Morphologically constrained modeling of spinous inhibition in the somato-sensory cortex

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ABSTRACT

Pyramidal neurons are covered by thousands of dendritic spines receiving excitatory synaptic inputs. The ultrastructure of dendritic spines shapes signal compartmentalization but ultrastructural diversity is rarely taken into account in computational models of synaptic integration. Here, we developed a 3D correlative light-electron microscopy (3D-CLEM) approach allowing the analysis of specific populations of synapses in genetically defined neuronal types in intact brain circuits. We used it to reconstruct segments of basal dendrites of layer 2/3 pyramidal neurons of adult mouse somatosensory cortex and quantify spine ultrastructural diversity. We found that 10% of spines were dually-innervated and 38% of inhibitory synapses localize to spines. Using our morphometric data to constrain a model of synaptic signal compartmentalization, we assessed the impact of spinous versus dendritic shaft inhibition. Our results indicate that spinous inhibition is locally more efficient than shaft inhibition and that it can decouple voltage and calcium signaling, potentially impacting synaptic plasticity.

INTRODUCTION

In the mammalian cortex, the vast majority of excitatory synapses are formed on dendritic spines, small membrane protrusions that decorate the dendrites of pyramidal neurons (PNs) (Ramon y Cajal, 1899; Hersch and White, 1981; Yuste and Bonhoeffer 2004). Dendritic spines are composed of a bulbous head connected to the dendritic shaft by a narrow neck (Arellano, 2007; Harris and Weinberg, 2012). They exist in a large variety of shapes and sizes along individual dendrites. Spine head volume can vary between 3 orders of magnitude (0.01-1.5 μm^3), neck length between 0.2 μm and 3 μm, and minimal neck diameter between 20 and 500 nm (Harris et al., 2015). Spine heads are typically contacted by an excitatory synaptic input and harbor an excitatory post-synaptic density (ePSD) that contains glutamatergic α-amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid (AMPA) and N-Methyl-D-aspartate (NMDA) neurotransmitter receptors, scaffolding proteins, adhesion molecules and a complex machinery of proteins undertaking the transduction of synaptic signals. The size of the spine head correlates with the size of the ePSD and the strength of synaptic transmission (Fifková and Anderson, 1981; Nusser et al., 1997; Matsuzaki et al., 2004; Bourne and Harris, 2011; Holderith et al., 2012). In addition to the ePSD, spines contain ribosomes, which mediate local protein synthesis, and endosomes, which play a critical role in membrane and receptor trafficking (Bailey et al., 2015; Kennedy and Hanus, 2019). The largest spines often contain a spine apparatus (SA), which contributes to calcium signaling and synaptic plasticity (Korkotian et al., 2014; Bailey et al., 2015), and some spines, especially in the upper layers of the cortex,

are also contacted by an inhibitory synaptic input (dually-innervated spines or DiSs; Kubota et
al., 2007). Spine necks are diffusional barriers that biochemically isolate spine heads from
their parent dendrite (Alvarez and Sabatini, 2007; Tønnesen et al., 2014; Adrian et al., 2017;
Yasuda, 2017). In addition, they can filter the electrical component of synaptic signals and
amplify spine head depolarization (Harnett et al., 2012; Yuste, 2013; Tønnesen and Nägerl,
2016; but see Svoboda et al., 1996; Takasaki and Sabatini, 2014; Popovic et al., 2015). Both
spine heads and spine necks are remodeled depending on neuronal activity (Matsuzaki et al.,
2004; Honkura et al., 2008, Araya et al., 2014) and in pathology (Androuin et al., 2018; Forrest
et al., 2018). While widely acknowledged, the extent of synaptic ultrastructural diversity along
individual dendrites has not been quantified, and the consequences of this variability on signal
compartmentalization and dendritic integration remain to be investigated.

Dendritic signaling can be modeled based on anatomical and biophysical parameters
(Poirazi and Papoutsi, 2020) using "realistic" multi-compartment models (Almog and
Korngreen, 2016). These models were pioneered by Wilfrid Rall following the seminal works
of Hodgkin and Huxley (Rall, 1970; Hodgkin and Huxley, 1952). They have provided a
powerful theoretical framework for understanding dendritic integration (Stuart and Spruston,
2015), spine function (Segev and Rall, 1988), inhibitory signaling (Gidon and Segev, 2012;
Doron et al., 2017) and electrical compartmentalization in spines (Yuste et al., 2000, Gulledge
et al., 2012; Tønnesen and Nägerl, 2016). However, spines and synapses are usually modeled
with *ad hoc* averaged biophysical parameters, which limits prediction accuracy. Modeling the
actual behavior of dendritic spines requires an accurate description of their ultrastructural
heterogeneity along identified dendrites of specific types of neurons. To acquire such data, it
is necessary to combine the nanometer resolution of electron microscopy (EM) with an
approach that allows the identification of the origin of dendritic spines (i.e. location on the
dendrite, type of dendrite and type of neuron) without obscuring the intracellular content. This
task is arduous: 1 mm^3 of mouse cortex contains over 50,000 of neurons, each of which
establishes approximately 8,000 synaptic connections with neighboring neurons, and these
synapses are highly specific, connecting multiple neuronal subtypes from various brain
regions (Schüz and Palm, 1989; Herculano-Houzel et al., 2013; Motta et al., 2019; Sanes and
Zipursky, 2020). Reconstructing selected dendritic spines and synaptic contacts along
dendritic trees requires either: enormous volumes of 3D-EM acquisitions using resource-
consuming approaches adapted from connectomics (Briggman and Denk, 2006;
Helmstaedter, 2013; Kasthuri et al., 2015; Karimi et al., 2020), or combining EM with a lower-
scale imaging modality, such as confocal or 2-photon light microscopy (LM), to guide 3D-EM
image acquisitions to the region of interest (ROI) (De Boer et al., 2015; Begemann and Galic,
2016; Müller-Reichert and Verkade, 2017). While very powerful *in vitro* (Watanabe et al., 2011;
Russell et al., 2017, de Boer et al., 2015), correlative light-electron microscopy (CLEM) is

difficult to implement in brain tissues (Collman et al., 2015; Lees et al., 2017; Maclachlan et
al., 2018). New protocols are required to facilitate the identification of targeted dendrites and
synapses in 3D-EM.

95   Here, we have developed a CLEM workflow combining confocal light microscopy with
serial block-face scanning EM (SBEM) and targeted photo-precipitation of 3,3-
diaminobenzidine (DAB) to facilitate ROI recovery. We applied this workflow to reconstruct
dendritic spines located exclusively on the basal dendrites of genetically-labelled PNs in layers
2/3 of the somatosensory cortex (SSC) of adult mice. We analyzed the variability of their
ultrastructure and estimated the electrical resistance of their neck. We also examined the
distribution and the morphology of inhibitory synapses. We specifically examined dendritic
spines receiving both excitatory and inhibitory inputs, which represented 10% of all spines
along basal dendrites. These dually-innervated spines (DiSs) exhibited wider heads and larger
ePSDs than singly-innervated spines (SiSs), and they were more electrically isolated from the
dendritic shaft than SiSs of comparable head size. We then used our measurements to
constrain a multi-compartment model of synaptic signaling and compartmentalization in
dendrites. We assessed the effects of individual excitatory and inhibitory signals on membrane
voltage and calcium concentration depending on inhibitory synapse placement (i.e. on a spine
head or on the dendritic shaft) and input timing. Our results challenge the view that spinous
inhibition strictly vetoes single excitatory inputs and rather suggest that it fine-tunes calcium
levels in DiSs. Our simulations indicate that a single inhibitory postsynaptic potential (IPSP)
evoked in a DiS within 10 ms after an excitatory postsynaptic potential (EPSP) can curtail the
local increase of calcium concentration without affecting the amplitude of membrane
depolarization. This decoupling effect could impact long-term synaptic plasticity in cortical
circuits.
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RESULTS

Combining light and electron microscopy to access the ultrastructure of targeted
populations of dendritic spines in brain slices
In the cortex, the morphology and distribution of dendritic spines vary depending on cortical
area and layer in which the cell body is located (Katz et al., 2009; Harris and Weinberg, 2012;
Bosch et al., 2015; Stuart and Spruston, 2015), and dendritic spines are differently regulated
depending on their location within dendritic trees — e.g. basal or apical dendrites (Globus et
al., 1973; Feldmeyer et al., 2012; Bian et al., 2015; Rajkovich et al., 2017; Karimi et al., 2020).
Therefore, it is critical to take into account both the cellular and dendritic context to
characterize the diversity of spine ultrastructure. To that aim, we developed a 3D-CLEM
workflow allowing the ultrastructural characterization of dendritic spines on genetically-defined

neuronal cell types and along identified types of dendrites in intact cortical circuits. In order to label specific subtypes of neurons, we used cortex-directed *in utero* electroporation (IUE) in mice. We electroporated neuronal progenitors generating layer 2/3 cortical PNs at embryonic day (E)15.5 with a plasmid expressing the fluorescent cytosolic filler tdTomato, granting access to the morphology of electroporated neurons, their dendrites and their dendritic spines in LM. We perfused adult mice with aldehyde fixatives, and collected vibratome sections of the electroporated area. To facilitate sample handling, we designed custom-made chambers allowing sample immersion in different solutions during confocal imaging and subsequent retrieval of the sample before EM preparation steps (Supplementary figure 1A). We enclosed 10-20 mm^2 fragments of brain sections in these chambers and acquired images of optically isolated basal dendrites of bright electroporated neurons with confocal microscopy (Figure 1A).

