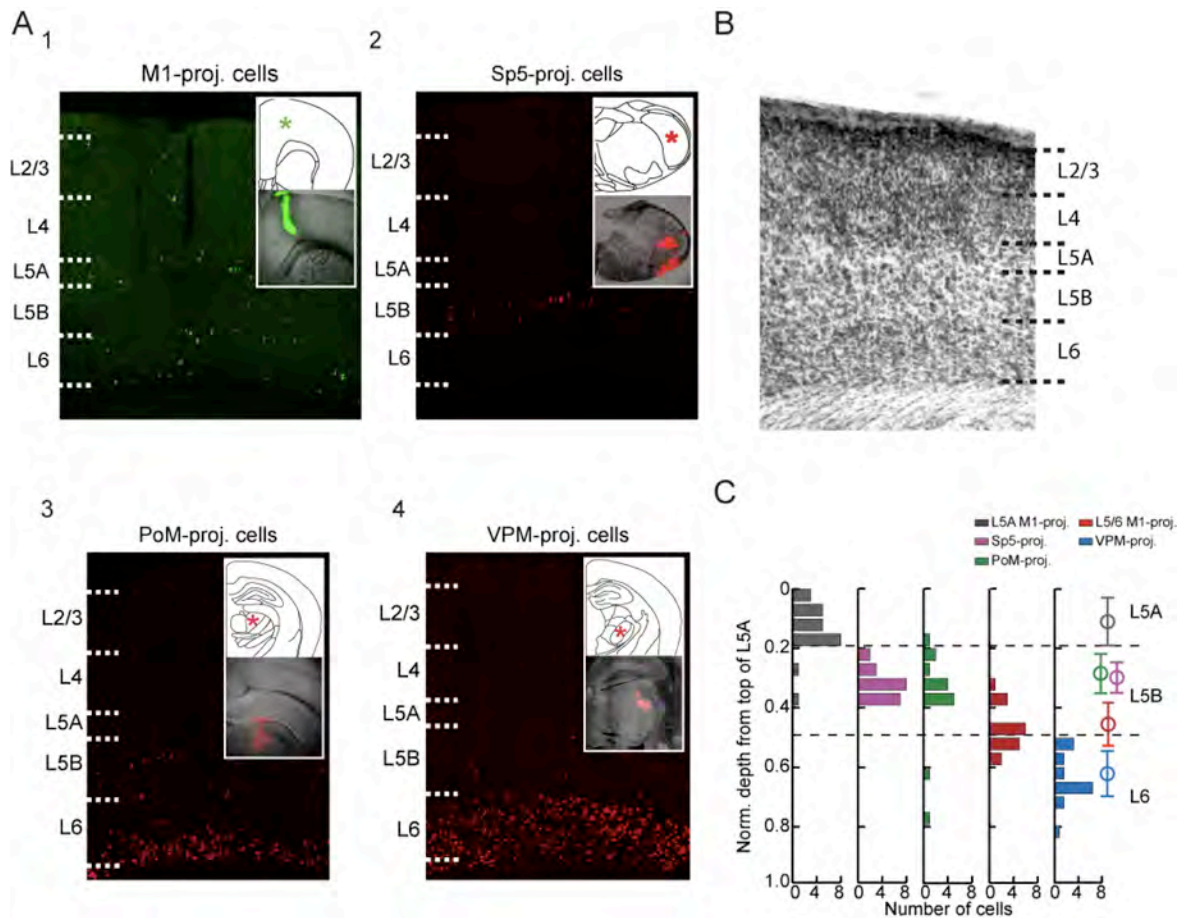


Figure 3.1 Laminal position of S1 pyramidal neurons in L5 and L6 is organized by projection target. (A) Retrobead injections into M1 (1) resulted in retrogradely labeled pyramidal neurons throughout layer 2-6. Labeled cells were especially numerous in L5A and the very bottom of L5 and top of L6. Red fluorescent retrobeads injected into the spinal trigeminal nucleus (2) exclusively labeled S1 neurons in L5B. Injection of retrobeads into the posteromedial (PoM) thalamus (3) labeled neurons in both L5B and L6, whereas retrobeads in the ventral posterior medial (VPM) thalamic nucleus (4) only labeled neurons in L6 of S1. In panels 1-4 an inset illustrates the injection target (top; marked with an asterisk) and an example injection for each target location (bottom). (B) Nissl stained tissue was used to demarcate laminar boundaries (see Methods). (C) Laminal depth measurements for each recorded cell were normalized to the top of L5A. Histograms illustrate the distribution of each population within L5 and L6. Open circles and error bars (right) indicate the mean and standard deviation, respectively, for each population.

Co-expression of both red and green fluorescent retrobeads was observed in a few neurons recorded from animals injected in M1 and Sp5 (n=2; data not shown). These cells were located in the deep part of L5B and were included in both the Sp5- and L5/6 M1-projecting data sets. No other cases of co-labeling were observed in this study. Morphological examination of all pyramidal neurons revealed anatomical differences between the populations that have been previously described by others (Chagnac-Amitai et al., 1990; Deschenes et al., 1994; Zhang and Deschenes, 1997; Hattox and Nelson, 2007; Le Be et al., 2007), and therefore were not quantified further.

Physiological properties of L5 and L6 pyramidal neurons

We compared the electrophysiological characteristics of L5A M1-projecting neurons (n=23), L5/6 M1-projecting neurons (n=19), Sp5-projecting neurons (n=24), L5B PoM-projecting neurons (n=15), and VPM-projecting neurons (n=19). As seen in juvenile mice (Hattox and Nelson, 2007), many subthreshold and action potential characteristics were strongly dependent on the projection target of the pyramidal neuron (Table 1; Figures 3.2&3.3). All statistical relationships are listed in Table 1.

Comparisons of passive properties demonstrated that VPM-projecting neurons had the highest input resistance ($241 \pm 9.1 \text{ M}\Omega$) and PoM- and Sp5-projecting neurons the lowest ($131 \pm 11.3 \text{ M}\Omega$ and $106 \pm 6.9 \text{ M}\Omega$, respectively; Table 1 & Figure 3.3A). These differences are inversely related to the size of the neurons, as Sp5- and PoM-projecting neurons are among the largest neurons in S1 and VPM-projecting neurons the smallest. L5A M1- and L5/6 M1-projecting neurons had significantly more hyperpolarized resting membrane potentials ($-65.2 \pm$

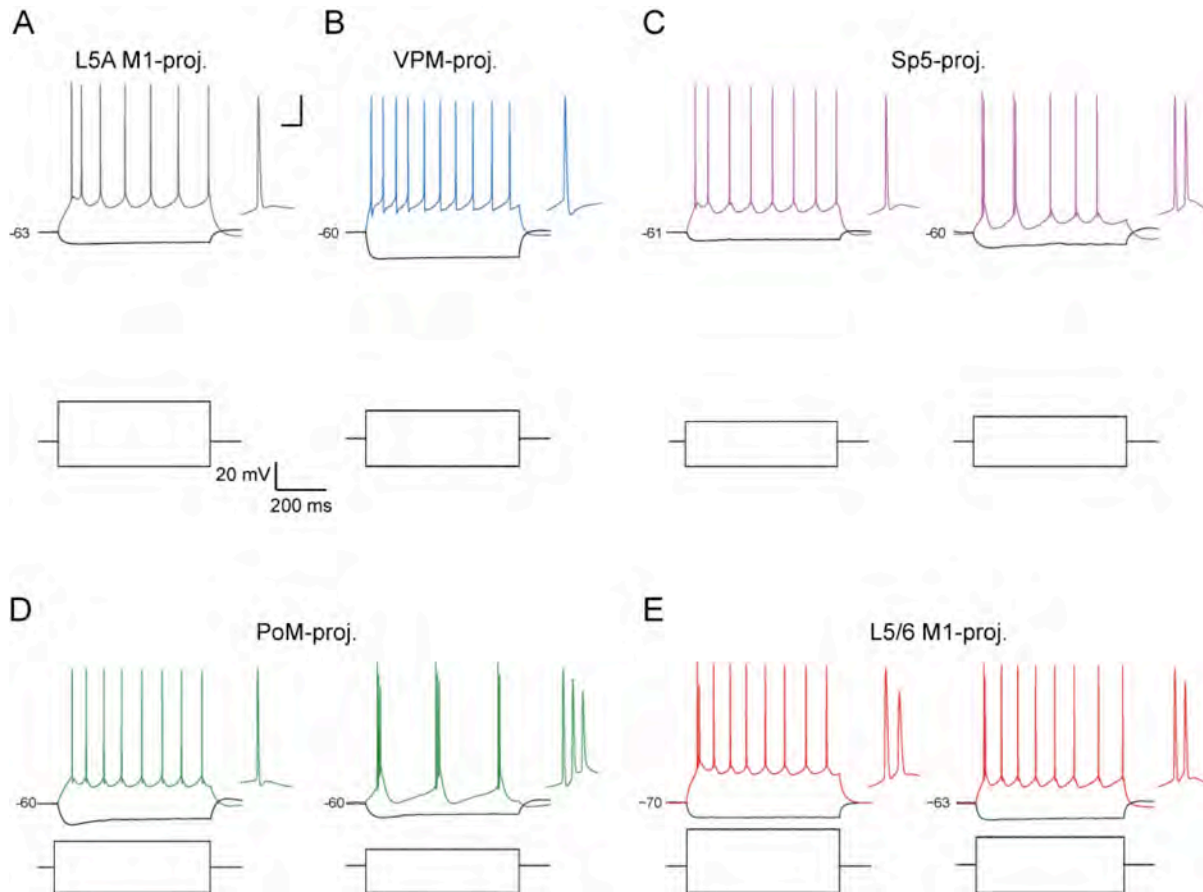


Figure 3.2 Examples of firing properties for S1 pyramidal neuron populations. (A) L5A M1-projecting neurons exhibited mostly regular-spiking behavior, and the first interspike interval (ISI) tended to be shorter than subsequent ISIs. (B) VPM-projecting neurons were also regular-spiking with large AHPs following each action potential. Sp5-projecting neurons (C) and PoM-projecting neurons (D) fired either single spikes that were followed by a depolarizing afterpotential (DAP; C&D, left), a mix of bursting and regular spiking (C, right) or completely bursting dynamics (D, right). (E) L5/6 M1-projecting neurons often fired an initial spike ‘doublet’, and in some cells subsequent spikes were followed by a DAP (right), whereas others were not (left). In each panel the first spike in the train is expanded and shown to the right. Scale bars for expanded spike in each panel are 10 ms and 10 mV.

0.8 mV and -68.8 ± 1.3 mV, respectively) than the other three pyramidal neuron types (Table 1). We also examined the presence of ‘sag’ current, which in pyramidal neurons is mediated by hyperpolarization-activated current (I_h ; Sheets et al, 2011) . PoM-projecting neurons exhibited the most sag current (4.22 ± 0.42 mV) that was significantly larger than sag currents in L5A M1-, VPM-, and L5/6 M1-projecting neurons (Table 1; Figure 3.3B&C). L5/6-M1 projecting neurons in particular had notably small sag currents (0.85 ± 0.26 mV).

Each population of pyramidal neurons exhibited distinct action potential (AP) features as well (Table 1). Sp5- and PoM-projecting neurons had the narrowest action potentials (AP half-width of 1.08 ± 0.05 mV and 1.05 ± 0.04 , respectively) as well as the most hyperpolarized threshold for AP initiation (-41.5 ± 0.6 mV and -42.3 ± 1.3 mV, respectively). By comparison, APs of VPM-projecting neurons were the broadest (AP half-width of 1.50 ± 0.06 ms) and had the most depolarized AP threshold (-33.8 ± 1.0 mV). L5A and L5/6 M1-projecting neurons had APs that were broader than Sp5- or PoM-projecting neurons, but narrower than VPM-projecting neurons (L5A M1: 1.31 ± 0.07 ms; L5/6 M1: 1.38 ± 0.10 ms), and AP thresholds that were more hyperpolarized than VPM-projecting neurons but more depolarized than Sp5- or PoM-projecting neurons (L5A M1: -37.8 ± 0.7 mV; L5/6 M1: -39.0 ± 1.3 mV).

Striking differences between pyramidal neuron subtypes were observed in the voltage trajectory following the action potential. VPM-projecting neurons were distinct in exhibiting a deep, fast afterhyperpolarization (AHP) that was not seen in any of the other cell types (Figure 3.2B and 3.3D&E). Sp5- and PoM-projecting neurons, on the other hand, displayed a small depolarizing afterpotential (DAP) following a spike (Figure 3.2C&D and 3.3F). DAPs were observed in many L5/6 M1-projecting and some L5A M1-projecting neurons as well (Figure 3.3F). In Sp5-, PoM-, and some L5/6 M1-projecting neurons a DAP was present following every

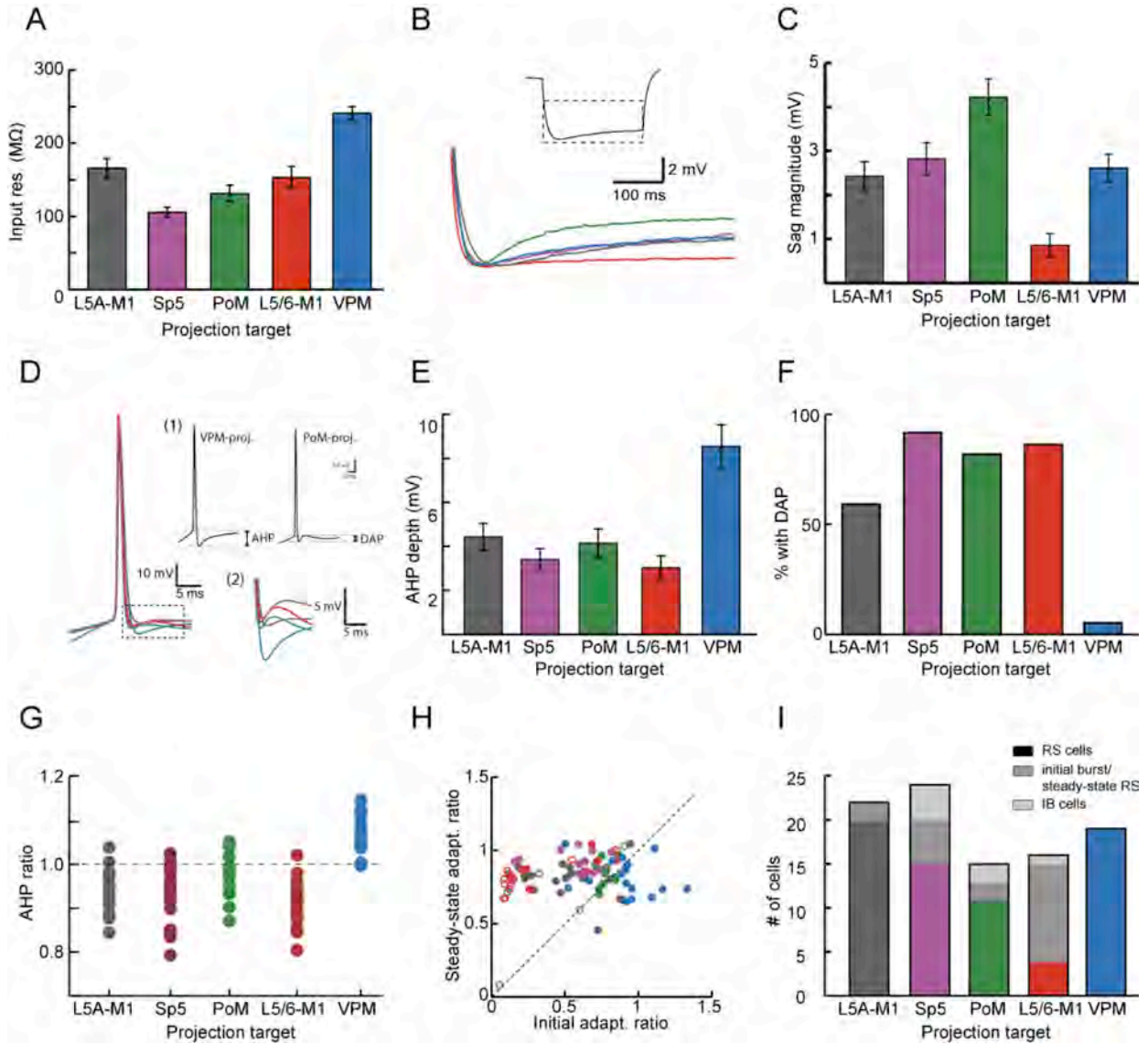


Figure 3.3 Comparison of intrinsic properties between S1 pyramidal neuron populations. (A) Population average of input resistance for each type of pyramidal neuron. (B) Illustration of sag currents. Traces are averaged voltage responses for all neurons in each population. (C) Average sag currents are largest in PoM-projecting neurons and smallest in L5/6 M1-projecting neurons. (D) First spike waveforms (*left*), averaged across neurons, highlight differences between populations. Inset (1) depicts measurements for AHP depth (*left*) and DAP magnitude (*right*). Inset (2) shows expansion of dashed box and highlights differences in the voltage trajectories for each cell type following an action potential. (E) Population average of AHP depth, and (F) the percentage of each population demonstrating DAPs. (G) Ratio of the first to second AHP depth in a spike train. (H) VPM-projecting neurons display distinct adaptation dynamics, as the initial adaptation ratio is higher than the steady-state adaptation ratio. (I)

Number of cells in each population that showed regular spiking (colored bars), initial bursts (medium gray bars), or intrinsic bursting (light gray bars). In panels A-I statistical relationships are not shown and instead are illustrated in Table 1. All bars and error bars indicate mean and SEM. Color scheme in B, D, and H is the same as bar plots in other panels.

spike (Figure 3.2C,D and 3.2E, right). For L5A M1-projecting and the remaining L5/6 M1-projecting neurons a DAP was only found following the first action potential in a train (Figure 3.2A and 3.2E, left).