A major challenge of CLEM in brain tissue is to recover the ROI in EM after imaging in LM. Several methods have been proposed to facilitate ROI recovery (De Boer et al., 2015; Begemann and Galic, 2016; Müller-Reichert and Verkade, 2017), but they come with caveats: (1) using only intrinsic landmarks has a low throughput (Luckner et al., 2018; Maclachlan et al., 2018); (2) filling target neurons with 3,3-diaminobenzidine (DAB) masks intracellular ultrastructure (Meißlitzer-Ruppitsch et al., 2009); (3) scarring the tissue with an infra-red laser to generate extrinsic landmarks, a.k.a. "NIRB", for "near-infrared branding" (Bishop et al., 2011; Maco et al., 2013; Cane et al., 2014; Blazquez-Llorca et al., 2015; Lees et al., 2017), produces landmarks with low pixel intensity in EM and can damage ultrastructure (Karreman et al., 2016; Luckner et al., 2018). To facilitate ultrastructural measurements in non-obscured identified dendrites, we took advantage of the photo-oxidability of DAB (Lübke, 1993; Grabenbauer et al., 2005). We immersed the samples in DAB solution and applied focalized UV light at user-defined positions (Figure 1A) to imprint osmiophilic DAB landmarks around targeted dendrites (Supplementary figure 1B-E) and pattern the tissue with localized electron-dense DAB precipitates (Figure 1B). After sample retrieval (Supplementary figure 1F), tissue sections were processed for SBEM and embedded in minimal amounts of epoxy resin in order to maximize sample conductivity and SBEM image quality (see Methods). In 3D-EM stacks, ROIs were recovered within the complex environment of brain tissues using both intrinsic landmarks such as blood vessels (Figure 1B), and high-contrast DAB precipitates (Figure 1C; see also supplementary figure 1G). We then segmented and reconstructed targeted dendrites in 3D, and registered whole dendritic portions in both LM and EM to identify each dendritic spine unequivocally using neighboring spines as dependable topographic landmarks (Figure 1D). CLEM-based 3D-reconstruction enabled the identification of dendritic spines that were not visible in LM or EM alone. In LM, the limited axial resolution prevents the identification of axially oriented spines, which are easily detected in 3D-EM (Karimi et al., 2020) (Figure 1D).

165 On the other hand, spines with the longest and thinnest necks are conspicuous in LM stacks,
166 but can be difficult to find in 3D-EM datasets without the cues provided by LM. The proportion
167 of spines recovered with CLEM versus LM alone could amount to up to 30% per ROI, and 5%
168 per ROI versus EM alone, highlighting the advantage of CLEM over unimodal microscopy
169 approaches.
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Spine ultrastructure along the basal dendrites of L2/3 cortical pyramidal neurons.
We used our CLEM workflow to quantify the full extent of the ultrastructural diversity of
dendritic spines along the basal dendrites of layer 2/3 PNs of the SSC of three adult mice. We
exhaustively segmented 254 μm of the basal dendritic arborization of four neurons and we
reconstructed a total of 390 individual spines (Supplementary table 1). As spine distance to
the soma spanned from 20 to 140 μm, with basal dendrites extending up to 150 μm
(Ballesteros-Yañez et al., 2006; Chen et al., 2012; Sarid et al., 2015; Kubota et al., 2015), we
consider that our dataset is representative of the whole spine population on these dendrites.
The average linear density of dendritic spines was 1.5 ± 0.3 spine.μm^{-1}. We then quantified
the following parameters for each spine: neck length, neck diameter, head volume, head
longitudinal diameter (referred to as "head length"), head orthogonal diameter (referred to as
"head diameter"), number of PSDs, and PSD area (Figure 2A; Supplementary table 2). In
agreement with previous reports in both basal and apical dendrites of mouse cortical and
hippocampal neurons (Knott et al., 2006; Arellano, 2007; Noguchi et al., 2011; Harris et al.,
2014), we found that ePSD area correlates linearly with the volume of the spine head (Figure
2B). We also observed a non-linear correlation between the length of the spine neck and its
diameter (Figure 2C): long spines (neck length > 2 μm) always had thin necks (neck diameter
< 0.2 μm), although short necks could also be thin. Furthermore, in spines with long necks,
spine heads were always stretched longitudinally with respect to the neck (i.e. prolate)
whereas they could also be stretched orthogonally (i.e. oblate) in shorter spines (Figure 2D),
with a possible impact on nanoscale ion flows (Bell et al., 2019; Lagache et al., 2019). By
contrast, there was no correlation between the position of the spine or between the inter-spine
distance and any of the morphological parameters we measured (data not shown), in line with
previous ultrastructural studies (Arellano, 2007). There was also no correlation between the
length or the diameter of the neck and the morphometry of the spine head or ePSD, which is
consistent with previous EM studies of L2/3 PNs of mouse neocortex (Arellano, 2007;
Ballesteros-Yáñez et al. 2006; but see Gulledge et al., 2012, and Noguchi et al., 2005 for
different conclusions in other brain areas).
Since our CLEM approach grants access to the cytosolic content of spines (Figure 3A),
we quantified the occurrence of SA, a complex stacked-membrane specialization of smooth
endoplasmic reticulum (SER) which contributes to calcium signaling, integral membrane

protein trafficking, local protein synthesis, and synaptic plasticity (Bourne and Harris, 2008;
Segal and Korkotian, 2014; Korkotian et al., 2014; Bailey et al., 2015; Kennedy and Hanus,
2019). In basal dendrites, about 54% of spines contained a SA (Figure 3B), which is
substantially higher than previous reports in the mature hippocampus (Bailey et al., 2015;
Chirillo et al., 2019). These spines were randomly distributed along the dendrites. They had
larger heads (Figure 3C), larger ePSDs (Figure 3D) and wider necks than spines devoid of
SA (Figure 3E), consistent with previous morphological studies of CA1 PNs (Bailey et al.,
2015; Spacek and Harris, 1997; Chirillo et al., 2019). The probability that a spine contained a
SA depending on spine head volume followed a sigmoid model (Figure 3F), predicting that all
spines with a head diameter larger than 1.1 μm (21% spines in our reconstructions) contain a
SA. Interestingly, SER was also often detected at the stem of half of spines devoid of SA (25%
of all spines), but these spines did not differ in their morphology from other spines devoid of
SA (data not shown).

Next, we used our ultrastructural data to estimate the electrical resistance of spine
necks using $R_{neck} = \rho W_{neck}$, where ρ is the cytosolic resistivity (set to 300 Ω.cm; Elbohouty et
al., 2013; Miceli et al., 2017) and W_{neck} is the diffusional neck resistance that restricts the
diffusion of molecules and charges between spine heads and dendritic shafts (Svoboda et al.,
1996). To quantify W_{neck}, for each spine we measured a series of orthogonal cross-sections
of the neck along its principal axis and integrated $W_{neck} = \int dl/A(l)$, where A(l) is the neck
cross-section area at the abscissa l along the neck axis. W_{neck} ranged from 2 μm^{-1} to 480 μm^{-1}
$^{-1}$ and R_{neck} from 8 MΩ to 1450 MΩ, with a median value of 188MΩ. These values are consistent
with previous estimations based on EM reconstructions and STED super-resolutive light
microscopy (Harris and Stevens, 1989; Tønnesen et al., 2014), and with direct
electrophysiological recordings (Jayant et al, 2017). It has been proposed that spine
apparatuses, which occupy some of the spine neck volume, could increase W_{neck} (Bourne and
Harris, 2008; Lagache et al., 2019; Kennedy and Hanus, 2019). Therefore, we subtracted SA
cross-section from A(l) when computing W_{neck} in SA+ spines (see Methods). This correction
increased W_{neck} by 13% ± 2% in SA+ spines (Supplementary figure 2). However, because of
their wider necks, W_{neck} of SA+ spines was still lower (59% in average) than W_{neck} of spines
devoid of SA (Figure 3G). These results suggest that, in addition to supplying large dendritic
spines with essential resources, the SA may adjust W_{neck} and influence spine
compartmentalization (Bailey et al., 2015; Chirillo et al., 2019; Kennedy and Hanus, 2019).

Excitatory and inhibitory synapses in dually-innervated spines.
We noticed that a significant proportion of dendritic spines were contacted by two distinct pre-
synaptic boutons (DiSs). DiSs have long been described in the literature as receiving both an
excitatory and an inhibitory synaptic contact (Jones and Powell, 1969; Beaulieu and Colonnier,

1985; Dehay et al., 1991; Fifková et al., 1992). In the somato-sensory cortex, DiSs are
contacted by VGLUT2-positive thalamocortical inputs (Kubota et al., 2007) and they are
sensitive to sensory experiences. The number of DiSs increases in response to sensory
stimulation and decreases in response to sensory deprivation (Micheva and Beaulieu, 1995;
Knott et al., 2002; Chen et al., 2012; van Versendaal et al., 2012), suggesting their importance
in synaptic integration and sensory processing. While ePSDs look asymmetrical and more
electron-dense than inhibitory PSDs (iPSDs) in transmission EM (Gray, 1959; Uchizono,
1965), the anisotropic resolution of SBEM does not allow the distinction of ePSDs and iPSDs
in most DiSs (Karimi et al., 2020). Therefore, to label inhibitory synapses in cortical PNs
(Figure 4A), we co-expressed tdTomato with small amounts of GFP-tagged gephyrin (GFP-
GPHN) (Chen et al., 2012; van Versendaal et al., 2012; Fossati et al., 2016; Iascone et al.,
2020), a core component of inhibitory postsynaptic scaffolds (Tyagarajan and Fritschy, 2014;
Krueger-Burg et al., 2017). We identified in LM spines containing a gephyrin cluster (Figure
4B) and we ascertained their dual-innervation in EM after back-correlating spine identity
between LM and SBEM acquisitions. To do so, we aligned reconstructed dendrites on LM
images (Figure 4C), and matched individual spines in both modalities (lettered in Figures 4B
and 4C). We identified iPSDs on DiSs based on GFP-GPHN cluster position in LM images. In
89% of DiSs (33/37), the excitatory (GFP-GPHN-negative) PSD and the inhibitory (GFP-
GPHN-positive) PSD could be clearly discriminated. However, in 11% of DiSs (4/37 DiSs),
distinguishing ePSD from iPSD was not obvious due to the coarse axial resolution of LM
imaging. To resolve ambiguities, we reconstructed the axons innervating the DiSs and
determined their identity based on their other targets in the neuropil, either soma and dendritic
shaft for inhibitory axons (Hersch and White, 1981; Isaacson and Scanziani, 2011; Boivin and
Nedivi, 2018), or other dendritic spines for excitatory axons (Karimi et al., 2020) (Figure 4D).
As a result, we could unequivocally determine the excitatory or inhibitory nature of each
synaptic contact on electroporated neurons, within ~10^5 µm^3 3D-EM acquisition volume.