DAP potentials in pyramidal neurons are calcium-dependent (Friedman et al., 1992) and may underlie the generation of bursting behaviors observed in some pyramidal neurons (Llano and Sherman, 2009). We therefore compared the proportion of pyramidal neurons that were either completely regular-spiking (RS), displayed an initial burst(s) followed by single spikes (B/RS), or exhibited bursting throughout the current step (intrinsic bursting [IB]; Figure 3.3I). VPM-projecting cells were always RS (19/19; Figure 3.2B). L5A-M1 projecting cells were almost always RS cells (22/23; Figure 3.2A) with one cell exhibiting an initial burst followed by regular-spiking behavior (B/RS; Figure 3.3I). Neurons projecting to Sp5 could be RS, B/RS, or IB cells (RS: 12/24; B/RS: 8/24; IB: 4/24; Figure 3.2C); the same was true for PoM-projecting neurons (RS: 9/15; B/RS: 4/15; IB: 2/15; Figure 3.2D). Interestingly, most L5/6 M1-projecting neurons were B/RS cells (11/19) as they often exhibited an initial spike ‘doublet’ (Figure 3.2E). Most of the remaining L5/6 M1-projecting neurons were RS (7/19) although one was an IB cell (1/19).

Finally, we examined changes in spike properties across the spike train. In VPM-projecting neurons AHP depth decreased from the 1st spike to subsequent spikes, resulting in an AHP ratio that was larger than 1 (Figure 3.3G). In the other cell types, the first AHP was

Table 1. Intrinsic properties of S1 pyramidal neurons

Property	PROJECTION TARGET					SIGNIFICANCE LEVEL		
	L5A M1 (1)	Sp5 (2)	PoM (3)	L5/6 M1 (4)	VPM (5)	p < 0.05	p < 0.01	p < 0.005
N	23	24	15	19	19			
<i>Passive properties</i>								
Input resistance (MΩ)	166 ± 13.7	106 ± 6.9	131 ± 11.3	153 ± 14.5	241 ± 9.1	2 : 4	1 : 2	5 : 1,2,3,4
V _{rest} (mV)	-65.2 ± 0.8	-62.3 ± 0.9	-62.4 ± 1.4	-68.8 ± 1.3	-61.0 ± 1.1	1 : 5		4 : 2,3,5
Sag current (mV)	2.43 ± 0.32	2.82 ± 0.36	4.22 ± 0.42	0.85 ± 0.26	2.61 ± 0.32	3 : 5	4 : 1,2,3	3 : 1,4 4 : 5
<i>Action potential (AP) features</i>								
AP half-width (ms)	1.31 ± 0.07	1.08 ± 0.05	1.05 ± 0.04	1.38 ± 0.10	1.50 ± 0.06	1 : 2,3	4 : 2,3	5 : 2,3
AP threshold (mV)	-37.8 ± 0.7	-41.5 ± 0.6	-42.3 ± 1.3	-39.0 ± 1.3	-33.8 ± 1.0	1 : 2,3,5		5 : 2,3,4
AP peak (mV)	43.0 ± 1.0	42.7 ± 1.7	39.8 ± 2.3	43.8 ± 1.7	41.6 ± 1.7	-----n.s.-----		
AHP depth (mV)	4.4 ± 0.6	3.4 ± 0.5	4.9 ± 0.8	3.1 ± 0.7	8.6 ± 1.0		5 : 3	5 : 1,2,4
% with depolarizing after-potential (DAP)	59% (13/22)	92% (22/24)	87% (13/15)	94% (15/16)	5% (1/19)	1 : 2	1 : 4	5 : 1,2,3,4
<i>Spiking properties</i>								
Rheobase (pA)	71.0 ± 9.6	77.5 ± 7.7	60.0 ± 5.5	96.3 ± 10.4	79.0 ± 7.6	-----n.s.-----		
Initial adaptation ratio	0.52 ± 0.04	0.50 ± 0.06	0.69 ± 0.07	0.38 ± 0.08	0.90 ± 0.05	3 : 4		5 : 1,2,4
Steady-state adaptation ratio	0.72 ± 0.02	0.88 ± 0.03	0.79 ± 0.04	0.71 ± 0.07	0.70 ± 0.04	2 : 1,4,5		
% with initial burst	4% (1/23)	33% (8/24)	27% (4/15)	58% (11/19)	0% (0/19)	1 : 2		1 : 4
						4 : 2,3		5 : 2,4
						3 : 5		
% with steady-state bursts	0% (0/23)	17% (4/24)	13% (2/15)	5% (1/19)	0% (0/19)	-----n.s.-----		

shallower than subsequent AHPs and AHP ratios were less than 1 (Figure 3.3G). VPM-projecting neurons also had different adaptation dynamics, with the inter-spike interval (ISI) gradually increasing across the spike train (Figure 3.2B). This resulted in initial adaptation ratios larger than the steady-state adaptation ratios (0.90 ± 0.05 vs. 0.70 ± 0.04 ; Figure 3.3H; Table 1). For each of the other cell types, the initial adaptation ratio was typically smaller than the steady-

state adaptation ratio. This was particularly evident in the neurons that exhibited an initial burst (Figure 3.3H; open circles). IB cells were not included in analysis of adaptation dynamics, as the irregular spike patterns within and between bursts yielded highly variable adaptation values.

Motor cortex input to L5 and L6 S1 pyramidal neurons

All animals were injected with a virus containing the ChR2 gene and retrobeads in M1. The resultant fluorescence in S1 therefore included both M1 axons, labeled with ChR2-mCherry, and M1-projecting neurons, labeled with retrobeads. In the same animal a second population of S1 neurons was labeled with retrograde tracers, either Sp5-, PoM-, or VPM-projecting neurons. Sp5-, PoM-, and VPM-projecting neurons were labeled and recorded in separate animals. In order to account for possible differences in ChR2 expression in the different animals, all experiments were conducted using a paired approach in which L5A M1-projecting neurons were recorded in the same slice as cells from one of the other pyramidal neuron populations: L5/6 M1-, Sp5-, PoM-, or VPM-projecting neurons (Figures 3.4&3.5). Paired comparisons could then be used to determine the strength of M1 inputs relative to a standard population (L5A M1 projecting neurons).

Recording from a L5A M1-projecting neuron and an Sp5-projecting neuron within the same slice demonstrated that optical stimulation of ChR2-M1 axons evoked a short-latency, excitatory post-synaptic current (EPSC) that was similar in amplitude for each cell (Figure 3.4B). Comparison across the population of paired recordings (n=12 pairs) determined that the average amplitude of M1 input was equivalent for each population (L5A M1-proj: 113.32 ± 25.97 pA, Sp5-proj: 138.09 ± 34.51 pA; $p=0.31$; Figure 3.4D). Similar results were found for L5A M1- and PoM-projecting neurons (n=9 pairs), indicating that L5A M1-projecting and PoM-projecting

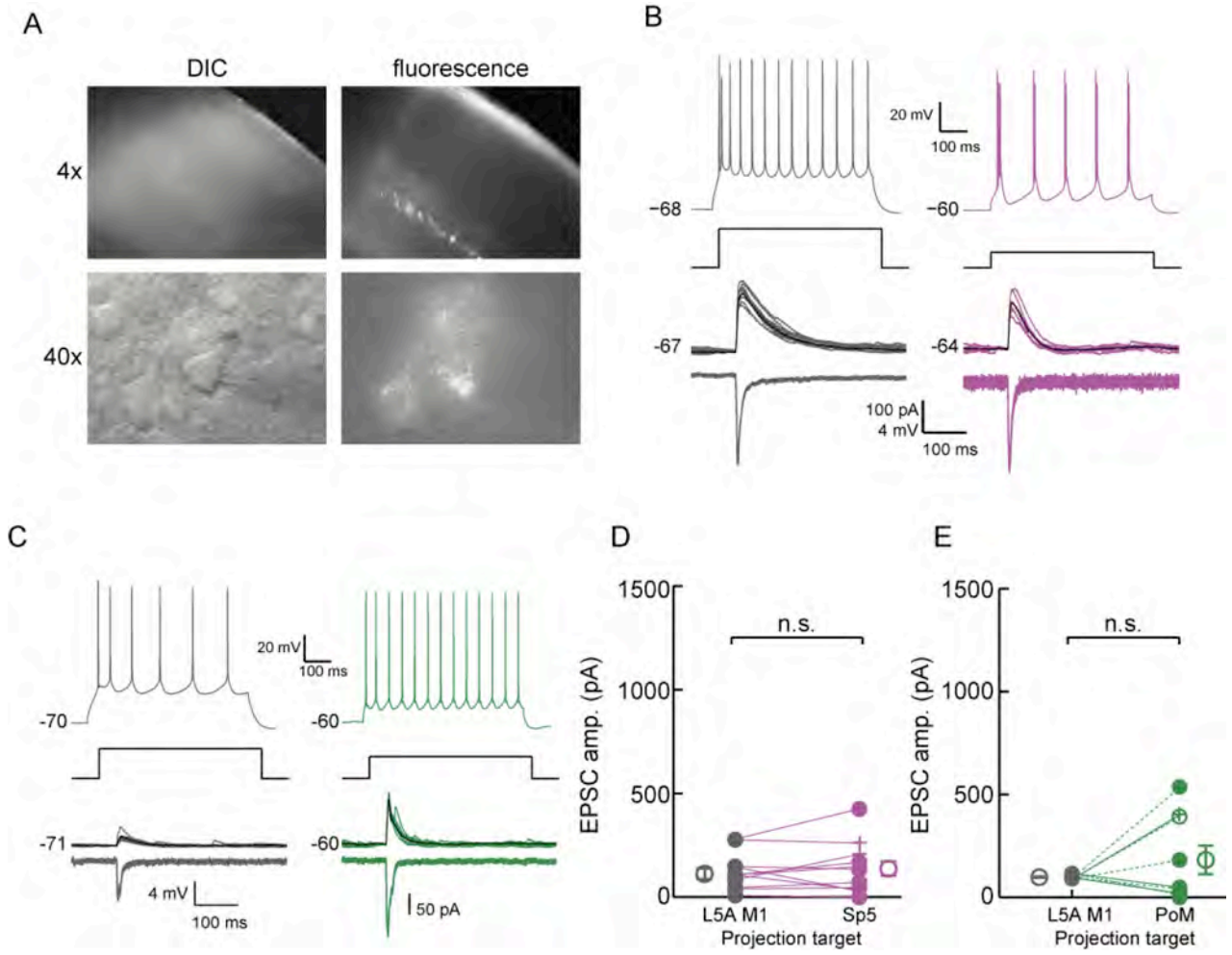


Figure 3.4 M1 input to L5A M1-, Sp5-, and PoM-projecting neurons is equivalent. (A) Example of a S1 slice at 4X magnification under DIC illumination (top left) and epifluorescence (top right) shows bright, retrogradely labeled neurons in L5B that project to Sp5. Diffuse fluorescent labeling is also observed, indicating presence of ChR2-expressing M1 axon terminals. Example of a recorded pyramidal neuron (bottom left) that expressed red fluorescent retrobeads (bottom right). (B) L5A M1-projecting neuron (gray; left) and Sp5-projecting neuron (magenta; right) recorded in the same slice showed characteristic current step responses (top), received similar excitatory input from M1 in current-clamp (middle) or voltage-clamp (bottom). In current-clamp multiple traces were overlaid and the average EPSP shown in black. EPSC traces show only the trial-averaged response. (C) L5A M1-projecting and L5B PoM-projecting neuron recorded in the same slice. Conventions as in (B). Cells were held at -80 mV for voltage-clamp recordings in (B) and (C). (D) Population averaged EPSC responses to M1 stimulation illustrate that Sp5-projecting neurons and (E) PoM-projecting neurons receive the same magnitude of M1 input as L5A M1-projecting neurons (L5A M1-proj. vs. Sp5-proj: $p=0.31$; L5A M1-proj. vs. PoM-proj: $p=0.27$). In D&E, closed circles denote RS neurons, open circles indicate B/RS cells, and plus signs are IB cells. Open circles and error bars to the left and right of the paired data are the population mean and SEM.

neurons also receive equivalent amount of input from M1 (L5A M1-proj: 97.59 ± 3.44 pA, PoM-proj: 181.55 ± 69.21 pA; $p=0.27$; Figure 3.4E). Sp5- and PoM-projecting cells that exhibited either initial bursts (Figure 3.4D&E, open circles) or were intrinsically bursting (Figure 3.4D&E, plus signs) received a comparable magnitude of M1 input as the cells that were regular spiking (Figure 3.4D&E, closed circles).

In contrast to Sp5- and PoM-projecting neurons, many L5/6 M1-projecting neurons had very large amplitude EPSC responses to optical stimulation (Figure 3.5A). Indeed, M1 input was often strong enough to evoke action potentials in the L5/6 M1-projecting neuron; action potentials were never observed in L5A M1-projecting neurons (Figure 3.5A; see also below). Across the population ($n=9$ pairs), L5/6 M1-projecting neurons on average received significantly stronger input from M1 than the L5A M1-projecting neurons (158.62 ± 50.14 pA vs. 700.62 ± 184.66 pA; $p=0.01$; Figure 3.5C). Many of the L5/6 M1-projecting neurons in our paired sample that received strong M1 input exhibited an initial spike burst (Figure 3.5C; open circles); however, large amplitude M1 inputs were observed in regular spiking cells as well (Figure 3.5C; closed circles).

M1 input to VPM-projecting neurons was also distinct, in this case because these cells received small input from M1 ($n=16$ pairs; Figure 3.5B&D). Across all paired recordings, the amplitude of M1 input to VPM-projecting neurons was significantly less than input to the L5A M1-projecting cells (161.55 ± 31.58 pA vs. 54.55 ± 10.88 pA; $p=0.008$).

The mean amplitude of M1 input to each population of paired L5A M1-projecting neurons did not differ ($p=0.43$), and therefore we pooled data for all pyramidal cell types in order to make direct comparisons among pyramidal neuron populations (Figure 3.6A). Group data showed similar results as the paired data, demonstrating that synaptic inputs from M1

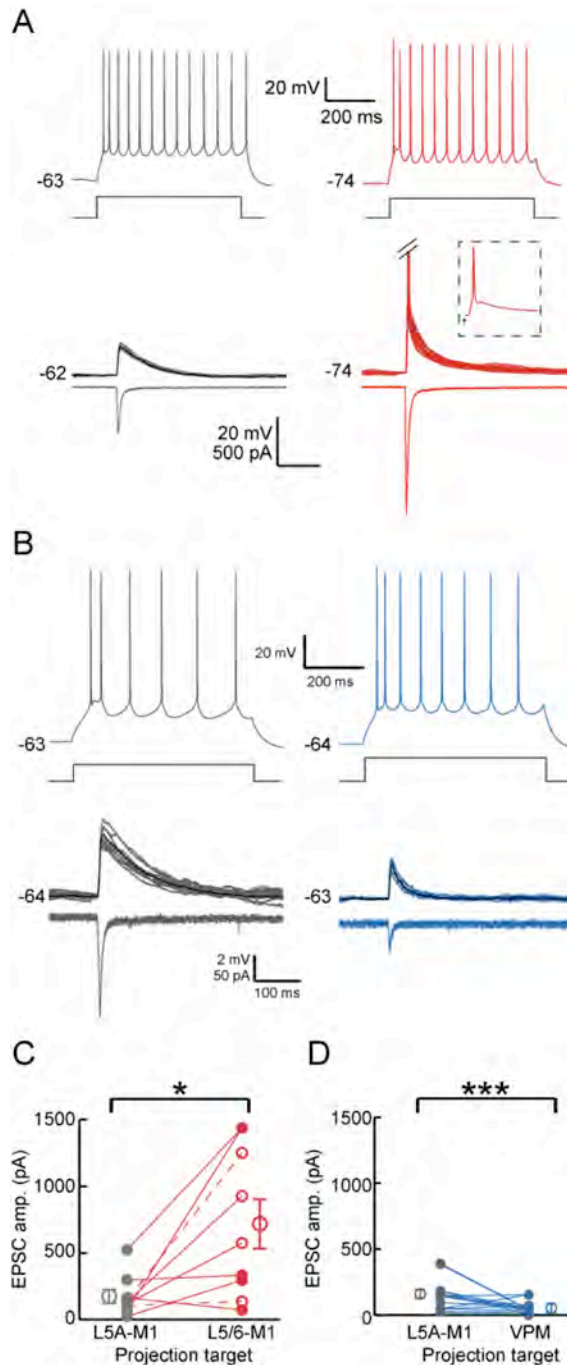


Figure 3.5 M1 provides strong input to L5/6 M1-projecting neurons and weak input to VPM-projecting neurons. (A) Example L5A M1-projecting (left, gray) and L5/6 M1-projecting neuron (right, red) responses to current steps (top). In this pair the L5/6 M1-projecting neuron received larger M1 input as seen in current-clamp (middle) and voltage-clamp (bottom). Note that the L5/6 M1-projecting neuron fired spikes on several trials (one trial shown in dashed box inset). Cells were recorded in the same slice. (B) Same as (A) but for a L5A M1-projecting neuron (gray; left) and VPM-projecting neuron (blue) recorded in the same slice. (C) On average, the L5/6 M1-projecting neurons received significantly larger input than the paired L5A M1-projecting cells ($p < 0.05$). (D) VPM-projecting

neurons received significantly less M1 input than the paired L5A M1-projecting neurons ($p < 0.005$). Conventions for (C) and (D) are same as Figure 3.4D&E.

differentially targeted populations of pyramidal neurons based on their axonal projections (ANOVA $p < 0.00001$). Notably, L5/6 M1-projecting neurons received significantly larger inputs than any other population, whereas VPM-projecting neurons received the smallest (Figure 3.6A).

Optical stimulation of ChR2-M1 terminals evokes suprathreshold responses in S1 pyramidal neurons

To determine if M1 inputs were strong enough to evoke suprathreshold responses in any of the retrogradely identified populations of neurons in this study we recorded in current-clamp and measured spiking probability in response to optical stimulation (Figure 3.6). Many pyramidal neurons responded by spiking, usually those in each population that received the largest amplitude of excitatory input from M1 (Figure 3.6B). For example, the L5/6 M1-projecting population had the highest proportion of neurons exhibiting suprathreshold responses ($n=9/19$). Interestingly, although L5A M1-, Sp5-, and PoM-projecting neurons received similar mean amplitudes of input from M1 (Figure 3.6A), L5A M1-projecting neurons never spiked ($n=0/23$) whereas many Sp5- and PoM-projecting neurons did ($n=6/24$ and $n=3/15$, respectively). VPM-projecting neurons received very small excitatory inputs from M1 (Figure 3.6A) and, despite having the largest input resistance (Figure 3.3A), rarely responded by firing action potentials ($n=1/20$).

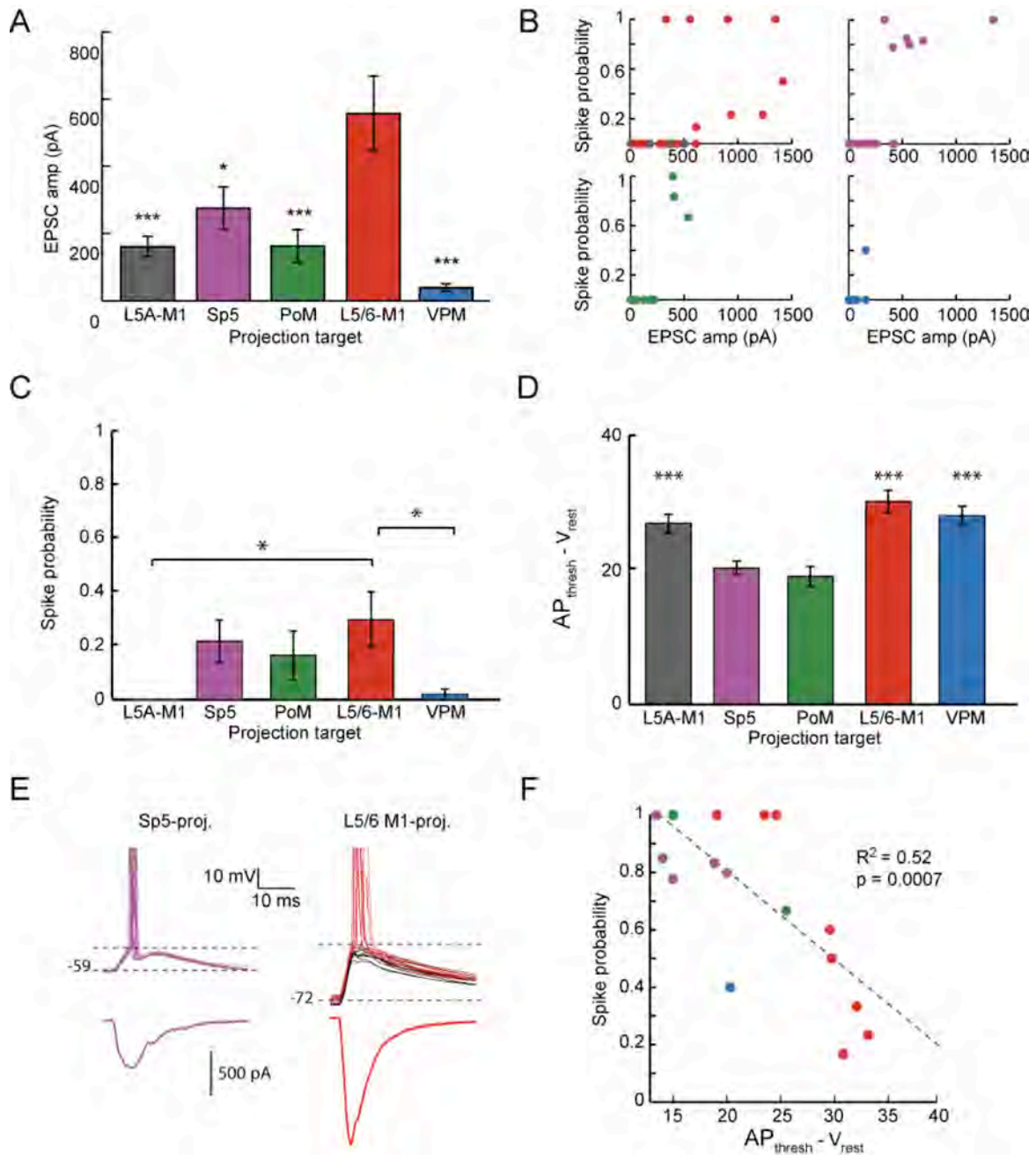


Figure 3.6 Optical stimulation of Chr2-M1 terminals evoked suprathreshold responses in multiple types of S1 pyramidal neurons. (A) Population average of EPSC amplitude evoked by M1 stimulation showed that L5/6 M1-projecting neurons received significantly larger M1 input than the other pyramidal neuron populations ($p < 0.005$). (B) Pyramidal neurons in each population with largest M1-evoked EPSC often showed non-zero spike probabilities when recorded in current-clamp. Note that upper left panel shows both L5A M1- (gray circles) and L5/6 M1-projecting neurons (red circles). Color conventions as in (A). (C) Average spike probability in L5/6 M1 projecting

neurons was significantly greater than in L5A M1- and VPM-projecting neurons ($p < 0.05$) but not Sp5- and PoM-projecting neurons ($p > 0.05$). (D) Average voltage difference from resting membrane potential to action potential threshold was significantly smaller in Sp5- and PoM-projecting neurons than other pyramidal neuron classes ($p < 0.005$). (E) Example Sp5-projecting neuron (left) that spiked on each trial (magenta), and a L5/6 M1-projecting neuron (right) that fired on some trials (red) but not others (black). (F) Scatterplot of the distance between rest and AP threshold and the M1-evoked spike probability for all neurons that spiked. Color conventions as in (D). Dashed line shows the linear, least squares fit.

Average spiking probability was highest in L5/6 M1-projecting neurons (0.30 ± 0.09), but was not significantly different than Sp5-projecting neurons (0.23 ± 0.08) or PoM-projecting neurons (0.19 ± 0.10 ; Figure 3.6C) despite those neurons receiving significantly less excitatory current from M1. Sp5- and PoM-projecting neurons reliably fired action potentials for EPSC amplitudes around 500 pA as did a few L5/6 M1-projecting neurons; other L5/6 M1-projecting neurons only spiked occasionally despite receiving excitatory inputs of 1000 pA or greater (Figure 3.6B).

Neurons differed with respect to the voltage difference between resting potential and AP threshold. Such differences were smallest for Sp5- and PoM-projecting pyramidal neurons ($p < 0.005$; Figure 3.6D). Differences within the populations reflect the fact that Sp5- and PoM-projecting neurons have lower voltage thresholds for AP generation (Table 1) whereas both L5A M1- and L5/6 M1-projecting pyramidal neurons exhibit more hyperpolarized resting membrane potentials (Figure 3.6E; Table 1). We hypothesized that the smaller voltage difference in Sp5- and PoM-projecting neurons compared to L5/6 M1-projecting neurons can account for differences in the likelihood that the cells would fire a spike in response to M1 input (Figure 3.6E). We confirmed this by comparing the voltage difference between resting and AP threshold with spike probability output. Neurons with the smallest differences, which tended to be Sp5-

and PoM-projecting neurons, had the highest spike probability, while those with larger voltage differences had lower probability of spiking, despite many of them receiving larger amplitude EPSCs (Figure 3.6F).

Neurons in the mid-depths of the infragranular layers, on average, received the largest M1 inputs (Figure 3.7A). Indeed, comparison of spike probability as a function of the laminar position showed that neurons mostly likely to fire action potentials in response to M1 synaptic input (i.e. Sp5-, PoM-, or L5/6 M1-projecting neurons) are primarily located within L5B or upper L6 (Figure 3.7B). This sub-laminar ‘zone’ within S1 may therefore be the most heavily influenced by increased activity in M1, such as occurs *in vivo* during whisking.

3.5 DISCUSSION

Here, we combined optogenetic techniques with retrograde tracers to investigate the anatomy, physiology, and magnitude of M1 input to selected classes of S1 pyramidal neurons in the infragranular layers. Retrograde tracers allowed unambiguous identification of the projection target for each recorded neuron. Injection of an AAV virus containing the gene for ChR2 permitted selective activation of synaptic inputs originating in M1.

Our results demonstrate a number of important findings. First, we found that pyramidal neuron subtypes that project to a particular target are organized within L5 and L6 such that, in most cases, little spatial overlap occurs among populations. Next, consistent with previous studies in juvenile mice, S1 pyramidal neurons exhibited a variety of physiological characteristics that correlated with the projection target of the neuron. L5A M1-projecting

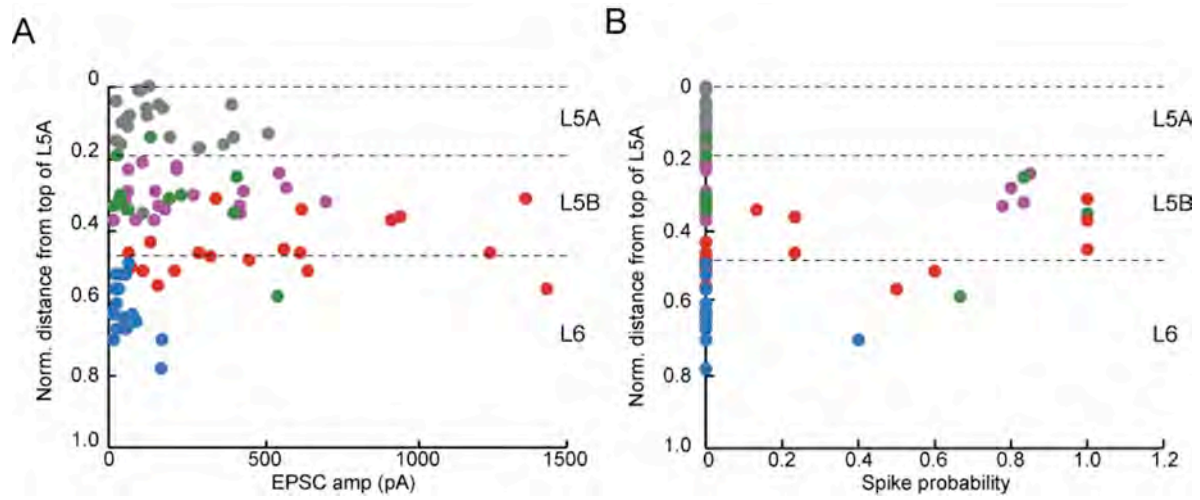


Figure 3.7 Activation of M1 primarily modulates L5B and upper L6 of S1. (A) Population graph of the amplitude of the M1 evoked EPSC as a function of the depth for each neuron indicates that M1 inputs are particularly strong around the border between L5 and L6. (B) The majority of the neurons that spike in response to optical stimulation of M1 axon terminals are located in L5B or the top of L6.

neurons exhibited classic pyramidal neuron characteristics that are consistent with known properties of corticocortical neurons. Intriguingly, VPM-projecting neurons possessed characteristics not typically associated with pyramidal neurons, such as high input resistance and fast, deep AHPs. Sp5-, PoM, and L5/6 M1-projecting neurons showed the most diversity with regards to intrinsic properties, as some neurons were regular spiking upon depolarization whereas others produced bursts of action potentials. Finally, M1 input was highly dependent on the projection target of the neuron, with VPM-projecting neurons receiving small input and L5/6 M1-projecting neurons receiving large input.

For this study we focused on S1 pyramidal neurons in L5 and/or L6 that projected to one of four targets (i.e. M1, Sp5, PoM, or VPM). Many other classes of pyramidal neurons exist in the infragranular layers, including neurons projecting to contralateral S1, secondary somatosensory cortex (S2), striatum, and the pons among others (White and DeAmicis, 1977; Wise and Jones, 1977; Wilson, 1987; Killackey et al., 1989; Petrof et al., 2012). Future studies will be needed to obtain a complete map of M1 inputs to all types of pyramidal neurons in S1. However, our results here demonstrate a remarkable degree of specialization and specificity regards to sub-laminar organization, intrinsic properties and spike dynamics, and M1 synaptic input to these S1 pyramidal neuron populations.

Comparison of L5A and L5/6 M1-projecting neurons in S1

In our study we distinguished between two populations of M1-projecting neurons on the basis of their non-overlapping locations within L5. The first population was restricted to L5A, residing just below the boundary between L4 and L5. These neurons were thin tufted pyramidal neurons (data not shown) and demonstrated typical corticocortical spike properties. L5A M1-

projecting neurons were relatively homogeneous in their intrinsic properties, exhibiting predominantly regular-spiking behavior and similar passive properties and spike adaptation dynamics as other corticocortical neurons (Hattox and Nelson, 2007).