In CLEM, we measured an average density of 1.4 ± 0.5 iPSDs per 10 µm of dendrite
on DiSs and 2.1 ± 1.2 iPSDs per 10 µm of dendrite on the dendritic shaft— amounting to 3.5
± 1.1 iPSDs per 10 µm of dendrite. We observed a homogeneous distribution of iPSDs located
either on spines or shaft, from 24 µm away from the soma to the dendritic tip, in contrast to
apical dendrites where spinous inhibitory synapses are distally enriched (Chen et al., 2012).
Along the basal dendrites of L2/3 cortical PNs, 38% of inhibitory contacts occurred in dendritic
spines, which is in the range of, or slightly higher than previously estimated using LM only
(Chen et al., 2012; Iascone et al., 2020). DiSs represented 10% ± 3% of all spines (Figure
5A). They had larger heads than SiSs (Figure 5B), in line with previous reports (Kubota et al.,
2007; Villa et al., 2016), and 86% ± 13% of them contained a SA (Figure 5C). DiSs also differed
in terms of neck morphology. They had longer necks than SiSs of comparable head volume

(V_{head} >0.05 µm^3), although neck length distribution was similar in the whole populations of
SiSs and DiSs (Figure 5D). DiSs also had lower D_{neck}/V_{head} ratio than SiSs (Figure 5E),
although D_{neck} distribution was similar between SiSs and DiSs (Supplementary figure 3),
suggesting that excitatory signals generated in DiSs are more compartmentalized than signals
of similar amplitude generated in SiSs. Accordingly, DiSs had a higher W_{neck} than SiSs of
comparable head size (52% larger in average) (Figure 5F). In spine heads, ePSDs on DiSs
were larger than ePSDs on SiSs (174% ± 113% of ePSD area) (Figure 5G), consistent with
the larger head size of DiSs. By contrast, iPSDs on DiSs were smaller than shaft iPSDs (53%
± 15% of shaft iPSD area) (Figure 5H), in line with previous reports in rat L5 PNs (Kubota et
al., 2015). The area of iPSDs on DiSs did not correlate with spine head volume (data not
shown). In 95% of DiSs, iPSDs were smaller than ePSDs (half the area, in average) (Figure
5I). Together, these results indicate that DiSs represent a specific population of dendritic
spines with distinctive ultrastructural features that could impact their functional properties.
289
Morphologically constrained model of synaptic signaling
Next, we wanted to assess the impact of spine diversity on synaptic signals. We used a
computational approach based on a multi-compartment "ball-and-stick" model of the neuronal
membrane (Rall, 1964; Gulledge et al., 2012). This model comprises an isopotential soma and
two dendritic compartments structured as cables featuring passive resistor-capacitor (RC)
circuits and conductance-based synapses. The two dendritic compartments correspond to the
dendrite receiving the synaptic inputs and to the remainder of the dendritic tree (Figure 6A1)
(Dayan and Abbot, 2001; Stuart et al., 2016). We constrained this model with morphological
parameters measured in CLEM (i.e. distance between spine and soma, neck resistance, head
volume, head membrane area, ePSD area and iPSD area for 390 spines, and dendritic
diameter), taking into account the structural shrinkage resulting from chemical fixation
(Supplementary figure 4). Excitatory and inhibitory synaptic conductances were modeled as
bi-exponential functions of time, with their rise and decay times tuned to the kinetics of different
receptor types: AMPA and NMDA receptors at ePSDs, and type A γ-aminobutyric acid
(GABA$_A$) receptors at iPSDs (Figure 6A2; see Methods). Individual synaptic conductances
were scaled proportionally to PSD areas (Nusser et al., 1998; Matsuzaki et al., 2004; Noguchi
et al., 2011). Voltage-dependent calcium channels (VDCCs) in spine heads were modeled
using Goldman–Hodgkin–Katz equations (Koch and Segev, 1998) and their conductance was
scaled proportionally to spine head areas.
We adjusted excitatory synaptic conductivity so that average amplitudes of both
synaptic currents and somatic depolarizations evoked by individual excitatory post-synaptic
potentials (EPSPs) fitted published electrophysiological values (Feldmeyer et al., 2006, Frik
et al., 2008, Lefort et al., 2009, Scala et al., 2018) (see Methods). After calibration of excitatory

synapses, maximal synaptic conductance ranged from 0.04 nS to 3.13 nS for g_{AMPA} (0.456 ± 0.434 nS) and from 0.04 nS to 3.42 nS for g_{NMDA} (0.498 ± 0.474 nS), in line with the literature (Hoffmann et al., 2015). We then adjusted inhibitory synaptic conductivity to set the mean conductance of dendritic inhibitory synapses to 1 nS (Gidon and Segev, 2012; Xu et al., 2013; Kobayashi et al., 2008). As a result, g_{GABA} ranged from 0.33 nS to 3.36 nS (1.00 ± 0.577 nS) for synapses located on the shaft, and from 0.19 nS to 1.56 nS (0.528 ± 0.277 nS) for inhibitory synapses located on spines.

We first examined the propagation of simulated EPSPs. We compared the evoked depolarization amplitude ΔV_{max} in three different compartments: in the spine head in which the EPSP was elicited, in the dendritic shaft close to the spine, and in the soma (Figure 6B). ΔV_{max} followed a log-normal distribution, reflecting the morphological variability of spines (Figure 6B). Due to the passive attenuation of electrical signals along dendritic processes, ΔV_{max} was sharply attenuated between the head of the spine and the dendritic shaft (51% attenuation in average) and about 5% of ΔV_{max} reached the soma (Figure 6B), in line with measurements performed in basal dendrites of L5 cortical PNs using voltage dyes, electrophysiology and glutamate uncaging (Popovic et al., 2015; Acker et al., 2016; see Williams and Stuart, 2003; Nevian et al., 2007; Larkum et al., 2009 for similar measurements in other systems). ΔV_{max} scaled with ePSD area in all compartments (Figure 6C).

To determine the contribution of morphological parameters to the variance of ΔV_{max}, we used a generalized linear model (GLM) (Pedregosa et al., 2011). We analyzed the contribution of the volume, diameter and resistance of spine necks and heads as well as the contribution of ePSD area and distance between spine and soma (L_{dend}) to the amplitude of the signals in the soma and in spine heads. In the soma, ΔV_{max} was mainly determined by A_{ePSD}, which accounted for 89% of its variance when EPSPs were elicited in SiSs (77% when they came from DiSs). The second determinant was L_{dend}, which accounted for 6.4% of the variance of ΔV_{max} for EPSPs generated in SiSs (14.7% in DiSs). The contribution of R_{neck} to the variance of ΔV_{max} in the soma was comparatively negligible (Supplementary table 3), indicating that the passive attenuation of EPSPs along dendrites dominates the contribution of R_{neck} to somatic depolarizations evoked in spines. In the heads of SiSs, A_{ePSD} and R_{neck} accounted for 60% and 19% of the variance of ΔV_{max}, respectively. In the heads of DiSs, the contribution of R_{neck} to ΔV_{max} was much higher, reaching 38% of the variance, while A_{ePSD} contribution dropped to 47% (Supplementary figure 5 and Supplementary table 3). In 56% of dendritic spines, R_{neck} was large enough (>145MΩ) to attenuate EPSP amplitude by >50% across the spine neck, and more than 90% of spine necks attenuated the signal by at least 10% (Figure 6D), suggesting that most spine necks constitutively compartmentalize electrical signals in the head of spines.

We also estimated the elevation of calcium ion concentration ($\Delta[Ca^{2+}]$) in spine heads
induced by an EPSP. $\Delta[Ca^{2+}]$ was similar in SiSs and DiSs and varied non-linearly with A_{ePSD}
(Figure 6E). A_{ePSD} was the main determinant of $\Delta[Ca^{2+}]$, accounting for 30% of the variance in
SiSs (45% in DiSs), followed by R_{neck} (9%; Supplementary table 3). As a single EPSP is not
sufficient to elicit a Ca^{2+} spike, we did not model Ca^{2+} transients outside of spine heads.
Overall, our model provides quantitative insights into the variability of EPSP amplitude
originating from spine diversity and highlights differences in the contribution of morphological
parameters to spine depolarization and calcium signals in DiSs and SiSs.