The second population of M1-projecting neurons was located near the L5/6 boundary. M1-projecting neurons have been previously shown to exist in the deep layers in rats (Zhang and Deschenes, 1997; Alloway et al., 2004). The physiology of L5/6 M1-projecting neurons was heterogeneous. Many displayed similar passive and spike dynamics as the L5A M1-projecting neurons; these cells may therefore also be corticocortical neurons that have M1 as their primary target. However, other L5/6 M1-projecting neurons instead resembled Sp5- and PoM-projecting neurons based on the presence of a DAP following each action potential and bursting tendencies. Given that we observed a few instances of co-labeled neurons in animals injected with retrobeads in M1 and Sp5, perhaps those M1-projecting neurons resembling Sp5-projecting neurons are actually corticofugal neurons (e.g. Sp5-projecting) that send a collateral to M1.

Based on our previous study, we hypothesized that a specific class of pyramidal neurons near the top of L6 received particularly strong input from M1. The finding that it was a subset of M1-projecting neurons only located around the L5/6 border was unexpected (see General Discussion). It is especially interesting that L5/6 M1-projecting neurons received significantly stronger input than L5A M1-projecting neurons. A previous study also found that M1-projecting neurons in L5A, as well as those in L2/3, do not receive especially strong input from M1 (Mao et al., 2011). Together, these results suggest a unique role for L5/6 M1-projecting neurons in serving as a sensory feedback signal to M1 during conditions when M1 activity is high.

Similarities between L5B PoM- and Sp5-projecting pyramidal neurons

Sp5-projecting neurons and PoM-projecting neurons in L5B were equivalent for all the intrinsic properties we measured and were intermixed within layer 5B. L5B PoM- and Sp5-projecting neurons, therefore, appear to be more similar to each other than to other types of L5 pyramidal neuron types, such as M1-projecting neurons and VPM-projecting neurons (this study), or callosal and corticostriatal neurons (Hattox and Nelson, 2007). Sp5-projecting neurons send a collateral to PoM, giving rise to the possibility that Sp5- and PoM-projecting neurons in L5B may be largely the same population of neurons (Deschenes et al., 1994). However, co-labeling within the same animal indicated that only a small percentage of Sp5- and PoM-projecting neurons project to both targets (Hattox and Nelson, 2007). Either way, our results suggest that PoM-projecting and Sp5 projecting neurons in layer 5B have similar or closely related functions.

Many Sp5- and PoM-projecting neurons exhibited bursting properties, either initially or continuously in response to current step injection. In the auditory system, only corticothalamic neurons located in L5B, not those in L6, exhibited bursting behaviors (Llano and Sherman, 2009), similar to our results in this study within S1. Throughout sensory and motor cortices, intrinsically bursting (IB) neurons are large, thick-tufted pyramidal neurons that reside in L5B and project subcortically, such as corticotectal and corticopontine neurons (Tseng and Prince, 1993; Wang and McCormick, 1993; Kasper et al., 1994; Rumberger et al., 1998). Across the cortex, therefore, neurons with intrinsic bursting behaviors appear to be restricted to pyramidal neurons in L5B that project to the thalamus, midbrain, or brainstem.

Interestingly, PoM- and Sp5-projecting neurons that were regular spiking (RS) still displayed different dynamics from regular spiking M1- or VPM-projecting neurons. In Sp5- and PoM-projecting neurons, every action potential is followed by a small depolarization (DAP).

DAPs are calcium-dependent (Friedman et al., 1992), and it has been suggested that the DAP underlies the bursting behavior in these neurons (Llano and Sherman, 2009). Therefore, RS neurons that exhibit DAPs may be capable of firing in bursts to certain types of stimuli and/or under certain conditions (Schwindt et al., 1997; Schwindt and Crill, 1999). This is in contrast to early electrophysiological studies, which classified neurons as either RS or IB cells (McCormick et al., 1985; Agmon and Connors, 1989). Thus, in the case of PoM- and Sp5-projecting neurons, RS and IB neurons may possess the same underlying dynamics and ionic currents and could represent a continuum of a single cell type, not two discrete populations (Schwindt et al., 1997). Furthermore, bursting behavior is developmentally regulated and is only seen in adult corticothalamic neurons (Kasper et al., 1994; Llano and Sherman, 2009). In juveniles, all L5B corticothalamic and corticotrigeminal neurons are regular spiking but still exhibit DAPs (Hattox and Nelson, 2007; Llano and Sherman, 2009).

PoM- and Sp5-projecting neurons received only moderate amount of synaptic input from M1, yet because of their intrinsic properties fired relatively easily in response to M1 synaptic stimulation. *In vivo*, L5B thick tufted neurons, which likely include PoM- and Sp5-projecting neurons, have the highest spontaneous firing rates of all S1 pyramidal neurons during both quiescent and whisking brain states (de Kock et al., 2007; de Kock and Sakmann, 2009), and also have robust whisker-evoked sensory responses (de Kock et al., 2007). In a similar manner to our results with M1 input, Sp5- and PoM-projecting neurons may therefore fire robustly to moderate amounts of synaptic input from many sources, both sensory and motor.

Unique properties of VPM-projecting neurons

Of the pyramidal neuron populations examined here, VPM-projecting neurons were the most distinct with regards to intrinsic properties (Figure 3.2; Table 1). VPM-projecting neurons had high input resistance, which is probably related to their compact size, and exhibited deep AHPs, slow adaptation dynamics, and were always regular-spiking. It is worth noting that the few L6 PoM-projecting neurons we recorded (n=2) resembled L6 VPM-projecting neurons in their intrinsic properties, not the L5B PoM-projecting neurons (data not shown). Therefore, as suggested for auditory cortex (Llano and Sherman, 2009), corticothalamic feedback in S1 may be laminar, not projection, specific with corticothalamic neurons in L5B (i.e. PoM-projecting only) having a different role than corticothalamic neurons in L6 (i.e. VPM- and PoM-projecting;).

M1 sends a dense projection to L6 of S1 (Zhang and Deschenes, 1998), where VPM-projecting neurons comprise about half of all neurons (Zhang and Deschenes, 1997). It was therefore unexpected that M1 input instead strongly contacted the M1-projecting neurons in the very top of L6, while the input to VPM-projecting neurons was actually the weakest of all the pyramidal neuron populations we examined in this study (Figure 3.6A). Interestingly, enhancement of M1 activity *in vivo* activates VPM-projecting neurons (Lee et al., 2008). This engagement of VPM-projecting neurons *in vivo* may require the conjunction of M1 input with another source of input, perhaps one that is sensory in nature (see General Discussion).

4.0 GENERAL DISCUSSION

4.1 SUMMARY OF FINDINGS

The function of the pathway from M1 to S1 continues to be examined, but these studies have laid a foundation for understanding how increased activity in M1, as occurs during whisking, will engage excitatory and inhibitory circuits in layers 2-6 of S1. Our findings demonstrate a broad excitation of excitatory and inhibitory neurons in the supragranular and infragranular layers, but a striking paucity of connections with barrel circuits in L4. Furthermore, our results suggest that while pyramidal cells in the superficial layers receive only weak excitation, specific pyramidal cell populations in the infragranular layers are driven strongly by M1. This increased excitation in deep layers is mirrored by stronger recruitment of inhibition, specifically of FS inhibitory interneurons in L5 and L6. Most inhibitory and pyramidal cells that receive direct M1 input also receive disynaptic inhibition, indicating an important role for inhibition in regulating excitatory connections between M1 and S1.

4.2 EFFECTS OF M1 ACTIVATION ON S1 CIRCUITS

4.2.1 Differential contributions of FS and SOM inhibitory interneurons

One main finding of this work is that M1 inputs ubiquitously contact two main inhibitory interneuron classes, FS and SOM inhibitory cells. Because M1 synapses onto both the perisomatic and apical dendrite of individual S1 pyramidal neurons (Petreanu et al., 2009), M1 may recruit somatic inhibition (i.e. FS cells) and dendritic inhibition (i.e. SOM interneurons) to regulate the excitatory drive to both subcellular domains.

Although both inhibitory interneuron subtypes were broadly contacted by M1, synapses onto FS cells were large but strongly depressing while SOM inhibitory interneurons received weak, facilitating synaptic input. These dynamics are strikingly similar to those for local cortical excitatory synapses and, in the case of FS cells, thalamocortical input. Therefore, FS inhibitory interneurons may regulate the excitatory drive from M1 in a similar manner to thalamocortical input; that is by sharpening the S1 response to the onset of M1 activity through feedforward somatic inhibition. In contrast, SOM inhibitory interneurons, due to the facilitating nature of excitatory synapses onto them, are maximally activated by high-frequency bursts of action potentials in pre-synaptic pyramidal neurons (Kapfer et al., 2007; Berger et al., 2010). Bursts of action potentials caused by excitatory input to the apical dendrite are inhibited by dendritic inhibition mediated, at least in part, by SOM interneurons (reviewed in Palmer et al, 2012). Therefore, the facilitating nature of excitatory inputs to SOM cells, both local and long-range, may serve to recruit dendritic inhibition and thereby regulate the prevalence and/or propagation of bursting activity in pyramidal neurons.

4.2.2 M1 contribution to state-dependent changes in S1 neurons

Cortical brain states during active behavior, relative to quiescent or anesthetized periods, are associated with increased neuromodulator actions, altered thalamocortical drive, and increased intercortical interactions. Specifically, in S1 this translates to decreased afferent drive from VPM and increased input from M1 and possibly other cortical areas. Heightened corticocortical activity is likely responsible for the increased levels of calcium activity in the apical dendrites of L5 pyramidal neurons during awake behaving states (Murayama and Larkum, 2009). In fact, a recent study found that calcium-dependent dendritic potentials occur in the apical dendrites of L5 S1 pyramidal neurons during active but not passive touch; furthermore, these dendritic potentials were dependent on input from M1 (Xu et al., 2012). The broad, but subthreshold M1 inputs to L2/3 and L5 S1 pyramidal neurons we reported in the first study may serve to heighten activity in the apical dendrites, which when combined with other inputs, perhaps sensory in nature, could contribute to the generation of dendritic calcium potentials. How increased activity in the apical dendrites *in vivo* translates to spike output at the soma is not yet clear, but a matter of ongoing research.

Our results in the second study add several important points regarding the function of M1 input to S1. First, M1 input to pyramidal neurons varies based on their projection target. Therefore, the function of the M1 to S1 pathway may broadly modulate S1 neurons, but also have circuit-specific effects through stronger inputs to certain types of pyramidal neurons. Secondly, the intrinsic properties also vary depending on the projection target of the neuron.

These results emphasize that M1 feedback does not equally engage all S1 neurons and therefore needs to be thought of in the context of specific circuits. For example, why do only pyramidal neurons that project subcortically exhibit intrinsic bursting? Perhaps M1 input during

active touch produces bursts of action potentials in PoM- and Sp5-projecting neurons but only weakly excites VPM-projecting neurons. Why would this be important for the system? To address this we will consider the specific type of projection neurons and their properties that we looked at in these studies.

4.3 M1 REGULATION OF CORTICAL AND SUBCORTICAL S1 PROJECTIONS

4.3.1 M1-S1 reciprocal circuitry

Regarding the reciprocal S1 to M1 pathway, optical mapping studies indicate that M1 neurons in L2/3 and L5A receive stronger input from S1 than L5B and L6 M1 neurons (Mao et al., 2011). Interestingly, S1-projecting neurons in L2/3 and L5A receive particularly strong input from S1 compared to other nearby pyramidal neurons (Mao et al., 2011). Together with our findings, this suggests that superficial S1-projecting neurons in M1 receive strong S1 input, while deep M1-projecting neurons in S1 receive strong M1 input (Figure 4.1A&B).

Whether these M1-S1 connections involve direct recurrent loops involving the same local neurons, or groups of neurons, is still unclear. For example, the upper layer M1 neurons that receive strong S1 input are neurons projecting back to S1 (Figure 4.1A), but what layers and neurons within S1 do they target? How do they interact with the L5/6 M1-projecting neurons in our study that receive strong M1 input (Figure 4.1B)? A number of circuit interactions are possible given the information we currently know. One possibility is that they could interact directly, forming a strong, direct recurrent loop (L2/3 and L5A M1 ↔ L5/6 S1; Figure 4.1C). A

second option is that they interact indirectly, through multiple laminar interactions within S1 and M1 (for example: L2/3 S1 → L2/3 M1 → L5/6 M1 → L5/6 S1). Finally, it is also unknown how M1-projecting neurons across different S1 layers interact with one another. Do L2/3, L5A, and L5/6 M1-projecting neurons preferentially synapse with one another, forming sub-networks in S1? Future studies that utilize more targeted expression of ChR2, either in specific layers or specific populations of projection neurons, are needed to answer each of these questions.

Based on known information, we suggest a circuit model wherein superficial layers of S1 and M1 are strongly connected from S1 to M1 and weakly connected in the reverse direction (Figure 4.1D). Such a pathway may serve to propagate sensory information from S1 to M1 (Ferezou et al., 2007). Conversely, the deep layers are strongly connected from M1 to S1 with a weaker connection in the other direction. In agreement with this idea, anatomical studies indicate that a dense projection exists from L5/6 of M1 to L5/6 of S1 (Zhang and Deschenes, 1998). Interestingly, M1 neurons in the deep layers that project to S1 are both callosal and corticofugal neurons (Veinante and Deschenes, 2003). The signal being relayed from M1 to S1 via M1 corticofugal neurons may be a copy of the motor command sent subcortically. However, an additional M1 to S1 circuit via M1 callosal neurons may also exist. The function of this circuit is less clear, but may carry to S1 information other than a copy of the subcortical motor command.

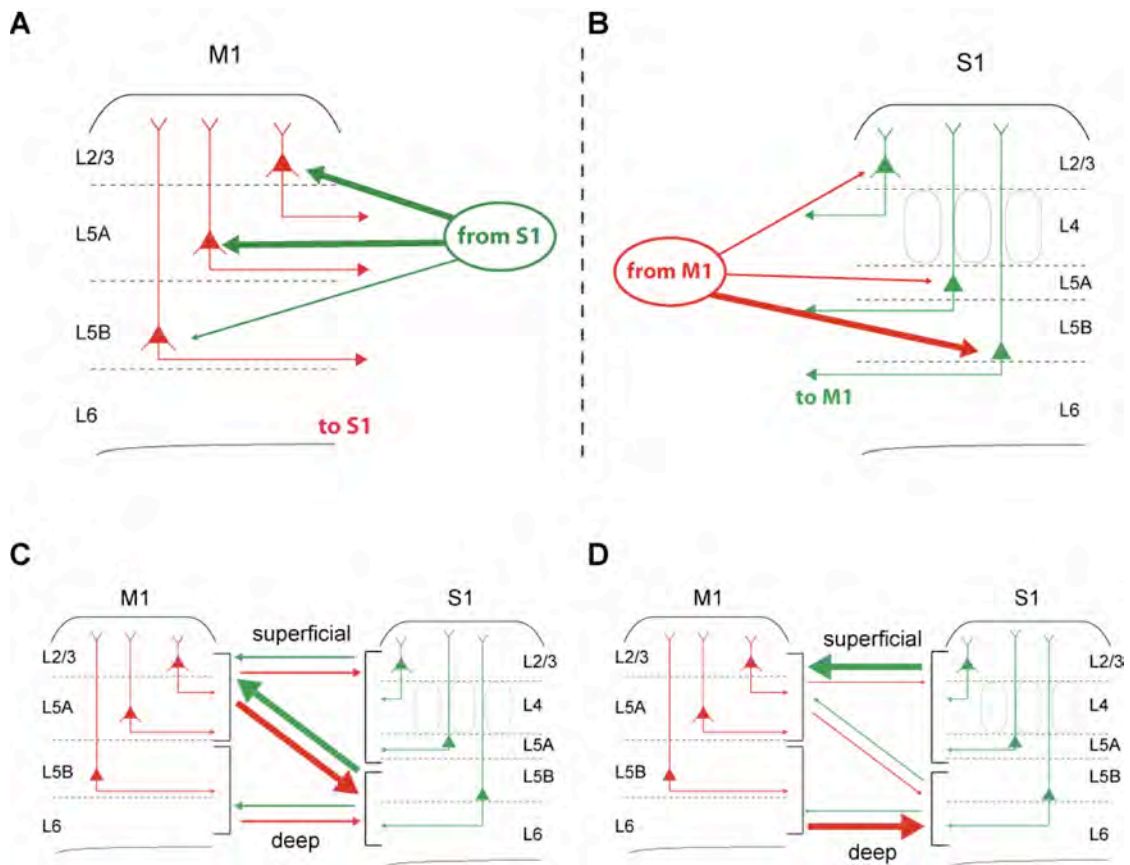


Figure 4.1 Schematic illustrating M1-S1 circuit interactions. (A) and (B) illustrate known information; (C) & (D) show possible models. (A) S1-projecting neurons in L2/3 and L5A of M1 receive strong S1 input. (B) L5/6 M1-projecting neurons in S1 receive strong M1 input. (C) Illustration of direct loops between superficial M1 and deep layer S1. (D) Proposed circuit model strongest connections are from superficial S1 to superficial M1, and deep M1 to deep S1 layers.

4.3.2 M1 influence on cortical feedback to VPM

VPM-projecting neurons *in vivo* are quiescent, exhibiting little or no spontaneous firing; neither do they reliably respond to whisker deflection, even when the animal is awake (Swadlow and Hicks, 1996). However, when activity in M1 is pharmacologically heightened, VPM-projecting neurons become more responsive to whisker deflections (Lee et al., 2008). Because in our study M1 input is weak and did not evoke action potentials in VPM-projecting neurons, our results, albeit in an *in vitro* preparation, argue against direct suprathreshold activation of VPM-projecting neurons by M1. However, M1 activity could excite VPM-projecting neurons indirectly in a number of different ways. For example, other S1 pyramidal neurons that are activated by M1, such as L5/6 M1-, Sp5-, or PoM-projecting neurons, may send a recurrent excitatory collateral onto VPM-projecting neurons, activating them via a disynaptic excitatory circuit within S1. However, corticothalamic neurons in L6 are relatively unconnected with cortical neurons in other layers and instead receive the majority of their excitatory input from other neurons in L6 (Zarrinpar and Callaway, 2006; Llano and Sherman, 2009).

Another possibility is that small input from M1 could combine with subthreshold input from another source, such as VPM, to evoke firing in VPM-projecting neurons. VPM-projecting neurons do receive input from VPM (White and Hersch, 1982), however this input alone also seems to be insufficient to evoke spiking activity since VPM-projecting neurons do not exhibit reliable suprathreshold responses to whisker deflections (see above). Therefore, perhaps VPM-projecting neurons act as ‘coincidence detectors’ by only firing when excitatory inputs from multiple sources are active. This would support the finding that VPM-projecting neurons become responsive to whisker deflection only when paired with activation of M1 (Lee et al., 2008). This idea would predict that VPM-projecting neurons might be more responsive during active touch,

when animals are actively moving their whisker while palpating an object thereby activating both the sensory and motor systems.

4.3.3 Cortical regulation of PoM and Sp5 circuits

An interesting difference between cortical feedback to PoM and VPM is the presence of L5B corticofugal neurons that send a collateral to PoM, but not VPM, en route to the brainstem. These corticothalamic synapses in PoM are very strong but also highly depressing (Reichova and Sherman, 2004; Groh et al., 2008). Therefore, the relatively high *in vivo* firing rates of PoM-projecting neurons and other subcortically projecting L5B neurons (de Kock et al., 2007) may cause these synapses to remain in a depressed, less effective state (Groh et al., 2008).

Intriguingly, if L5B PoM-projecting neurons transition between regular-spiking and intrinsically bursting dynamics *in vivo*, this could change the transfer of information at the L5B-PoM corticothalamic synapses in two ways, similar to the effects of thalamic bursting on cortical neurons (Swadlow and Gusev, 2001; Llano and Sherman, 2009). First, the high frequency spiking *within* a burst could increase the probability of information propagating through a depressed synapse; secondly, the relatively long time interval *between* bursts could allow for recovery of the synapse from synaptic depression (Llano and Sherman, 2009). Combination of these two features would allow for increased communication from S1 to PoM during states when PoM-projecting neurons are in burst mode. Whether the S1 to Sp5 synapse shows similar synaptic properties is unknown. If so, a similar model could regulate S1 input to Sp5 as well. Finally, if M1 input to PoM- and Sp5-projecting neurons changes the incidence of burst firing,

via its input to the apical tuft in L1, it could regulate the information transfer between S1 and its subcortical projections.

4.4 GENERAL IMPLICATIONS

Some models of the cerebral cortex assume that cortical connectivity is random and indiscriminate; that is, a given axon will synapse with equal probability on all neurons within its target zone. In this case, the actual number of synapses on a given neuron simply reflects, statistically, the number of contacting terminals and the number of available post-synaptic sites. Because pyramidal cell morphology is relatively uniform, at least when compared to that of interneurons, such models often consider pyramidal neurons, especially neighboring ones, as a single population. However, accumulating evidence demonstrates that pyramidal neurons differ with respect to their intrinsic properties and their local and extrinsic projections. Based on these properties, recent studies have found that excitatory connections among local S1 pyramidal neurons are not random, but instead can form subnetworks based on either similar physiological properties (Otsuka and Kawaguchi, 2008), number of shared pre-synaptic neurons (Perin et al., 2011), or the projection target of the neurons (Brown and Hestrin, 2009; Otsuka and Kawaguchi, 2011). Incorporating non-uniform connections between excitatory neurons can greatly alter the dynamics and properties of a cortical network (Litwin-Kumar and Doiron, 2012).

The findings of the studies in this thesis argue strongly against an indiscriminate connection model regarding long-range corticocortical connections. Indeed, present findings indicate a high degree of specificity not only in the connections from M1 to excitatory versus

inhibitory S1 neurons but also to different sub-classes of interneurons and of pyramidal cells. In the case of the latter, projection target, and hence effects on other brain regions, may be the most important factor. Our studies do not distinguish whether the larger inputs onto some classes of S1 pyramidal neurons are due to larger unitary inputs and/or the convergence of a larger number of M1 neurons onto a single post-synaptic S1 neuron, although both have interesting implications.

Our findings of M1 to S1 specificity are entirely consistent with rules of cortical circuitry originally formalized by White (1989) based on his careful light and electron microscope studies of VPM input to L4 neurons and to pyramidal neurons projecting to M1, striatum, and the thalamus. First, axon terminals from any extrinsic source will form synaptic contacts with every neuronal type within its axonal projection field that is capable of receiving that type of synapse. Second, neuron types will receive characteristic patterns of synaptic input regarding the actual numbers and proportions of synapses. Regarding our findings, a functional counterpart of these rules is that inputs from M1 may generally modulate S1 circuitry by synaptically contacting multiple cell types at different depths within a cortical column, but such inputs will also engage to different degrees specific cell populations based on their extrinsic, and possibly local, projections.

APPENDIX A

POSTNATAL MATURATION OF SOMATOSTATIN-EXPRESSING INHIBITORY CELLS IN THE SOMATOSENSORY CORTEX OF GIN MICE

A.1 ABSTRACT

Postnatal inhibitory neuron development affects mammalian brain function, and failure of this maturation process may underlie pathological conditions such as epilepsy, schizophrenia and depression. Furthermore, understanding how physiological properties of inhibitory neurons change throughout development is critical to understanding the role(s) these cells play in cortical processing. One subset of inhibitory neurons that may be affected during postnatal development is somatostatin-expressing cells. A subset of these cells is labeled with green-fluorescent protein (GFP) in a line of mice known as the GIN line. Here, we studied how intrinsic electrophysiological properties of these cells changed in the somatosensory cortex of GIN mice between postnatal ages P11 to P32+. GIN cells were targeted for whole-cell current clamp recordings and ranges of positive and negative current steps were presented to each cell. The results showed that as the neocortical circuitry matured during this critical time period, multiple intrinsic and firing properties of GIN inhibitory neurons, as well as those of excitatory (regular-spiking [RS]) cells, were altered. Furthermore, these changes were such that the output of GIN

cells, but not RS cells, increased over this developmental period. We quantified changes in excitability by examining the input-output relationship of both GIN and RS cells. We found that the firing frequency of GIN cells increased with age, while the rheobase current remained constant across development. This created a multiplicative increase in the input-output relationship of the GIN cells, leading to increases in gain with age. The input-output relationship of the RS cells, on the other hand, showed primarily an additive shift with age, but no substantial change in gain. These results suggest that as the neocortex matures, inhibition coming from GIN cells may become more influential in the circuit and play a greater role in the modulation of neocortical activity.

A.2 INTRODUCTION

The second and third postnatal weeks are a period of massive developmental change within the rodent neocortex. This is the time of sensory onset for multiple sensory modalities, including the visual, auditory, and somatosensory systems (Welker, 1964, Ehret, 1976, Hensch, 2005). This time window represents a ‘critical period’ for the brain, as the organization of neural circuitry that develops within this time remains relatively fixed throughout adulthood. Diseases such as schizophrenia, which do not appear until late adolescence in humans, are thought to be a manifestation of pathological alterations in developmental processes, potentially in inhibitory neurons (Lewis et al., 2005). Understanding the development of cortical circuits during this period is therefore an important component to understanding both normal and pathological cortical function.

Maturation of inhibitory circuitry is of particular importance for shaping cortical activity during postnatal development (Hensch, 2005). Investigation of inhibitory interneuron development in multiple cortical areas reveals that during the second to third postnatal weeks, inhibitory cells undergo changes in their physiological, molecular, and anatomical characteristics and do not reach maturation until the third week of postnatal development or later (Goldberg et al., 2011, Lazarus and Huang, 2011, Oswald and Reyes, 2011). In mice and rats, the animals first begin actively using their whiskers around postnatal day 14, signaling the onset of active sensory processing within the somatosensory system (Welker, 1964, Mosconi et al., 2010). During the first few weeks of postnatal development, inhibition onto pyramidal neurons in somatosensory cortex increases in amplitude and frequency (Kobayashi et al., 2008), and alterations in sensory experience, such as whisker removal, cause substantial disruptions to the intrinsic and synaptic properties of inhibitory interneurons (Jiao et al., 2006, Daw et al., 2007, Lee et al., 2007, Chittajallu and Isaac, 2010, Jiao et al., 2011).

Understanding maturation of inhibitory circuits is important for determining how neurons control synaptic integration and communication within and amongst cortical areas during development. Most studies investigating how development of inhibitory neurons affects cortical circuitry have focused on one type of inhibitory neuron, known as the fast-spiking (FS) or parvalbumin-expressing (PV) cells. These cells represent a major subtype of inhibitory neurons, but only constitute approximately half of inhibitory neurons as a whole (Markram et al., 2004). A second major class of inhibitory neurons is the somatostatin-expressing (SOM) inhibitory cells. However, the developmental trajectory of SOM cells in the somatosensory system is largely unknown. One recent study found that the intrinsic properties of SOM cells in the visual

cortex change during the second postnatal week and suggested SOM cells may be involved in ocular dominance plasticity (Lazarus and Huang, 2011).

The role of SOM inhibitory cells in the neocortex has yet to be fully understood. Because the properties of these cells differ significantly from other inhibitory cells, such as PV cells, it is thought that they could play a unique role in neocortical processing. In contrast to PV cells, which target the perisomatic regions of their downstream cells, SOM cells primarily target the dendrites of pyramidal neurons. They may therefore be important for regulating and modulating incoming synaptic inputs. Recent *in vivo* recordings from superficial SOM cells in mouse somatosensory cortex have found that activity in SOM cells is suppressed during both passive whisker deflection and during active whisking states (Gentet et al., 2012). This loss of dendritic inhibition may function to allow excitatory inputs on the distal dendrites to summate and propagate to the soma more effectively. This may be particularly relevant during awake states when activity in the dendrites is enhanced (Murayama and Larkum, 2009). Activation of SOM cells has also been shown to effectively prevent pyramidal neurons from producing ‘bursts’ of action potentials that are generated in the apical dendrite through active dendritic currents (Larkum et al., 1999, Gentet et al., 2012, Lovett-Barron et al., 2012). In addition, SOM cells are extensively coupled to each other through gap junctions (Gibson et al., 1999, Fanselow et al., 2008). Alterations in intrinsic properties with age could change how SOM cells act as an electrically coupled network, such as by changing their tendency to correlate or synchronize their activity (Amitai et al., 2002, Long et al., 2005, de la Rocha et al., 2007).

Across the SOM interneuron population there exists a degree of variability in the morphology, physiological characteristics, and co-expression of additional neuropeptides. Whether this reflects variability within a single class or is sufficiently diverse to warrant multiple

subtypes is unclear (Ma et al., 2006, Sugino et al., 2006, McGarry et al., 2010). This issue has been partially addressed by the creation of transgenic mouse lines that express a fluorescent molecule, such as green fluorescent protein (GFP), in particular subsets of inhibitory neurons. Here, we used such a mouse line to study SOM inhibitory cells by utilizing a line of mice that expresses GFP in approximately one-third of SOM cells (Oliva et al., 2000). In these mice, the GFP-positive inhibitory neurons (“GIN”) express the neuropeptide, somatostatin, exhibit adapting responses to intracellular current steps, and are often of Martinotti morphology, with an axon traveling up to layer 1 and ramifying extensively (Oliva et al., 2000, Halabisky et al., 2006, Ma et al., 2006, Fanselow et al., 2008).

Understanding the normal trajectory of GIN cell maturation serves several purposes. First, it helps clarify the physiological role(s) these cells can play during different stages of postnatal development by indicating how intrinsic properties of GIN cells change over age, how readily these cells are excited and what their firing characteristics are once activated. Second, it will help us learn how to best distinguish between types and subtypes of inhibitory neurons even as their physiological characteristics, which are often used as factors to distinguish between cell types (Kawaguchi, 1995, Kawaguchi and Kubota, 1996, Kawaguchi and Kondo, 2002, Markram et al., 2004, Ascoli et al., 2008), change systematically with age. Finally, the way circuitry of excitatory and inhibitory neurons changes at different stages of maturation may help us determine causes and mechanisms of abnormal cortical development (Dani et al., 2005, Lewis et al., 2005, Yizhar et al., 2011).

Here, we characterized the intrinsic physiological properties of GIN cells and regular-spiking (RS) excitatory cells in the mouse primary somatosensory cortex during a developmental period (P11-P32+) that encompasses changes in multiple intrinsic cellular properties in

somatosensory cortex (Maravall et al., 2004, Jiao et al., 2006, Goldberg et al., 2011, Jiao et al., 2011) as well as whisking onset. We found that changes in sub- and suprathreshold properties occurred in both cell types, collectively resulting in an increase in GIN cell excitability and a reduction in RS cell excitability as a function of age. These results suggest that as the neocortical circuit develops, GIN cells may increasingly participate in sensory processing within somatosensory cortex.

A.3 MATERIALS AND METHODS

Slicing procedure

All experiments were carried out in compliance with the University of Pittsburgh School of Medicine animal use policies and were approved by the University of Pittsburgh Institutional Animal Care and Use Committee. Thalamocortical slices from mouse somatosensory cortex were prepared as previously described (Agmon and Connors, 1991). In addition, a subset of cells was recorded from coronal sections of the same brain region. No systematic differences between these two slicing angles were observed, so the data were pooled.

Recording procedures

Whole cell current-clamp recordings were performed using micropipettes (4–7 M Ω) filled with internal solution containing (in mM): 135 K-gluconate, 4 KCl, 2 NaCl, 10 HEPES, 0.2 EGTA, 4 ATP-Mg, 0.3 GTP-Tris, and 14 phosphocreatine-Tris (pH 7.25, 280-290 mOsm). Membrane potentials reported here were not corrected for the liquid junction potential.