Spatial interplay of excitatory and inhibitory signals
We used our model to compare the effects of spinous and dendritic shaft inhibition. *In vitro*
uncaging experiments have shown that inhibitory contacts located on DiSs could weaken local
calcium signals (Chiu et al., 2013), but the consequences on synaptic excitation are still
unclear (Kubota et al., 2015; Higley et al., 2014). To understand how spine ultrastructure and
iPSD location influence synaptic integration, we modeled the interaction between one IPSP
and one EPSP under the constraint of our morphological measurements. Assessing the extent
of signal variability originating from spine morphological heterogeneity requires a large number
of simulations (N≥1000). Therefore, we used a bootstrapping method (Efron and Tibshirani,
1994) (see Methods) to derive N≥1000 sets of parameters from our dataset of 390 spines and
62 shaft iPSDs (Supplementary file 1), and provide unbiased estimations of the mean and
variance of the signals. Importantly, the strength of inhibition depends on the reversal potential
of chloride ions (E_{Cl^-}). In healthy mature layer 2/3 cortical PNs, E_{Cl^-} typically varies between
the resting membrane potential, V_{rest} = -70 mV, and hyperpolarized values (-80 mV) (Doyon
et al., 2016). When E_{Cl^-} = -70 mV, active inhibitory synapses generate a local increase of
membrane conductance, which "shunts" membrane depolarization induced by concomitant
EPSPs. When E_{Cl^-} < -70 mV, the driving force of Cl^- ions is stronger, GABAergic inputs can
hyperpolarize the cell membrane and IPSPs counter EPSPs. These two situations are
respectively termed "shunting inhibition" and "hyperpolarizing inhibition" (Gidon and Segev,
2012; Doyon et al., 2016).
We first assessed the impact of "shunting" IPSPs elicited in the dendritic shaft on
individual EPSPs depending on the inter-synaptic distance (Figure 7A). For each iteration of
the model (i.e. N=3700 sets of realistic morphological values for spine size, and iPSD location
and size), we varied the distance Δx between a spine receiving an EPSP and a shaft iPSD
activated simultaneously. For Δx>0, the iPSD was located between the spine and the soma
(i.e. "on-path" inhibition, from the viewpoint of the soma), and for Δx<0, the iPSD was located
distally to the spine (i.e. "off-path" inhibition). To quantify inhibition, we compared the amplitude

of individual EPSPs in the absence $(\Delta V_{max,E})$ or presence $(\Delta V_{max,E+I})$ of inhibition, and

computed the drop in voltage $inh_V = 1 - \Delta V_{max,E+I}/\Delta V_{max,E}$. $Inh_V = 0$ indicates that the

electrical signal was not affected, $inh_V = 1$ indicates that it was completely inhibited.

Importantly, due to the electrical properties of the cell membrane, inh_V depends on where the

signal is measured. In the soma, inh_V was maximal (20% in average) when the iPSD was

located on-path and it decreased exponentially with Δx when the iPSD was located off-path

(exponential decay length: L$_{soma,off}$ = 30 μm) (Figure 7B), highlighting the impact of proximal

inhibitory synapses on distal excitatory inputs (Hao et al., 2009; Gidon and Segev, 2012). In

the spine where the EPSP was elicited, inh_V decayed exponentially with Δx for both on-path

and off-path inhibition (Figure 7C), with respective decay lengths L$_{spine,on}$ = 30 µm and L$_{spine,off}$

= 11 µm (also see Supplementary figure 6 for the dependence of decay lengths on the

dendritic cross-section). Therefore, shaft inhibition could affect excitatory signals within 40 µm,

which corresponds to approximately 50 spines considering the spine density and dendritic

diameter that we measured in basal dendrites of L2/3 cortical PN.

Next, we focused on DiSs. We modeled N=3700 DiSs and shaft iPSDs (Δx = 0) at

random locations on the dendrite, and we compared two configurations per iteration of the

model: (1) the ePSD of the DiS and the shaft iPSD were activated simultaneously (Figure

8A1); (2) both the ePSD and the iPSD of the DiS were activated simultaneously (Figure 8A2).

In the soma, the effect of shaft and spinous inhibition was comparable: inh_V was centered on

17%, and reached up to 50% (Figure 8B), in line with somatic recordings following coincident

uncaging of glutamate and GABA in acute brain slices (Chiu et al., 2013). By contrast, in the

head of all DiSs, spinous inhibition was more efficient than shaft inhibition despite the smaller

size of spinous iPSDs compared to shaft iPSDs. Spinous inh_V was centered on 10% and

reached up to 35%, whereas dendritic inh_V was centered on 3% and did not reach more than

23% (Figure 8C). These results support the notion that the placement of inhibitory synapses

structures the detection and integration of excitatory signals (Bono et al., 2017; Boivin and

Nedivi, 2018; Chiu et al., 2019), and highlight the role of spinous inhibition in local synaptic

signaling.

We then run the simulations with a lower chloride reversal potential (E$_{Cl-}$ = -80 mV) for

which GABAergic inputs can hyperpolarize the cell membrane. In numerous simulations, we

observed that hyperpolarization could take over depolarization (i.e. inh_V > 1) (Supplementary

figure 7A). This was the case in the soma for 37% of simulations, and in the spine where the

EPSP was generated for 10% of simulations. With E$_{Cl-}$ = -80 mV, the median inh_V imposed

by spinous inhibition in spine heads was 45%, in line with a previous morphologically

constrained model (Kubota et al., 2015) and with the fact that a hyperpolarizing IPSP may

block multiple EPSPs depending on iPSD placement (Boivin and Nedivi, 2018). Interestingly,

lowering E_{Cl^-} affected spinous and shaft inhibition differently. In the soma, shaft inhibition became more efficient than spinous inhibition (Supplementary figure 7B) due to the larger area of shaft iPSDs. However, in the heads of most DiSs, spinous inhibition remained more efficient than shaft inhibition (Supplementary figure 7C-F). Altogether, our results indicate that spinous inhibition is stronger than shaft inhibition in DiSs, and that their relative weight is modulated by the driving force of chloride ions.

Temporal interplay of excitatory and inhibitory signals

Since the efficacy of inhibition depends on the membrane potential at the onset of the IPSP (Koch et al., 1983; Segev and Rall, 1988; Hao et al., 2009; Doron et al., 2017), we addressed the effect of input timing on inhibition efficacy in DiSs. We simulated the interaction of one EPSP and one IPSP generated with a time difference of Δt (Figure 9A). For $\Delta t<0$ (IPSP before EPSP), IPSPs decreased EPSP amplitude (ΔV_{max}) (Figure 9B1). For $\Delta t>0$ (IPSP after EPSP), IPSPs had no effect on ΔV_{max} but abruptly decreased the tail of the EPSPs (Figure 9B2) (Koch et al., 1983). We first compared how spinous and shaft inhibition reduced EPSP duration by comparing the 80-to-20% decay time of the summed signals (τ_{E+I}) to that of uninhibited EPSPs (τ_E). Decay times were minimal at $\Delta t = +4$ ms and decreased by 64% and 78% with spinous and shaft inhibition respectively (Figure 9C), which could shorten the integration window of EPSPs and increase the temporal precision of synaptic transmission (Gabernet et al., 2005). The variance of τ_{E+I} was mainly determined by the area of iPSDs (Supplementary table 4). We then quantified how the timing of inhibition affected EPSP amplitude using $inh_V(\Delta t) = 1 - \Delta V_{max,E+I}/\Delta V_{max,E}$. In spine heads, inh_V was an asymmetrical function of Δt (Koch et al., 1983, Segev and Rall, 1988) and it was maximal at $\Delta t = -4$ ms for spinous inhibition, and at $\Delta t = -6$ ms for dendritic shaft inhibition. Overall, spinous inhibition was stronger than shaft inhibition, decreasing ΔV_{max} by 26.3% and 16.2%, respectively (median values in Figure 9D; see also Supplementary figure 8A for $inh_V(\Delta t)$ with higher E_{Cl^-}). Interestingly, R_{neck} had a negligible contribution to the variance of inh_V in the case of hyperpolarizing inhibition, but a major one (37%) in the case of shunting inhibition (Supplementary table 4), suggesting that spine necks compartmentalize IPSPs differently depending on E_{Cl^-}.

Then, we examined the effect of timed inhibition on calcium signaling in spines, using $inh_{[Ca^{2+}]}(\Delta t) = 1 - \Delta[Ca^{2+}]_{max,E+I} / \Delta[Ca^{2+}]_{max,E}$. We observed that $inh_{[Ca^{2+}]}$ peaked at $\Delta t = 0$ ms for both spinous and shaft inhibition. More precisely, spinous inhibition reduced calcium transient amplitude by 10% in average, reaching >36% in the top 10% simulations, while shaft inhibition reduced it by 8.6% in average and >28% in the top 10% simulations (Figure 9E). These values are in the range of $inh_{[Ca^{2+}]}$ measured with double uncaging experiments (Chiu

457 et al., 2013). Importantly, IPSPs could decrease the amplitude of calcium transients, within a
458 short time-window (Δt between 0 and +10 ms) in which depolarization amplitude was not
459 affected (Figure 9D-E), thereby decoupling calcium signalling from electrical activity in DiSs.
460

DISCUSSION

In the present study, we developed a novel 3D-CLEM workflow allowing the ultrastructural
characterization of specific populations of dendritic spines in genetically defined types of
neurons. We used this workflow to exhaustively reconstruct spines and synaptic contacts
along the basal dendrites of fluorescently labelled L2/3 cortical PNs of the SSC and to provide
a quantitative description of their diversity. We input our measurements in a computational
model to analyze the variability of electrical and calcium synaptic signals originating from spine
ultrastructural diversity, and to characterize the spatio-temporal integration of excitatory and
inhibitory inputs. Our results shed light on unique properties of DiSs, which represent 10% of
all spines and 38% of all inhibitory synapses along the basal dendrites of L2/3 cortical PNs.
We show that while individual inhibitory synapses distributed along dendritic shafts can be
powerful enough to block several EPSPs, spinous inhibitory synapses affect excitatory signals
more efficiently in DiSs. We also show that the activation of a spinous inhibitory synapse within
a few milliseconds after an EPSP can decouple voltage and calcium signals in DiSs, which
could impact calcium-dependent signaling cascades that drive spine plasticity.
The molecular composition of synapses and their biophysical properties are reportedly
heterogeneous along dendrites and across dendritic trees (Yuste et al., 2000; Berglund et al.,
2008; Losonczy et al., 2008; Stuart and Spruston, 2015). However, most computational
models so far have used *ad hoc* or averaged values as parameters for dendritic spines and
excitatory synapses (Sarid et al., 2007; Almog and Korngreen, 2014; Müllner et al., 2015), and
considered that all inhibitory synapses were located along the dendritic shaft (Markram et al.,
2015). This limits their accuracy (Jadi et al., 2014). Hence, there is a critical need for cell type-
and dendritic type-specific measurements of synaptic ultrastructure and placement, and for
detailed quantification of synaptic diversity beyond the µm scale in intact brain circuits. The
correlative workflow we propose provides a solution to such demand. It is applicable to any
type of tissue and allows anatomical measurements of any kind of genetically labelled cells
and organelles. One technical limitation is the need for chemical fixation, which may distort
tissue morphology (Korogod et al., 2015; Rostaing et al., 2006). Therefore, it may be
necessary to correct for tissue shrinkage based on a morphological comparison with physically
fixed tissues (Supplementary figure 4B3) in order to reliably depict *in vivo* situations. Future
development of aldehyde-free cryo-CLEM methods will be important to grant access to cellular
and synaptic ultrastructure in close-to-native environments.

Applying 3D-CLEM to the basal dendrites of L2/3 cortical PNs allowed us to
quantitatively describe the landscape of synaptic diversity and to characterize the
ultrastructural features of a scarce population of dendritic spines receiving both excitatory and
inhibitory synaptic inputs (DiSs). In the cortex, DiSs are mostly contacted by VGluT2-positive
excitatory thalamo-cortical inputs (Kubota et al., 2007) and they receive inhibition from
somatostatin-expressing and parvalbumin-expressing interneurons (Kubota et al., 2016; Chiu
et al., 2013), which are the two main sources of inhibitory inputs to the basal dendrites of layer
2/3 cortical PNs (Tremblay et al., 2016; Kuljis et al., 2019). *In vivo* 2-photon imaging
experiments have shown that DiSs are among the most stable spines along the dendrites of
layer 2/3 PNs (Villa et al., 2016). The inhibitory synapse in DiSs is smaller and more labile
than inhibitory synapses along dendritic shafts, and it is very sensitive to sensory experience
(Knott et al., 2002; Chen et al., 2012; van Versendaal et al., 2012; Villa et al., 2016). Whisker
stimulation induces a lasting increase in the occurrence of iPDSs in spines of the barrel cortex
(Knott et al., 2002) and monocular deprivation destabilizes iPSDs housed in spines of the
visual cortex (Chen et al., 2012; van Versendaal et al., 2012; Villa et al., 2016), suggesting
their role in experience-dependent plasticity. Our morphological and computational analysis
provides new insights into the biophysical properties of DiSs. We show that DiSs have larger
heads and larger ePSDs than SiSs, and most often contain a spine apparatus. However, the
ratio between mean spine neck diameter and spine head volume (or ePSD area) was smaller
in DiSs than in SiSs, and DiSs had longer necks than SiSs of comparable head volume, so
that EPSPs of similar amplitudes encounter a higher neck resistance in DiSs than in SiSs.
Thus, DiSs are uniquely compartmentalized by their ultrastructural features and the presence
of an inhibitory synapse.
Our model predicts that IPSPs occurring in DiSs within milliseconds after an EPSP can
curtail calcium transients without affecting depolarization, thereby locally decoupling voltage
and calcium signaling. This is expected to impact the induction of long-term forms of synaptic
plasticity, such as long-term potentiation (LTP) or long-term depression (LTD), which underlie
learning and memory (Bourne and Harris, 2008; Nabavi et al., 2014; Hayashi-Takagi et al.,
2015; Segal, 2017). The induction of LTP versus LTD is determined by the magnitude and
time course of calcium flux, with brief, high calcium elevation generating LTP, sustained
moderate calcium elevation generating LTD, and low calcium level inducing no plasticity
(Lisman, 1989; Feldman, 2012; Graupner and Brunel, 2012). Therefore, a small reduction in
the amplitude of calcium transients may limit spine potentiation, or even cause depression
(Hayama et al., 2013; Higley, 2014; Nakahata and Yasuda, 2018). In the cortex,
thalamocortical inputs may contact DiSs on the basal dendrites of L2/3 both directly (excitatory
connection) and indirectly through feed-forward inhibition *via* parvalbumin-expressing fast-
spiking interneurons (Gabernet et al., 2005; Kimura et al., 2010). The delay between thalamo-

cortical excitatory and feed-forward inhibitory signals is typically +1 ms to +3 ms (Gabernet et

al., 2005), within the 10 ms time window for voltage-calcium decoupling in DiSs. Therefore,

the presence of inhibitory synapses in DiSs could prevent synaptic potentiation and thereby

increase the temporal precision of cortical response to sensory stimulation (Knott et al., 2002;

Wehr and Zador, 2003; Gabernet et al., 2005). On the contrary, the removal of spine inhibitory

synapses could favor synaptic potentiation during experience-dependent plasticity such as

monocular deprivation to strengthen inputs from the non-deprived eye (Frenkel et al., 2006;

Chen et al., 2012; van Versendaal et al., 2012).

Our understanding of synaptic and dendritic computations is intimately linked to the

quantitative description of synaptic distribution, ultrastructure, nano-organization, activity and

diversity in neural circuits. The CLEM workflow we propose opens new avenues for the

ultrastructural characterization of synapses with defined molecular signature characterizing

their identity or activation profile in response to certain stimuli or behaviors. Another milestone

to better model the biophysics of synaptic integration will be to combine EM and quantitative

super-resolution LM to measure the density and nano-organization of molecular species (e.g.

AMPARs, NMDARs, voltage-dependent calcium channels) in specific populations of synapses

in intact brain circuits. Combining circuit and super-resolution approaches through CLEM will

be critical to refine large-scale circuit models (Markram et al., 2015; Sarid et al., 2015; Bhatia

et al., 2019; but see Almog and Korngreen, 2016) and bridge the gap between molecular,

system and computational neurosciences.

MATERIALS AND METHODS

Animals and *in utero* cortical electroporation

All animals were handled according to French and EU regulations (APAFIS#1530-

201508261150869 1v3). *In utero* cortical electroporation was performed as described

previously (Fossati et al., 2019). Briefly, pregnant Swiss female mice at E15.5 (Janvier Labs,

France) were anesthetized with isoflurane (3.5% for induction, 2% during the surgery) and

subcutaneously injected with 0.1 mg/kg of buprenorphine for analgesia. The uterine horns

were exposed after laparotomy. Electroporation was performed using a square wave

electroporator (ECM 830, BTX) and tweezer-type platinum disc electrodes (5mm-diameter,

Sonidel). The electroporation settings were: 4 pulses of 40 V for 50 ms with 500 ms interval.

Endotoxin-free DNA was injected using a glass pipette into one ventricle of the mouse

embryos at the following concentrations: pH1SCV2 TdTomato: 0.5 µg/µL and pCAG EGFP-

GPHN: 0.3 µg/µL. All constructs have been described before (Fossati et al., 2016).

Cortical slice preparation

Electroporated animals aged between postnatal day P84 and P129 were anesthetized with
ketamin 100 mg/kg and xylazin 10 mg/kg, and intracardiacally perfused with first 0.1 mL of
heparin (5000 U.I/mL, SANOFI), then an aqueous solution of 4% w/v paraformaldehyde (PFA)
(Clinisciences) and 0.5% glutaraldehyde (GA) (Clinisciences) in 0.1 M phosphate-buffered
saline (PBS). The fixative solution was made extemporaneously, and kept at ice-cold
temperature throughout the perfusion. The perfusion was gravity-driven at a flow rate of about
0.2 ml/s, and the total perfused volume was about 100 ml per animal. Brains were collected
and post-fixed overnight at 4°C in a 4% PFA solution. 30 μm-thick coronal brain sections were
obtained using a vibrating microtome (Leica VT1200S).
575

Fluorescence microscopy of fixed tissue
Slices containing electroporated neurons were trimmed to small (5-10 mm^2) pieces centered
on a relatively isolated fluorescent neuron, then mounted in a custom-made chamber on #1.5
glass coverslips. The mounting procedure consisted in enclosing the slices between the glass
coverslip and the bottom of a cell culture insert (Falcon, ref. 353095) adapted to the flat
surface with a silicon O-ring gasket (Leica) and fixed with fast-curing silicon glue
(Supplementary figure 1A). Volumes of GFP and tdTomato signals were acquired in 12 bits
mode (1024x1024 pixels) with z-steps of 400 nm using an inverted Leica TCS SP8 confocal
laser scanning microscope equipped with a tunable white laser and hybrid detectors and
controlled by the LAF AS software. The objective lenses were a 10X PlanApo, NA 0.45 lens
for identifying electroporated neurons and a 100X HC-PL APO, NA 1.44 CORR CS lens
(Leica) for higher magnification images. GFP-GPHN puncta with a peak signal intensity at
least four times above shot noise background levels were considered for CLEM.
589