Recordings were conducted at 32°C. When patching, cell-attached seal resistances were 1 GΩ or greater and series resistance after achieving whole cell configuration was 5-20 MΩ. Data were collected using a Multiclamp 700B amplifier and pClamp10 software (Molecular Devices). Data were analyzed using in-house programs written in Matlab (The MathWorks, Natick, MA; E. Fanselow).

Cell visualization and identification

Cells were viewed under infrared-differential interference contrast illumination using a Nikon FN-1 microscope and a Dage IR-1000 CCD camera. GIN cells were identified by visualization of GFP under epifluorescence illumination. Regular-spiking (RS) pyramidal cells did not express GFP in these mice and were therefore targeted for recording based on having a triangular-shaped soma. During recording, RS pyramidal cells were positively distinguished from regular-spiking inhibitory interneurons as having strongly adapting spike firing rates during suprathreshold current steps, small amplitude AHPs, and during a train of action potentials the first AHP was smaller than subsequent AHPs (Porter et al., 2001; Beierlein et al., 2003). All cells were recorded in layers 2 and 3 of somatosensory cortex. Layer 2/3 cells were defined as those between the bottom of layer 1, identified by its lack of cell bodies, and the tops of the barrels, which demarcate layer 4.

Cells recorded

Recordings from a total of 269 GIN cells and 200 RS cells from a total of 170 mice were analyzed for this study. Numbers of cells for each cell type and age can be found in Table 1. All mice were of the GIN strain (Jackson Labs, Bar Harbor, ME; (Oliva et al., 2000)) and were aged

P11-50. Cells from those aged > P31 were combined and are identified here as P32+. In addition, we combined P18 and P19 cells in to one group, identified as P19 in the text and on our plots. Both male and female mice were used. The morphologies of GIN cells have been described extensively in other publications, and these cells have been shown to express the peptide, somatostatin (SOM; (Oliva et al., 2000; Halabisky et al., 2006; Ma et al., 2006; Fanselow et al., 2008)).

Measures of cellular properties

Once patched, we presented each cell with a series of negative and positive current steps, ranging from -100 pA to 300 pA in 10 or 20 pA steps. The following measures were made from the voltage traces recorded during these current steps.

Adaptation ratio: inverse of mean of the last three interspike intervals during a 200 pA step, divided by the inverse of the initial interspike interval in that step.

Action potential threshold: voltage at the time of the peak of the 3rd derivative of the voltage trace for the first action potential in a given current step.

After-hyperpolarization (AHP) magnitude (mV): voltage at the most negative trough of the AHP (within 4 ms of the action potential peak) minus the action potential threshold voltage.

AHP slope (mV/ms): positive slope of the AHP for 8 ms following the most negative trough in the AHP.

Firing frequency, initial (F_i ; Hz): inverse of the first interspike interval during a 200 pA current step.

Firing frequency, steady state (F_{ss} ; Hz): average of the inverse of the last 3 interspike intervals in a 200 pA current step.

Half-width (ms): width of the action potential halfway between the threshold voltage and the peak voltage.

Input-output gain (F-I slope; Hz/pA): slope of the linear portion of each frequency-current (F-I) curve calculated from responses to suprathreshold current steps. Frequency was defined as the inverse of the interspike intervals throughout the entire current step.

Input resistance (M Ω): magnitude of voltage deflection (mV)/magnitude of current step (pA); calculated as the slope of the V/I plot for positive and negative voltage deflections between ± 8 mV.

Membrane time constant (τ_m ; ms): calculated from a monoexponential curve best fitting the falling phase of the voltage deflection in response to negative current steps. Only voltage deflections < 8 mV negative to the resting membrane potential were used.

Rebound depolarization (mV): The height of the voltage peak within 200 ms of the end of a current step. Current steps with voltage deflections from -70 to -15 mV were used.

Resting membrane potential (V_{rest} ; mV): average voltage in the 20 ms prior to a current step.

Rheobase current (pA): We plotted the number of action potentials as a function of input current and extrapolated to find the current required to elicit a single action potential

Sag magnitude (mV): calculated from negative-going current steps as the difference between the peak negative voltage deflection and the average voltage for the last 50 ms of the 600 ms current step. Only traces in which the maximum voltage deflection was >15 mV more negative than the resting membrane potential were analyzed.

Threshold voltage (mV): calculated as the voltage at the time of the maximum of the 3rd derivative of the action potential voltage trace.

Statistical analyses

For quantification of RS and GIN cell properties, measurements were made from negative or positive current steps as described above. When results were plotted as a function of age, the resulting data were fit with linear, exponential, 2nd degree polynomial or sigmoidal functions, as appropriate, using SigmaPlot (San Jose, CA). P11 and P32+ values were compared for each type of measurement using a 2-tailed t-test and results are presented in Table 2.

A.4 RESULTS

In this study, we quantified intrinsic physiological properties of GIN and regular-spiking (RS) pyramidal cells in layer 2/3 of somatosensory cortex ranging in age from P11 to P32+. We recorded from local pyramidal neurons (RS cells) in order to directly compare how GIN and RS cells develop relative to one another within the same network. The development of intrinsic properties of pyramidal cells has been reported previously in multiple brain regions (McCormick and Prince, 1987; Maravall et al., 2004; Frick et al., 2007; Oswald and Reyes, 2008), and the results we present here for RS cells are largely in agreement with these previous studies. In contrast, the changes in GIN properties we observed have not been reported for the somatosensory cortex and differed somewhat from those described for visual cortex (Lazarus and Huang, 2011).

Subthreshold intrinsic properties

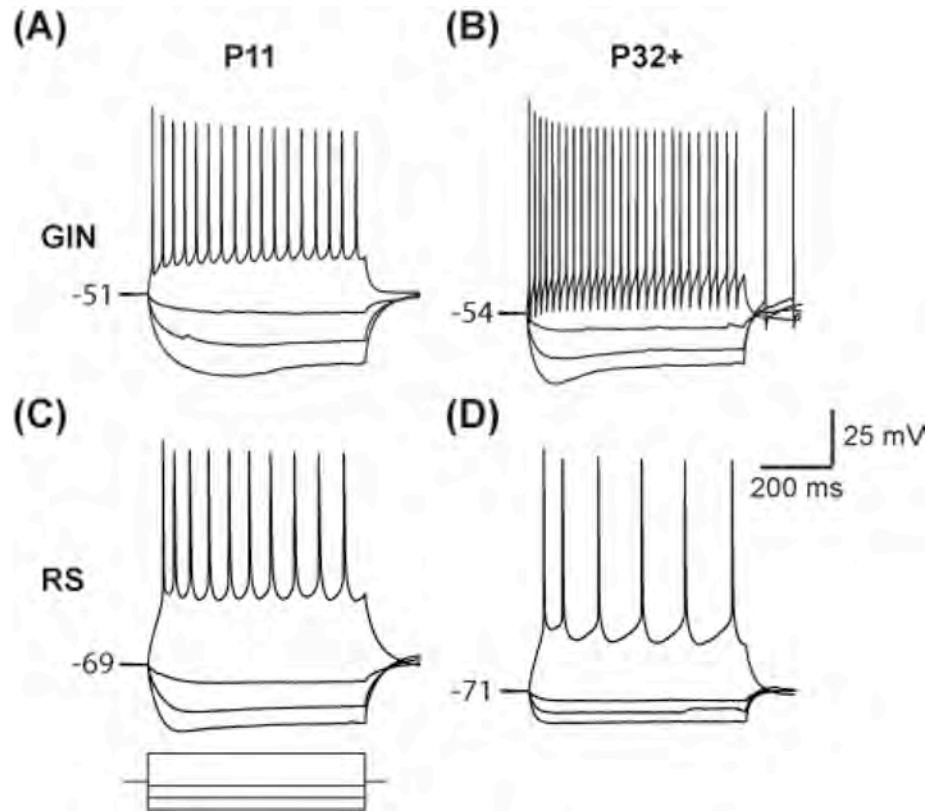


Figure A. 1 Responses of GIN and RS cells to current steps. (A) Voltage responses to current steps injected into a P11 GIN cell. (B) Voltage responses to current steps injected into a P32+ GIN cell. (C) Top: Voltage responses to current steps injected into a P11 RS cell. Bottom: Current steps representing -100, -60, -20, and +100pA current steps, each of 600ms duration. The same amplitude current steps were used for A-D. (D) Voltage responses to current steps injected into a P32+ RS cell.

We observed that multiple subthreshold characteristics of GIN cells develop over the time window examined in this study, P11-32+ (Figure A.1). First, we examined several intrinsic properties of these cells: the input resistance (R_{in}), membrane time constant (τ_m), rheobase current, resting membrane potential (V_{rest}), and the threshold voltage for action potential generation (V_{thresh}). We found that several, but not all, of these features changed over this age range (Figure A.2). The R_{in} of GIN cells decreased significantly from P11 to P32+ (P11: 379 ± 32 M Ω ; P32+: 244 ± 12.1 M Ω , $p < 0.001$). This change occurred rapidly in GIN cells, as R_{in} decreased mainly between P11 (379 ± 32 M Ω) and P12 (247 ± 22 M Ω) then remained steady until P32+ (Figure A.2A). The τ_m in GIN cells also decreased from P11 to P12 then remained steady until P32+ (P11: 56 ± 4.6 ms; P12: 32 ± 4 ms; P32+: 23.2 ± 1.7 ms, $p < 0.001$; Figure A.2B). The rheobase current in GIN cells increased from P11 to P12 and then remained constant (P11: 14 ± 2 pA; P12: 37 ± 7 pA, P32+: 28 ± 5 pA, $p < 0.05$; Figure A.2C).

In RS cells, the R_{in} and τ_m also decreased during this period, but did so more gradually than in GIN cells (R_{in} : P11, 326 ± 35 M Ω ; P32+, 160 ± 9.7 M Ω , $p < 0.001$; τ_m : P11, 32 ± 4.9 ms; P32+, 14 ± 1.9 , $p = 0.009$; Figure A.2A&B). The rheobase current in RS cells increased threefold, from 30 ± 4 pA at P11 to 90 ± 21 pA at P32+ (Figure A.2C). This increase in rheobase current also occurred more gradually over the second postnatal week than in GIN cells. V_{rest} did not change significantly throughout development in GIN cells (P11: -57.2 ± 1 mV; P32+: -59.2 ± 0.8 mV, $p = 0.135$; Table 2) or RS cells (P11: -64.9 ± 1.3 mV; P32+: -64.9 ± 2.1 mV; $p = 0.978$; Table 2). In addition, V_{thresh} remained constant across the age range investigated here in both GIN cells (P11: -40 ± 0.5 mV; P32+: -41 ± 0.6 mV, $p = 0.511$; Table 2) and RS cells (P11: -40 ± 0.9 mV; P32+: -40 ± 1.3 , $p = 0.633$; Table 2).

Table 2. Summary of cell counts

Cell Type	Postnatal Age (days)									Totals
	P11	P12	P13	P14	P15	P16	P17	P18-19	P32+	
GIN	16	14	41	66	48	46	10	6	22	269
RS	11	4	42	37	39	20	22	12	13	200

Collectively, these results suggest that during the age range investigated, the intrinsic excitability of both GIN and RS cells decreased. But it should be noted that within the same age range, GIN cells remained more excitable than RS cells, exhibiting higher input resistance, higher τ_m , lower rheobase current, and more depolarized V_{rest} than RS cells across this developmental period.

Another characteristic feature of GIN cells is the presence of a pronounced ‘sag current’ in response to hyperpolarizing current steps (Oliva et al., 2000; Halabisky et al., 2006; Ma et al., 2006) Figure A.1B & A.3A). In GIN cells, sag currents are mediated by I_h currents (Ma et al., 2006). We observed a strong increase in the magnitude of the sag current with age (P11: 3 ± 0.6 mV; P32+: 5.6 ± 0.5 mV, $p < 0.01$; Figure A.3A&B). A related feature observed in GIN cells is the presence of a rebound depolarization when the cell is released from hyperpolarization, which in some cases results in the generation of a rebound spike (Figure A.1B & A.3A). This depolarization is also mediated by I_h currents in GIN cells (Ma et al., 2006). We observed that the amplitude of rebound depolarization also increased with age (P11: 2 ± 0.8 mV; P32+: 4.4 ± 0.4 , $p < 0.05$; Figure A.3C), and showed a similar developmental time course as the change in sag current magnitude. RS cells also showed a small I_h -mediated sag current (Maravall et al., 2004), but the magnitude of the sag current and rebound depolarization decreased with age in RS cells (Figure A.3A-C), consistent with previous results (Maravall et al., 2004). Together, these findings suggest that I_h currents increased in GIN cells and decreased in RS cells from ages P11 to P32+.

Table 3. Cell properties at P11 and P32+

Property	GIN			RS		
	P11	P32+	p-value	P11	P32+	p-value
Adaptation Ratio	0.6 ± 0.03	0.5 ± 0.03	*	0.4 ± 0.03	0.4 ± 0.06	0.162
AHP magnitude (mV)	-9.6 ± 0.5	-14.7 ± 0.7	***	-6.4 ± -1.1	-11.8 ± -1.5	*
AHP slope (mV/ms)	0.7 ± 0.1	1.9 ± 0.15	***	0.4 ± 0.03	0.8 ± 0.2	0.189
Action potential threshold, 1st sp ke (mV)	-40 ± 0.5	-41 ± 0.6	0.511	-40 ± 0.9	-40 ± 1.3	0.633
Firing frequency, initial (Hz)	61 ± 2	96 ± 2.6	***	59 ± 2.6	60 ± 5.1	0.856
Firing frequency, steady state (Hz)	38 ± 3	42.4 ± 3.2	**	20 ± 1	25 ± 1.9	0.11
Gain (Hz/pA)	0.2 ± 0.02	0.44 ± 0.02	***	0.14 ± 0.02	0.28 ± 0.05	*
Half-width (ms)	1.4 ± 0.05	0.8 ± 0.05	***	1.8 ± 0.08	0.96 ± 0.09	***
Input-output gain (Hz/pA)	0.2 ± 0.02	0.4 ± 0.02	***	0.14 ± 0.02	0.28 ± 0.05	0.037
Input resistance (MΩ)	379 ± 32	244 ± 12.1	***	326 ± 35	160 ± 9.7	***
Membrane time constant (ms)	56 ± 4.6	23.2 ± 1.7	***	32 ± 4.9	14 ± 1.9	0.009
Rebound depolarization (mV)	2 ± 0.8	4.4 ± 0.4	*	3.1 ± 0.7	1.4 ± 0.4	0.033
Rheobase current (pA)	14 ± 2	28 ± 5	*	30 ± 4	90 ± 21	*
Sag magnitude (mV)	3 ± 0.6	5.6 ± 0.5	**	2.7 ± 0.6	1.02 ± 0.3	*
V _{rest} (mV)	-57.2 ± 1	-59.2 ± 0.8	0.135	-64.9 ± 1.3	-64.9 ± 2.1	0.978

*p<0.05, **p<0.01, ***p<0.001

Action potential properties

GIN cells exhibit a distinct action potential and afterhyperpolarization (AHP) waveform, which is likely due to expression of several types of potassium currents (Zhang and McBain, 1995; Rudy and McBain, 2001; Halabisky et al., 2006; Ma et al., 2006). We therefore sought to determine whether action potential properties of GIN cells changed with age, which would suggest differential expression of these currents across development. We measured three properties related to the shape of the action potential: action potential half-width, the AHP magnitude, and the slope of the voltage depolarization following the AHP. The average action potential half-width in GIN cells decreased with age, from 1.4 ± 0.05 ms at P11 to 0.8 ± 0.05 ms at P32+ ($p<0.001$; Figure A.4A). Narrowing of the action potential with age was also observed in RS cells

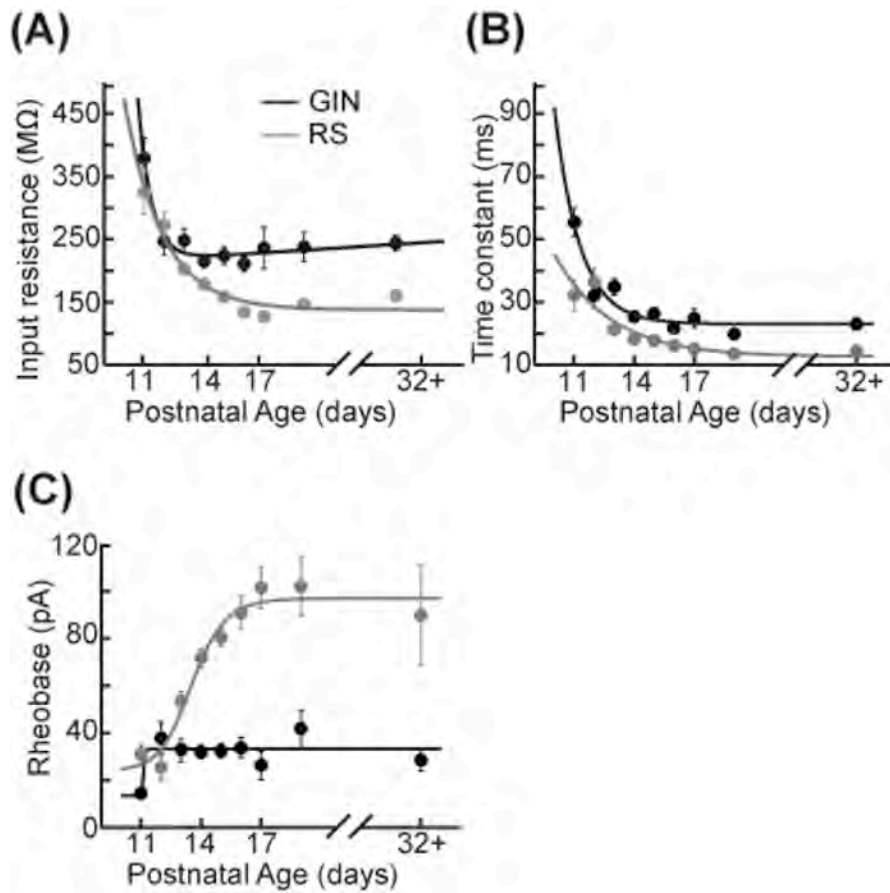


Figure A. 2 Intrinsic properties of GIN cells and RS cells change over development. (A) Average input resistance of GIN cells (black) and RS cells (gray) decreases as a function of age. (B) The average membrane time constant also decreases with age in both the GIN cells (black) and the RS cells (gray). (C) The rheobase current increases from P11 to P12 in the GIN cells, then remains constant into maturity (black). The rheobase current in the RS cells increases significantly with age (gray). Data for each cell type was fit with an exponential function (A&B) or a sigmoidal function (C).

(P11: 1.8 ± 0.08 ms; P32+: 0.96 ± 0.09 , $p < 0.001$; Figure A.4A), but at all ages recorded the half-width of GIN cells was shorter than for RS cells of the same age.

GIN cells can display a complex 'triphasic' AHP shape following an action potential that consists of both a rapid 'early' component, followed by a slower 'late' component (Halabisky et al., 2006; Ma et al., 2006). We found that both of these features were almost completely absent at P11 in the GIN cells but developed throughout the second and third postnatal weeks (Figure A.1A&B; Figure A.3B). Specifically, the trough of the AHP became increasingly negative with age (P11: -9.6 ± 0.5 mV; P32+: -14.7 ± 0.7 mV, $p < 0.001$; Figure A.4B&C). We then quantified the repolarization of the voltage following the AHP and found that the slope increased almost threefold from P11 (0.7 ± 0.1 mV/ms) to P32+ (1.9 ± 0.15 , $p < 0.001$; Figure A.4B&C). Therefore, in GIN cells, although the AHP became increasingly negative with age, the voltage repolarized more quickly so that by ~ 8 ms following the AHP the voltage was more depolarized in mature animals than younger animals (Figure A.4B). In contrast, RS cells had a smaller AHP magnitude across all ages than GIN cells (Figure A.4C). We did observe a significant increase in the AHP magnitude with age in RS cells (P11: -6.4 ± -1.1 ; P32+ -11.8 ± -1.5 , $p < 0.05$), but no significant change in AHP slope for RS cells (P11: 0.4 ± 0.03 mV/ms; P32+: 0.8 ± 0.2 ; $p = 0.189$).

Firing properties

Finally, we examined how the propensity of GIN cells to fire in response to a current step changed as a function of age. We started by quantifying the firing of the GIN cells throughout the duration of suprathreshold current steps across each age (Figure A.5A). GIN cells across all ages exhibited adaptation throughout the current steps, such that the initial firing rate was higher

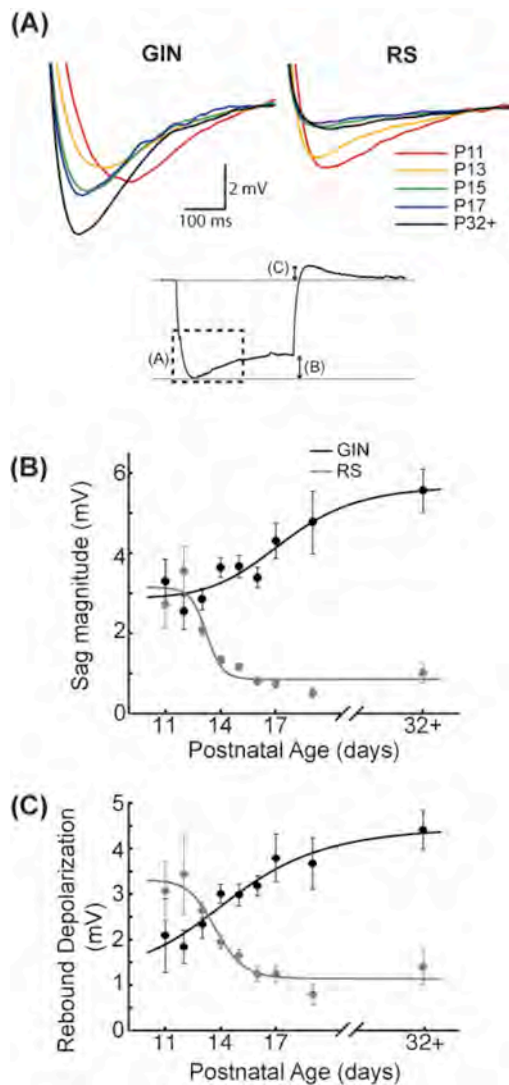


Figure A. 3 Sag currents increase in GIN cells with age, but decrease in RS cells. (A) Top: Average sag current traces for GIN cell traces (left) and RS cell (right) by age. The colors correspond to the following ages: P11 (red), P13 (orange), P15 (green), P17 (blue) and P32+ (black). Bottom: Response of a GIN cell to a hyperpolarizing current step. The dotted box denotes the region illustrated above in (A). The arrows describe the sag magnitude (shown in (B)) and the rebound depolarization (shown in (C)). For descriptions of how these quantities were measured, see METHODS. (B) The magnitude of the sag current in GIN cells increases with age (black), whereas the sag current decreases with age in the RS cells (gray). (C) The rebound depolarization also increases in GIN cells

(black) and decreases in RS cells, following a similar time course as the sag current (gray). Data for each cell type was fit with a sigmoidal function (B&C).

than the steady state firing rate (Figure A.5A&C). Both the initial (P11: 61 ± 2 Hz; P32+: 96 ± 2.6 Hz, $p < 0.001$) and the steady state firing rate (P11: 38 ± 3 Hz; P32+: 42.4 ± 3.2 Hz, $p < 0.01$) increased significantly with age. The ratio of the two values, the adaptation ratio, decreased across development (P11: 0.6 ± 0.03 ; P32+: 0.5 ± 0.03 , $p < 0.05$; Figure A.3C). This is likely due to the fact that the initial firing rate increased more with age than the steady state firing rate, causing GIN cells to exhibit more adaptation in older animals. In comparison, we observed no significant change in the initial firing rate, steady state firing rate, or adaptation ratio across ages in RS cells (Table 2; Figure A.5B&C).

In order to fully understand how the firing dynamics of GIN cells and RS cells changed during development, we examined the input-output relationship of both cell types from P11 to P32+. When we compared the input-output relationship across age, GIN cells showed a 'multiplicative' increase in firing rate from P11 to P32+ (Figure A.5D). This was primarily because whereas the firing rates increased with age, the rheobase current stayed constant from P12 through P32+ (Figure A.2C; Figure A.5D). To quantify this change, we calculated the slope of the linear portion of the input-output relationship, or the "gain" of the cell. The gain of the GIN cells increased significantly across development (P11: 0.2 ± 0.02 Hz/pA; P32+: 0.44 ± 0.02 Hz/pA, $p < 0.001$; Figure A.5F). RS cells also showed a significant increase in gain from P11 (0.14 ± 0.02 Hz/pA) to P32+ (0.28 ± 0.05 Hz/pA, $p < 0.05$; Figure A.5F). However, the dominant change in the input-output curve of RS cells was a rightward shift with age (Figure A.5E). This was due to the increase

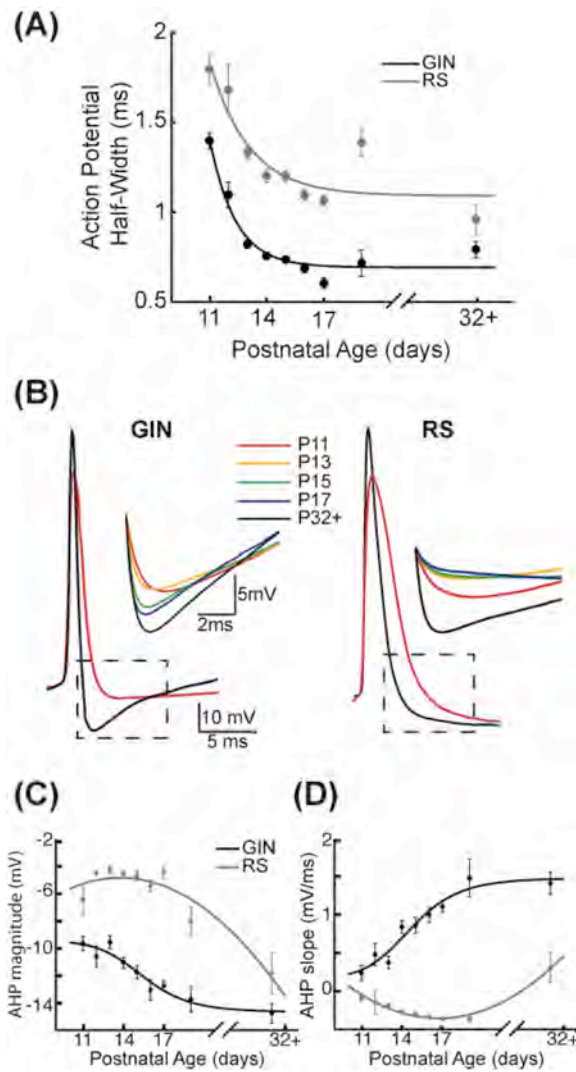


Figure A. 4 Action potential properties change as a function of age in GIN and RS cells. (A) Across development, action potential half-width decreases in GIN (black) and RS cells (gray). (B) Left: Overlapped action potential waveforms of a P11 GIN cell (red) and a P32+ GIN cell (black). Dashed black box indicates region shown above, which illustrates the change in the afterhyperpolarization (AHP) with age in GIN cells. Right: Overlapped action potential waveforms of a P11 RS cell (red) and a P32+ RS cell (black). Dashed black box indicates region shown above, which illustrates the change in the afterhyperpolarization (AHP) with age in RS cells. The colors correspond to the following ages: P11 (red), P13 (orange), P15 (green), P17 (blue) and P32+ (black). (C) Quantification of AHP magnitude (measured as the most negative point of the AHP) for GIN cells (black) and RS cells (gray). (D) Quantification of the slope of the voltage for 8ms following the AHP for GIN cells (black) and RS

cells (gray). Data for each cell type was fit with a sigmoidal function (C&D; black) or a 2nd degree polynomial (C&D; gray).

in rheobase current with age in RS cells (Figure A.2C), resulting in more current being required to drive RS cells to fire at a given rate as the neurons matured.

Discussion

In this study, we sought to determine whether physiological features of GIN inhibitory cells changed as a function of age. Our data demonstrated that multiple intrinsic physiological characteristics of L2/3 GIN cells and RS cells in somatosensory cortex changed across the developmental time period investigated here (P11-P32+). GIN and RS cells showed decreased intrinsic excitability over the second and third weeks of postnatal development, based on features such as input resistance, membrane time constant and rheobase current. However, despite these changes, the net change in GIN cell output during this time was an increase in firing rate as a function of age. This was due to an increase in initial and steady state firing rates with age, which resulted in a multiplicative increase in the input-output relationship. In contrast, RS cells did not show substantial changes in firing rate with age. Together, these results suggest that, for the same level of input current, GIN cells became more active as the animal matured through the age range investigated, whereas RS cells became less readily activated.

The results we reported here for GIN cells are consistent with developmental changes to intrinsic properties reported for pyramidal neurons in the current study and elsewhere (Maravall et al., 2004) and for other types of inhibitory interneurons (Goldberg et al., 2011) in the upper layers of

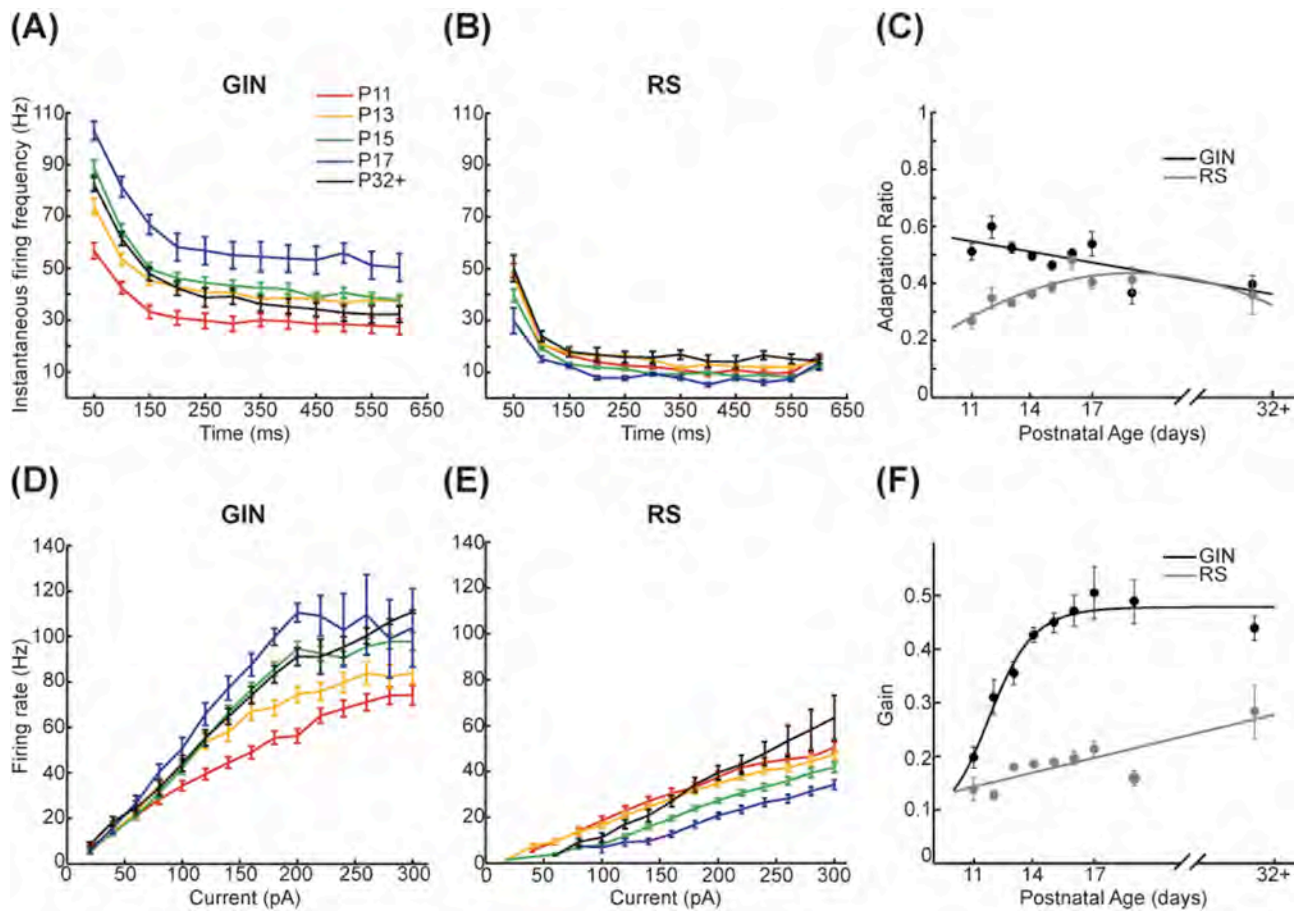


Figure A. 5 GIN cells show increased firing rates with age. (A) Instantaneous firing rate across a 200pA current step of 600ms duration for GIN cells. (B) Instantaneous firing rate across a 200pA current step of 600ms duration for RS cells. (C) The adaptation ratio (1^{st} ISI/average of last 3 ISIs) decreased slightly with age in GIN cells (black) but did not change in RS cells (gray). (D) The frequency/current (f/I) curve of GIN cells calculated from 20 to 300 pA across age showed a multiplicative increase. (E) The f/I curve of RS cells calculated from 20 to 300pA across age shows primarily a rightward shift. (F) The gain (calculated as the slope of the linear portion of the f/I curve) for GIN cells (black) increased with age, but not in RS cells (gray). The colors in A, B, D, and E correspond to the following ages: P11 (red), P13 (orange), P15 (green), P17 (blue) and P32+ (black). Data for each cell type was fit by a linear fit (C, black; F, gray), a 2^{nd} degree polynomial (C, gray), or a sigmoidal function (F, black).

the somatosensory cortex. Therefore, this may be a general developmental trend across many cell types within this neocortical region. It should be noted that a recent study investigated maturation of SOM and PV cells in the visual cortex during the first few weeks of postnatal development (Lazarus and Huang, 2011). Interestingly, they observed different trends than we reported here, as SOM cells in the visual cortex displayed increases in input resistance and membrane time constant with age. These different developmental trajectories seen by Lazarus and Huang (2011) could reflect differences between cortical areas. Additionally, these authors investigated SOM and PV cells from ages P15-P30, and grouped the data into bins of multiple days, including P15 to P17. Because the decreases we observed in SOM cell input resistance and membrane time constant occurred at earlier ages (greatest changes prior to day 14), it is difficult to compare across these studies.