Placement of DAB fiducial landmarks
Following confocal imaging, slices were immersed in a solution of 1 mg/mL 3,3'-
diaminobenzidine tetrahydrochloride (DAB, Sigma Aldrich) in Tris buffer (0.05 M, pH 7.4). The
plugin "LAS X FRAP" (Leica) was used to focus the pulsed laser in the tissue in custom
patterns of 10-to-20 points using 100% power in 4 wavelengths (470 to 494nm) for 30s-60s
per point at 3 different depths: the top of the slice, the depth of the targeted soma, then the
bottom of the slice (surface closest to the objective). DAB precipitates were imaged in
transmitted light mode. Slices were subsequently rinsed twice in Tris buffer and prepared for
electron microscopy.
599

Tissue preparation for serial block-face scanning electron microscopy (SBEM)
Using a scalpel blade under a M165FC stereomicroscope (Leica), imaged tissue slices were
cut to ~1mm^2 asymmetrical pieces of tissue centered on the ROI, and then kept in plastic

baskets (Leica) through the osmification and dehydration steps. Samples were treated using
an osmium bridging technique adapted from the NCMIR protocol (OTO) (Deerinck et al.,
2010). The samples were washed 3 times in ddH_2O and immersed for 1 hour in a reduced
osmium solution containing 2% osmium tetroxide and 1.5% potassium ferrocyanide in ddH_2O.
Samples were then immersed for 20 minutes in a 1% thiocarbohydrazide (TCH) solution
(Electron Microscopy Science) prepared in ddH_2O at room temperature. The samples were
then post-fixed with 2% OsO_4 in ddH_2O for 30 minutes at room temperature and colored *en*
bloc with 1% aqueous uranyl acetate at 4 °C during 12 hours. Post-fixed samples were
subjected to Walton's *en bloc* lead aspartate staining at 60 °C for 30 minutes (Walton, 1979).
After dehydration in graded concentrations of ice-cold ethanol solutions (20%, 50%, 70%,
90% and twice 100%, 5 minutes per step) the samples were rinsed twice for 10 minutes in
ice-cold anhydrous acetone. Samples were then infiltrated at room temperature with graded
concentrations of Durcupan (EMS) prepared without plastifier (components A, B, C only). In
detail, blocks were infiltrated with 25% Durcupan for 30 minutes, 50% Durcupan for 30
minutes, 75% Durcupan for 2 hours, 100% Durcupan overnight, and 100% fresh Durcupan
for 2 hours before being polymerized in a minimal amount of resin in a flat orientation in a
sandwich of ACLAR® 33C Films (EMS) at 60 °C for 48 hours. Samples were mounted on
aluminum pins using conductive colloidal silver glue (EMS). Before curing, tissue blocks were
pressed parallel to the pin surface using a modified glass knife with 0° clearance angle on an
ultramicrotome (Ultracut UC7, Leica), in order to minimize the angular mismatch between LM
and SEM imaging planes. Pins then cured overnight at 60°C. Samples were then trimmed
around the ROI with the help of fluorescent overviews of the ROI within their asymmetrical
shape. Minimal surfacing ensured that superficial DAB landmarks were detected at the SBEM
before block-facing.
627

SBEM acquisition
SBEM imaging was performed with a Teneo VS microscope (FEI) on the ImagoSeine imaging
platform at Institut Jacques Monod, Paris. The software MAPS (Leica) was used to acquire
SEM images of targeted volumes at various magnifications. Acquisition parameters were:
1,7830 kV, 500 ns/px, 100 pA, 40 nm-thick sectioning and 8200x8200 pixels resolution with
either 2.5 nm or 25 nm pixel size for high- and low-magnification images, respectively. Placing
an electromagnetic trap above the diamond knife to catch discarded tissue sections during
days-long imaging sessions was instrumental to achieve continuous 3DEM acquisitions.
636

Image segmentation
Dendrites were segmented from SBEM stacks using the software Microscopy Image Browser
(MIB) (Belevich et al., 2016). 3D reconstruction was performed with the software IMOD

(Kremer et al., 1996; http://bio3d.colorado.edu/imod/). 3D spine models were imported in the
software Blender (www.blender.org) for subsampling and the quantification of spine section
areas along their main axis was done with in-house python scripts. Other measurements were
performed using IMOD and in-house python scripts.
644

Tissue preparation for tissue shrinkage estimation
Two female mice (21 days postnatal) were used for the analysis of tissue shrinkage induced
by chemical fixation. Mice were decapitated and their brains were rapidly removed. The brains
were transferred to an ice-cold dissection medium, containing (in mM): KCl, 2.5; $NaHCO_3$, 25;
NaH_2PO_4, 1; $MgSO_4$, 8; glucose, 10, at pH 7.4. A mix of 95% O_2 and 5% CO_2 was bubbled
through the medium for 30 min before use. 300-µm-thick coronal brain sections were obtained
using a vibrating microtome (Leica VT1200S). Small fragments of the SSC were cut from
those slices and fixed either by immersion in an ice-cold PBS solution containing 4% PFA and
0.5% GA, or in frozen with liquid nitrogen under a pressure of 2100 bars using a high pressure
freezing system (HPM100, Leica). For HPF-frozen samples, the interval between removal of
the brain and vitrification was about 7 min. Cryo-substitution and tissue embedding were
performed in a Reichert AFS apparatus (Leica). Cryo-substitution was performed in acetone
containing 0.1% tannic acid at -90°C for 4 days with one change of solution, then in acetone
containing 2% osmium during the last 7h at -90°C. Samples were thawed slowly (5°C/h) to -
20°C and maintained at -20°C for 16 additional hours, then thawed to 4°C (10°C/h). At 4°C
the slices were immediately washed in pure acetone. Samples were rinsed several times in
acetone, then warmed to room temperature and incubated in 50% acetone-50% araldite
epoxy resin for 1h, followed by 10% acetone-90% araldite for 2h. Samples were then
incubated twice in araldite for 2h before hardening at 60°C for 48h. As for chemically fixed
sections, they were post-fixed for 30 min in ice-cold 2% osmium solution, rinsed in PBS buffer,
dehydrated in graded ice-cold ethanol solutions and rinsed twice in ice-cold acetone, before
undergoing the same resin infiltration and embedding steps as HPF-frozen samples. After
embedding, ultrathin sections were cut in L2/3 of the SSC, orthogonally to the apical dendrites
of pyramidal neurons, 200-300 µm from the pial surface using an ultramicrotome (Ultracut
UC7, Leica). Ultra-thin (pale yellow) sections were collected on formwar-coated nickel slot
grids, then counterstained with 5% uranyl acetate in 70% methanol for 10 min, washed in
distilled water and air dried before observation on a Philips TECNAI 12 electron microscope
(Thermo Fisher Scientific).
673

Measurement of shrinkage correction factors
Ultra-thin sections of both HPF-frozen tissues and chemically-fixed tissues were observed
using a Philips TECNAI 12 electron microscope (TFS). Cellular compartments contacted by a

pre-synaptic bouton containing synaptic vesicles and exhibiting a visible electron-dense PSD

at the contact site, but no mitochondrion within their cytosol were identified as dendritic spine

heads. Cross-section areas of random spine heads and the curvilinear lengths of their PSD

were quantified in both conditions using the softwares MIB and IMOD. N = 277 spine head

sections were segmented in HPF-frozen cortical slices from two female mice, and N = 371

spine head sections were segmented in chemically fixed cortical slices originating from the

same two mice. Chi-square minimization was used between spine head cross-section area

distributions in HPF or OTO conditions to compute average volume shrinkage and correction

factors. PSD areas were not corrected as they exhibited no shrinkage.

Computation of the diffusional neck resistance

The diffusional resistance of spine necks W_{neck} was measured as follows. Using IMOD, we

first modeled in 3D the principal axis of each spine neck as an open contour of total length

L_{axis} connecting the base of the neck to the base of the spine head. Using Blender, we

interpolated each spine neck path linearly with 100 points. We named P(l) the plane that

bisected the spine neck model orthogonally to the path at the abscissa l, and A(l) the spine

neck cross-section within P(l). In spines containing a spine apparatus (SA), we corrected A(l)

by a scaling factor $\beta(l) = 1-(D_{SA}/D_{spine})^2(l)$, where $D_{SA}/D_{spine}(l)$ is the local ratio of SA and neck

diameter. We measured D_{SA}/D_{spine} orthogonally to the neck path in 10 SA+ spines and in three

different locations per spine on SBEM images: at the spine stem (l/L_{axis} = 0.1), at the center of

the spine neck (l/L_{axis} = 0.5), and at the stem of the head (l/L_{axis} = 0.9). D_{SA}/D_{spine} was

44% ± 11%, 31% ± 8% and 37% ± 8% respectively, and fluctuations were not statistically

significant. We then divided each SA+ spine neck in thirds and scaled their neck cross-section

areas along neck axis $A_{SA+}(l) = \beta(l)A(l)$ before computing $W_{neck} = \int dl/A(l)$ for all spines, using

Simpson's integration rule.