Intrinsic cellular properties during postnatal development

Previous studies have examined the properties of pyramidal neurons during the age range studied here and have observed a similar decrease in intrinsic excitability of these cells with age (McCormick and Prince, 1987; Maravall et al., 2004; Oswald and Reyes, 2008). Interestingly, although we observed changes to these properties in GIN and RS cells as a function of age, GIN cells consistently exhibited higher R_{in} , higher τ_m , and lower rheobase current than RS cells of the same age. This illustrates that the relative intrinsic excitability between GIN and RS cells is preserved at all ages. This may be important for maintaining the proper ratio of excitation to inhibition throughout development (Zhang et al., 2011).

In excitatory neurons and other inhibitory interneurons, such as the fast-spiking cells, changes in input resistance are driven primarily by increased expression of potassium leak currents (Goldberg et al., 2011; Guan et al., 2011), which could have caused the change in input resistance we reported for GIN cells. It is currently unknown whether the amount of synaptic input or gap junction coupling changes over development in GIN cells, but either could also contribute to the changes in intrinsic properties observed here.

A prominent feature of GIN cells is the presence of a strong ‘sag’ current and related rebound depolarization during and after negative current pulses, respectively (McGarry et al., 2010; Ma et al., 2012). Both of these properties are mediated by I_h currents in GIN cells (Ma et al., 2006). We found a pronounced increase in the magnitude of the sag current and rebound depolarization in GIN cells, illustrating that the expression of I_h is likely increasing in GIN cells over development. I_h currents often underlie the generation of rhythmic activity (McCormick and Pape, 1990; Maccaferri and McBain, 1996; Pape, 1996; Luthi and McCormick, 1998; Griguoli et al., 2010). GIN cells in both the neocortex and hippocampus show a strong preference for firing at theta frequencies (Chapman and Lacaille, 1999b, a; Klausberger et al., 2003; Lawrence et al., 2006; Fanselow et al., 2008; Griguoli et al., 2010). This could result partially from the expression of I_h current (Griguoli et al., 2010). A developmental increase in I_h current could cause the theta preference found in GIN cells to mature during this period of postnatal development.

Maturation of GIN action potential waveform

A dominant factor that controls the action potential waveform is the expression of voltage-gated potassium channels (Rudy and McBain, 2001). In particular, the potassium channel family, Kv3,

has been shown to underlie the narrow action potential waveform in fast-spiking (FS) inhibitory interneurons (Rudy and McBain, 2001; Lien and Jonas, 2003; Goldberg et al., 2011). The action potential half-width observed in FS cells decreases during the second and third weeks of postnatal development, and this property is paralleled by increased expression of Kv3 over development (Goldberg et al., 2011). It is possible that developmental increases in potassium currents could also underlie the decreased action potential half-widths observed in GIN cells in the current study. Although Kv3 channels are observed in deep layer SOM inhibitory interneurons, they are not present in superficial SOM cells (Chow et al., 1999), which were recorded here. The specific types of potassium channels expressed in superficial SOM cells are not known, so it is difficult to know which potassium currents could underlie the developmental changes in action potential waveform we observed in our study.

We demonstrated that the voltage trajectory following the AHP trough depolarized more rapidly in older GIN cells than in younger ones. The ability of mature GIN cells to repolarize more rapidly following an action potential may underlie the increased firing rates observed at older ages. I_h currents can act on the shape and duration of the AHP and increase the slope of the membrane potential during the interspike interval (McCormick and Pape, 1990; Maccaferri and McBain, 1996; Pape, 1996; Griguoli et al., 2010). The changes observed here in the AHP shape and slope therefore further support the idea that I_h currents increase in GIN cells over the second and third postnatal weeks.

GIN cells display multiplicative increases in the input-output relationship with age

We assessed the firing output of GIN and RS cells over development by examining changes in firing rates. The initial and steady state firing rates of GIN cells increased with age. Because the rheobase current remained largely constant in GIN cells, the increase in firing rate resulted in a multiplicative increase in the input-output curve of GIN cells as a function of age. Such gain changes can arise from alterations in the rate and balance of excitatory and inhibitory synaptic inputs to a neuron (Chance et al., 2002; Mitchell and Silver, 2003; Larkum et al., 2004). Another way to modulate the gain of a neuron is to affect a process that is spike-dependent, such as the action potential AHP (Mehaffey et al., 2005; Diaz-Quesada and Maravall, 2008). This will increase or decrease the firing rate of a neuron without a concomitant change in rheobase current. Our observed changes to the AHP in GIN cells as a function of age could account for the increased gain we observed in these cells.

In contrast, in RS cells the more prominent change in firing rate over development was an increase in rheobase current, accompanied by a rightward shift of the input-output curve (Oswald and Reyes, 2008). Additive and subtractive shifts in input-output curves can be caused by alterations in intrinsic neuronal excitability (Silver, 2010), which do not cause changes in gain. The changes in intrinsic properties (such as input resistance) of RS cells are likely the cause of the subtractive change in the input-output relationship observed in these cells over development.

Characterizing subtypes of inhibitory neurons

Establishing relevant distinctions between subtypes of inhibitory neurons, while critical for characterizing their role(s) in the neocortex, has been difficult (Gupta et al., 2000; Markram et al., 2004; Ascoli et al., 2008; Guerra et al., 2011). This has been aided, but not entirely

ameliorated, by the use of lines of mice that express fluorescent proteins in inhibitory neurons of interest. It is not clear whether GIN cells form a homogeneous population. These cells have been subdivided in to three (McGarry et al., 2010) and four (Halabisky et al., 2006) groups based on physiological and morphological characteristics. In our study, we showed that physiological factors frequently used to characterize inhibitory neurons vary with age. Therefore, in order to use these factors to accurately categorize GIN neurons it is necessary to take age into account. Here, we focused on changes in the physiological features of GIN cells, but other features, such as morphological characteristics, can also be considered when subdividing GIN cells (McGarry et al., 2010). It would be interesting to know whether the morphological features of GIN cells also develop over this time window, and whether they change in concert with the physiological changes we have described. In fact, developmental morphological changes could constitute possible mechanisms for some of the physiological changes we observed. For example, changes in overall dendritic length with age could account for the decreased input resistance in GIN cells we report in this study.

Impact of GIN cell maturation on cortical circuitry

This study represents the first in-depth analysis showing how the physiology of GIN cells in layers 2/3 of somatosensory cortex changes during the second and third postnatal weeks. This encompasses a period of massive reorganization of the neocortex correlated with the onset of sensory input and plasticity. Our results suggest that GIN cells become more active in the cortical network around P14, which corresponds to the approximate time of whisking onset in mice and rats (Mosconi et al., 2010). Although a precise function for GIN cells, and more

broadly SOM cells, in sensory processing is still being investigated, there are several implications of our results.

First, changes in intrinsic properties of SOM cells can affect how these cells communicate amongst themselves via electrical connections, or gap junctions. Gap junctions are common between SOM cells (Gibson et al., 1999; Fanselow et al., 2008), and are involved in the generation of synchronous activity between SOM interneurons (Beierlein et al., 2000; Fanselow et al., 2008). The ability of electrically coupled cells to synchronize depends on both the properties of the coupling as well as the intrinsic properties of the individual cells (Amitai et al., 2002; Lewis and Rinzel, 2003; Mancilla et al., 2007; Hayut et al., 2011). Therefore, as the intrinsic properties of SOM cells mature during postnatal development the ability of SOM cells to synchronize their activity through gap junctions may change. Factors such as input resistance, membrane time constant, and AHP size and shape could all influence gap junction-mediated synchrony between GIN cells. However, determining how the changes we saw in these factors as a function of age would individually or collectively affect GIN-GIN synchrony would require detailed computational and experimental investigations that are beyond the scope of this study.

Second, SOM cells participate in cortical ‘UP-states’ (Fanselow and Connors, 2010), brief periods of depolarization and higher neuronal activity observed in a range of neocortical regions and layers. UP-states have been described *in vitro* (Sanchez-Vives and McCormick, 2000; Shu et al., 2003; Fanselow and Connors, 2010), *in vivo* during anesthesia (Steriade et al., 1993; Sanchez-Vives and McCormick, 2000) and in sleep and waking states (Steriade et al., 2001; Petersen et al., 2003). The function of UP-states is not well understood, but they represent

a high-conductance state with neuronal activity that is reminiscent of firing observed in intact animals (Destexhe and Pare, 1999; Destexhe et al., 2003; Rudolph et al., 2005; Fanselow and Connors, 2010). Work by Fanselow and Connors (2010) suggested that GIN cells could be involved in terminating UP-states. If, as our current results suggest, GIN cells play a larger role in cortical activity as an animal progresses through the second and third weeks of postnatal development, then the propensity for GIN cells to terminate UP-states might also increase during the same period. Indeed, Fanselow and Connors (2010) demonstrated a strong inverse correlation between UP-state duration and age, as well as a decrease in UP-state frequency with increasing age from P13 to P17. Such an ability to terminate UP-states could potentially extend to termination of other periods of heightened activity within the cortex, such as seizures. In fact, loss of SOM, dendrite-targeting inhibitory cells of the hippocampus has been associated with increased seizure activity (Buckmaster and Jongen-Relo, 1999; Cossart et al., 2001).

Third, SOM cells influence the activity of the surrounding neuronal network in multiple ways. These mechanisms would likely be affected by the type of age-driven changes in the input-output relationship (gain) of GIN cells that we observed. For example, because SOM interneurons directly inhibit FS cells (Gibson et al., 1999; Ma et al., 2012), an increase in SOM cell gain could paradoxically increase local excitatory neuron activity through disinhibition. In contrast, it has been shown that activation of even a few pyramidal cells exerts strong disynaptic inhibition onto surrounding pyramidal cells via SOM interneurons (Kapfer et al., 2007; Silberberg and Markram, 2007). In this way, increases in SOM cell gain may serve to increase SOM-mediated disynaptic self-inhibition of pyramidal cells, resulting in a decrease in cortical excitability. Note that these two methods of regulating cortical activity are not mutually exclusive, but may instead work antagonistically to fine tune cortical activity.

It will be useful in future studies to determine how developmental changes to synaptic connections complement development of the intrinsic properties we have described here, and, in turn, affect the influence SOM cells have on neighboring neurons. Excitatory inputs onto SOM cells are initially weak, but display strong short-term facilitation (Gibson et al. 1999, Fanselow et al, 2008), which suggests these cells are engaged during periods of heightened neuronal activity (Kapfer et al., 2007; Silberberg and Markram, 2007; Fanselow et al., 2008; Hayut et al., 2011). Interestingly, inhibitory output from SOM cells initially exhibits weak depression followed by weak facilitation (Ma et al., 2012). Together, these short-term dynamics suggest that SOM cells would be able to increase and then sustain their inhibitory influence on downstream targets during sensory processing. This is in contrast to FS cells, which receive initially strong but rapidly depressing synapses from upstream neurons and make depressing connections onto target neurons (Beierlein et al., 2003; Ma et al., 2012). This suggests that FS cells, while exerting strong inhibitory output at first, would decrease their influence on surrounding cells as ongoing neuronal activity progressed. These opposing short-term dynamics suggest SOM and FS cells may be specialized for distinct modes of inhibitory processing: FS cells are optimized for fast, transient inhibition, whereas SOM cells can provide somewhat delayed, but sustained, inhibition (Tan et al., 2008). The extent to which each of these functions is fulfilled at a given point during development remains to be determined. Developmental changes in the short-term dynamics of synapses onto and from SOM cells would dictate the degree to which these cells could be activated by high-frequency inputs and the time course over which they could affect their downstream targets. Additionally, inhibitory synapses between SOM cells and fast-spiking (FS) cells can produce synchronous activity between these different types of inhibitory interneurons (Hu et al., 2011). Therefore, developmental regulation of the inhibitory output from SOM cells

may impact the ability of SOM cells to synchronize with FS cells through inhibitory synapses. Further investigation of SOM input and output properties (e.g. synaptic dynamics) as a function of age will be critical to our understanding of the role of SOM cells during this postnatal time period.

Finally, multiple studies *in vitro*, *in vivo*, and using computational models have suggested unique roles for SOM inhibitory interneurons during cortical sensory processing (Tan et al., 2008; Vierling-Claassen et al., 2010; Hayut et al., 2011; Ma et al., 2012). For example, inhibition onto the apical dendrite of pyramidal cells can inhibit pyramidal cell firing by influencing active currents in this dendritic sub-region (Larkum et al., 1999, 2001; Perez-Garci et al., 2006; Gentet et al., 2012; Lovett-Barron et al., 2012). This suggests that SOM cells may be involved in regulating excitatory synapses onto such apical dendrites, including long-range projections from other cortical and subcortical areas (Petreanu et al., 2007; Petreanu et al., 2009). In contrast, FS cells, which terminate on the perisomatic regions of their targets, would not directly control these inputs. Thus, developmentally mediated increases in SOM cell activity due to increases in gain might more closely regulate long-range inputs to somatosensory cortex at later ages. In addition, SOM cells exhibit different behaviors *in vivo* than other inhibitory interneurons, including FS cells. Specifically, it has been shown that SOM cells are preferentially active during quiet wakefulness, and are hyperpolarized in response to sensory input, whereas FS cells respond robustly to sensory input (Gentet et al., 2012). If, as these findings collectively suggest, increased SOM cell activity gates out excitatory distal dendritic input during sensory processing, then it would appear from our study that such dendritic input (e.g. from motor cortex) becomes increasingly gated as a function of age. Whether and why more excitatory dendritic input might

be useful early in the development of the sensorimotor system, or a detriment at older ages, is not yet known. In any case, it is clear that SOM and FS cells possess distinct intrinsic and synaptic properties that may contribute to their different behaviors *in vivo*. The development of these properties in both FS and SOM cells happens concurrently, over the second and third postnatal weeks, and may contribute to the dichotomous roles of these subsets of interneurons in neocortical processing.

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