Multi-compartment electrical model

All simulations were implemented in Python using NEURON libraries (Hines and Carnevale,

2001) and in-house scripts. Ordinary differential equations were solved with NEURON-default

backward Euler method, with Δt = 0.05 ms. Scripts and model definition files are available in

a GitHub repository: https://github.com/pabloserna/SpineModel. Biophysical constants were

taken from the literature as follows: membrane capacitance C_m = 1μF/cm^2 (Doron et al., 2017);

cytosolic resistivity ρ = 300 Ω.cm (Miceli et al., 2017; Barbour, 2017); synaptic conductivities

were modeled as bi-exponential functions $g(t) = Ag_{max}(e^{-t/t_2} - e^{-t/t_1})$ where A is a

normalizing constant and t_1 and t_2 define the kinetics of the synapses: GABAergic

conductance (t_1, t_2) = (0.5, 15) ms, AMPAR-dependent conductance (t_1, t_2) = (0.1, 1.8) ms,

NMDAR-dependent conductance (t_1, t_2) = (0.5, 17.0) ms (ModelDB:

https://senselab.med.yale.edu/ModelDB/). The magnesium block of NMDA receptors was

modeled by a voltage-dependent factor (Jahr and Stevens, 1990). Remaining free parameters

comprised: the leaking conductivity g_m (or, equivalently, the membrane time constant T_m); the

peak synaptic conductance per area: g_{AMPA}, g_{NMDA}, g_{GABA}; the total membrane area of the

modeled neuron. These parameters were adjusted so that signal distributions fitted published

electrophysiological recordings (Feldmeyer et al., 2006; Kobayashi et al., 2008; Hull et al.,

2009; Avermann et al., 2012; Scala et al., 2018). In more detail, we first set up one "ball-and-

stick" model per segmented spine (N=390). The dendrite hosting the modeled spine was

generated as a tube of diameter $d_{dendrite}$ = 0.87 µm, and length $L_{dendrite}$ = 140 µm. This dendrite

is split in three parts, the 2 µm-long middle one harboring the modeled spine. To account for

the passive electrical effects of neighboring spines, the membrane surfaces of both the

proximal and distal sections of the studied dendrite were scaled by a correction factor alpha =

1 + <A_{spine}>d_{spine}/$\pi d_{dendrite}$ = 3.34, with the density d_{spine} =1.63 spine.µm^{-1} and the average spine

membrane area <A_{sp}> = 3.89 µm^2 . We calibrated synaptic conductances type by type, by

fitting the signals generated in the whole distribution of 390 models to published

electrophysiological recordings. The AMPA conductances of all excitatory synapses were set

proportional to ePSD area and scaled by the free parameter g_A. In each model, we activated

the AMPAR component of excitatory synapses and monitored the amplitude of resulting

EPSCs in the soma. The average EPSC amplitude was adjusted to 58 pA (Feldmeyer et al.,

2006; Hull et al., 2009), yielding a scaling factor g_A = 3.15 nS/µm^2, which takes into account

the average number of excitatory contacts per axon per PN in L2/3 of mouse SSC: $N_{ePSD/axon}$=

2.8 (Feldmeyer et al., 2006). The average AMPA synaptic conductance was 0.42 nS. The

leakage resistance was fitted to 65 MΩ (Hoffmann et al., 2015), yielding a total membrane

surface of the modeled neurons: $A_{mb, total}$ = 18550 µm^2. The NMDA conductances of all

excitatory synapses were set proportional to ePSD area and scaled by the free parameter g_N.

In each model, we activated both NMDA and AMPA components of excitatory synapses and

fitted the amplitude ratio between the average AMPA+NMDA and AMPA-only responses to

1.05 (Hull et al., 2009), yielding g_N = 3.4 nS/µm^2. The GABA conductances of all inhibitory

synapses were set proportional to iPSD area and scaled by the free parameter g_G. In this

case, we set the neuron to a holding potential of 0 mV and the reversal potential of chloride

ions (E_{Cl-}) to -80 mV. Then, we activated shaft inhibitory synapses and monitored the

amplitude of resulting IPSCs in the soma. The amplitude of the average GABAergic

conductance was set to 1 nS (Doron et al., 2017; Hoffmann et al., 2015; Xu et al., 2013),

yielding a scaling factor g_G = 5.9 nS/µm^2, which takes into account the average number of

inhibitory contacts per axon per PN in L2/3 of mouse SSC: $N_{iPSD/axon}$ = 6 (Hoffmann et al.,

2015). Considering inhibition, since E_{Cl-} is regulated on timescales exceeding 100 ms (Boffi et

al., 2018) and we modeled signals in the 10 ms timescale, we could set E_{Cl^-} as a constant

parameter of our steady state model. Calcium influxes were modeled in spines as a result of

the opening of NMDARs and voltage-dependent calcium channels (VDCCs). Since we

simulated signals that remained below the threshold for eliciting dendritic spikes (Sarid et al.,

2015; Stuart and Spruston, 2015), we did not include VDCCs in dendrites, and monitored

calcium transients exclusively in spine heads. The dynamics of L-, N- and Q-type VDCCs were

obtained from ModelDB (accession n°: 151458), and their conductivities were scaled to the

head membrane area of each spine, A_{head}, excluding synaptic area(s). VDCC-type ratios and

calcium conductivities were adjusted by fitting the average amplitude of calcium concentration

transients to 20% of the NMDA conductance (Bloodgood et al., 2009). Calcium uptake from

cytosolic buffers was set to 95% to yield an average amplitude of Ca^{2+} concentration transients

of 0.7 µM (Sabatini et al, 2002).

Bootstrapping

To simulate a large number of spine-spine interactions with limited redundancy, our

distribution of spines was expanded using a "smooth" bootstrapping method (Efron and

Tibshirani, 1994). Specifically, the dataset (i.e. a matrix of dimensions N x N_f) was re-sampled

to generate a new matrix of dimension M x N_f, where N is the number of spines, N_f is the

number of selected features, and M is the final number of synthetic spines. M rows were

randomly selected in the original dataset and zero-centered, feature-dependent Gaussian

noise was added to each element of the matrix (excluding absolute quantities, e.g. number of

PSDs or presence of SA). To determine appropriate noise amplitude for each parameter, a

synthetic set of M=500 spines was generated from the original dataset, including Gaussian

noise with an arbitrary amplitude σ, on one selected parameter. This new feature distribution

was compared to the original distribution using a 2-sample Kolmogorov-Smirnov test (KS-

test), and this procedure was repeated 1000 times for each set value of σ. A conservative

noise level (σ = 10%) was sufficient to smear parameter distributions while the fraction of

synthetic sets that were statistically different from the original set ($p < 0.05$, KS-test) remained

0 over 1000 iterations. σ = 10% was valid for all relevant features, and we assumed that such

a small noise amplitude would minimally interfere with non-linear correlations in our dataset.

Synthetically generated spines were then used to simulate elementary synaptic signaling

using in-house python scripts. We also used bootstrapping to estimate standard deviations in

our simulations.

Statistics

785 No statistical methods were used to predetermine sample size. We used a one-way ANOVA

786 on our 4 datasets (Supplementary table 1) to test that inter-neuron and inter-mice variability

were small enough to pool all datasets together (Supplementary table 2). We used Kolmogorov-Smirnov test to determine that all measured morphological parameters followed a log-normal distribution (Supplementary table 2). We used Mann–Whitney U test for statistical analyses of morphological parameters, except when comparing the probability for SiSs and DiSs to harbor SA, for which we used Pearson's χ^2 test. All results in the text are mean ± SD. In figures 6, 7 and 9, we plot medians as solid lines, as they better describe where log-normal distributions peak. Shaded areas represent 68% confidence intervals, which span approximately one standard deviation on each side of the mean.

ACKNOWLEDGEMENTS

We thank members of the Triller and Charrier laboratories (IBENS, Paris, France), Vincent Hakim and Boris Barbour for insightful discussions. This work was supported by INSERM, the Agence Nationale de la Recherche (ANR-13-PDOC-0003 and ANR-17-ERC3-0009 to C.C.), the European Research Council (ERC starting grant 803704 to C.C.), the Labex Memolife (901/IBENS/LD09 to O.G.), the company NIKON France via the CIFRE program (convention n° 2015/1049 to O.G.) and the European Union's Horizon 2020 Framework Programme for Research and Innovation under the Specific Grant Agreement No. 785907 (Human Brain Project SGA2 to P.S.). We acknowledge the ImagoSeine imaging facility (Jean-Marc Verbavatz and Rémi Leborgne, Jacques Monod Institute, Paris, France) for SBEM availability, service and support. We thank Liesbeth Hekkings (ThermoFisher Scientific) for technical support. We are grateful to the IBENS Imaging Facility (France BioImaging, supported by ANR-10-INBS-04, ANR-10-LABX-54 MEMO LIFE, and ANR-11-IDEX-000-02 PSL* Research University, ''Investments for the future''; NERF 2011-45; FRM DGE 20111123023; and FRC Rotary International France).

AUTHOR CONTRIBUTIONS

O.G. developed the 3D-CLEM workflow, performed the experiments, segmented images and analyzed data. P.S. coded the model and analyzed data. N.A. and M.F. carried out the *in utero* electroporations. P.R. trained O.G. in EM and carried out part of the TEM imaging. O.G., P.R., A.T. and C.C. designed the study. O.G., P.S. and C.C. interpreted the results, prepared the figures, and wrote the manuscript. A.T. and C.C. provided funding.

COMPETING INTERESTS

The authors declare that no competing interests exist.

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FIGURES AND LEGENDS

Figure 1. CLEM imaging of identified spines within intact cortical circuits.

(A) Visualization of basal dendrites of a pyramidal neuron expressing cytosolic TdTomato in layers 2/3 (L2/3) of adult mouse somatosensory cortex. Diaminobenzidine (DAB) was photo-precipitated using focused UV light to insert correlative landmarks (pink dots in yellow circles).

(B) Transmitted light image of the same field of view after DAB photo-precipitation. DAB precipitates are highlighted with yellow circles. Blood vessels are outlined with purple dashed lines.

(C) Composite scanning EM (SEM) image displaying DAB patterning at the depth of the neuron of interest (yellow circles). Slight mismatch between LM and SEM observation planes resulted in DAB landmarks appearing in different z-planes during block-facing; the white line represents stitching between z-shifted images. In C1, landmarks are arranged as in B. C2 is a close-up on the soma of the electroporated neuron, labelled with three DAB landmarks (arrowheads). C3 corresponds to an orthogonal (x,z) view of the SEM stack along the green dashed line in C2. The superficial DAB layer enabled ROI targeting, and the deeper layer enabled retrospective identification of the target neuron.

(D) 3D-reconstruction of dendrites of interest from the overview SEM stack. DAB landmarks are reconstructed in blue (in yellow circles). The red rectangle outlines the portion of dendrite represented in E and F.

(E) Z-projection of the confocal stack corresponding to the portion of dendrite reconstructed in D. Letters identify individual spines.

(F) 3D-EM reconstruction. Individual dendritic spines are manually segmented and randomly colored. Spines that were detected in EM but not in LM are labelled in red.

Scale bars: A, B, C1, D: 10 µm; C2, C3: 5 µm; E, F: 2 µm.

Figure 2. Spine morphometry along basal dendrites of layer 2/3 cortical pyramidal neurons.

(A) 3D-reconstruction of a dendritic spine from a SBEM stack. Dendritic shaft is in light green, spine neck in turquoise, spine head in blue and PSD surface in red. The following parameters were measured : PSD area, head diameter, neck diameter and neck length. Scale bar: 300 nm.

(B) Linear correlation of PSD area and spine head volume. R^2=0.82.

(C) Plot of the minimal spine neck diameter as a function of spine neck length. Spearman correlation coefficient is -0.58.

(D) Neck length as a function of spine head orientation, as quantified by the ratio of spine head diameter (D_{head}) over its length (L_{head}). $D_{head}/L_{head} < 1$ corresponds to a prolate spine head, which shape is stretched longitudinally with respect to the neck. $D_{head}/L_{head} > 1$ characterizes an oblate spine head, oriented orthogonally to the neck. The dashed line corresponds to $D_{head}/L_{head} = 1$. We did not observe any oblate spine with a long neck (non-existent spine morphology).

Figure 3. Ultrastructural comparison of spines with and without a spine apparatus.

(A) TEM images of spines either devoid of spine apparatus (SA-, left) or containing a spine apparatus (SA+, right, yellow arrowhead). Scale bars: 500nm.

(B) Proportion of SA- and SA+ spines. Histogram represents mean ± SD (N=390 spines).

(C) Distribution of mean head diameter for SA- and SA+ spines. N=179 and 221, respectively (p < 10^{-38}).

(D) Distribution of ePSD area. (p < 10^{-40})

(E) Distribution of mean neck diameter. (p < 10^{-12})

(F) Probability of harboring a SA as a function of spine head volume. Blue: experimental data. Orange: sigmoid fit.

(G) Distribution of the diffusional resistance of the spine neck (W_{neck}) calculated based on neck morphology. (p < 10^{-5}).

 ***p<0.001 calculated using Mann-Whitney test.

Figure 4. Identification of excitatory and inhibitory synapse on DiSs using CLEM.

(A) Confocal image of basal dendrites of a cortical L2/3 PN that was electroporated with cytosolic TdTomato and GFP-GPHN to label inhibitory synapses. The magenta rectangle outlines the region enlarged in B.

(B) Enlargement of a portion of the dendrite in A harboring several dendritic spine (lettered). Spine "e" contains a cluster of GFP-GPHN (asterisk) and corresponds to a putative dually-innervated spine (DiS).

(C) 3D-EM reconstruction of the same dendritic fragment as in B. Dendritic shaft is colored in purple; individual spines and PSDs are colored randomly. Spines visible in EM but not in LM are labelled in red. The inhibitory PSD (colored in green) on spine "e" is identified based on the position of the GFP-GPHN cluster (asterisk in B and C). GFP-GPHN-negative PSDs are defined as excitatory.

(D) 3D-EM reconstruction of spine "e" (yellow) with its presynaptic partners (magenta and green). As the "green" axon also targets a neighboring dendritic shaft (blue), it is defined as inhibitory. Scale bars: A: 10 µm; B, C, D: 1 µm.

Figure 5. Anatomical properties of DiSs.

(A) Proportion of spines harboring 0, 1 or 2 synaptic contacts, quantified with CLEM. Histograms represent mean ± SD from, 390 spines in 8 dendrites.

(B) Quantification of mean spine head diameter for SiSs (blue) and DiSs (red). ($p < 10^{-4}$).

(C) Proportion of SiSs and DiSs harboring a SA. ($p < 10^{-10}$ using Pearson's χ^2 test)

(D-F) Quantification of neck length (D), the ratio between mean neck diameter and head volume (E) and the diffusional neck resistance (W_{neck}) (F) between SiSs and DiSs (solid lines,

N=349 and 37, respectively) and between DiSs with SiSs of similar head volume (spines with $V_{head}>0.05$ μm^3, dashed lines, N=186 and 34, respectively).

(G) Quantification of iPSD area in DiSs and dendritic shafts. N=37 and 62, respectively (p < 10^{-6}).

(H) Quantification of ePSD area in SiSs or DiSs. (p < 10^{-5}).

(I) Plot of iPSD area as a function of ePSD area in individual DiSs. The dashed line (y = x) highlights that the ePSD is larger than the iPSD in most of DiSs. N=37.

p-values were computed using Mann-Whitney test (B, D-H) or Pearson's χ^2 test (C). Only significant (p<0.05) p-values are shown (*p<0.05; **p<0.01; ***p<0.001).

Figure 6. Morphologically-constrained model of synaptic signaling.

(A) Schematic of the circuit model (A1) and representative time-course of excitatory (magenta) and inhibitory (green) conductance based on the kinetics of AMPA, NMDA and GABA$_A$ receptors (A2). All compartments include passive resistor-capacitor circuits to model cell membrane properties and optionally include an active conductance that models voltage-dependent currents (VDC). All modeled spines feature an excitatory synapse with glutamatergic AMPA and NMDA currents. Spines and dendritic compartments can also

feature an inhibitory synapse with GABAergic currents. All conductances were scaled to PSD area (see Methods).

(B) Simulation of the time-courses of membrane depolarization following an EPSP, taking into account spine diversity (i.e. R_{neck}, ePSD area and distance to soma, as measured in CLEM). Membrane voltage is monitored over time in the spine head (blue), in the dendritic shaft in front of the spine (orange) and in the soma (green). Median voltage transients are plotted as solid lines. Shaded areas represent 68% confidence intervals, which span approximately one standard deviation on each side of the mean.

(C) Amplitude of evoked depolarization (ΔV_{max}) as a function of ePSD area at three distinct locations: head of SiSs (blue) or DiSs (magenta) where the EPSP was elicited, dendritic shaft 1 μm from the spine (orange) or soma (green).

(D) Attenuation of the amplitude of depolarization between the spine head and the dendrite as a function of the resistance of the neck (R_{neck}). The attenuation was calculated as: α = 1 - $\Delta V_{max, shaft}$/$\Delta V_{max, spine}$. Red cross: mean value of α.

(E) Estimated amplitude of intracellular calcium concentration transients $\Delta[Ca^{2+}]_{max}$, following activation of NMDA receptors and VDCCs as a function of ePSD area. Three spiking outliers are not represented.

Figure 7. Effect of the distance between excitatory and inhibitory synapses on the integration of coincident EPSPs and IPSPs.

(A) Sketch: an EPSP was elicited in the spine and an IPSP in the shaft at a distance Δx from the spine.

(B-C) Voltage inhibition, inh_V, calculated in the soma (B) or in the spine head (C) as a function of Δx, for N=3700 iterations of the model. On-path inhibition: $\Delta x > 0$; off-path inhibition: $\Delta x < 0$. Solid lines represent medians. Shaded areas represent 68% confidence intervals, which span approximately one standard deviation on each side of the mean.

Figure 8. Effect of dendritic and spinous inhibition on EPSPs.

(A) Sketches: an EPSP (arrow) was elicited in a bootstrapped DiS placed randomly along the dendrite, and an IPSP ($\top$ symbol) was elicited either in the dendritic shaft at $\Delta x = 0.7$ μm from the stem of the spine (A1, blue) or directly in the spine head (A2, orange).

(B-C) Quantification of the inhibitory impact, inh_V, in the soma (B) and in the spine head (C) for N=3700 iterations of the model. Blue: dendritic inhibition; Orange: spinous inhibition.

Figure 9. Effect of input timing on EPSP and IPSP integration.

(A) Schematic: excitatory AMPA and NMDA conductances were activated at t=0. The inhibitory GABAergic conductance was activated at an interval Δt before or after the onset of excitation.

(B) Examples of the time-course of depolarization in the spine head for Δt = +5 ms (B1) and Δt = -5 ms (B2) (purple curves) compared to no inhibition (magenta curves) for E_{Cl^-} = -80 mV and V_{rest} = -70 mV. Arrows represent the onset of excitatory and inhibitory inputs (magenta and green arrows, respectively). τ_E represents the 80-to-20% decay time (case with no inhibition).

(C) Ratio of 80-to-20% decay time of membrane depolarizations in the presence of inhibition (τ_{E+I}) over that without inhibition (τ_E) as a function of Δt for dendritic (blue) or spinous (orange) inhibition. Solid lines represent medians. Shaded areas represent 68% confidence intervals, which span approximately one standard deviation on each side of the mean.

(D) Voltage inhibition in the spine head, inh_V, induced by dendritic (blue) or spinous (orange) IPSPs as a function of Δt.

(E) Inhibition of the calcium influx in the spine head, $inh_{[Ca^{2+}]}$, induced by dendritic (blue) or spinous (orange) IPSPs as a function of Δt.