

Cell Type–Specific Thalamic Innervation in a Column of Rat Vibrissal Cortex

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This is the concluding article in a series of 3 studies that investigate the anatomical determinants of thalamocortical (TC) input to excitatory neurons in a cortical column of rat primary somatosensory cortex (S1). We used viral synaptophysin-enhanced green fluorescent protein expression in thalamic neurons and reconstructions of biocytin-labeled cortical neurons in TC slices to quantify the number and distribution of boutons from the ventral posterior medial (VPM) and posteromedial (POM) nuclei potentially innervating dendritic arbors of excitatory neurons located in layers (L)2–6 of a cortical column in rat somatosensory cortex. We found that 1) all types of excitatory neurons potentially receive substantial TC input (90–580 boutons per neuron); 2) pyramidal neurons in L3–L6 receive dual TC input from both VPM and POM that is potentially of equal magnitude for thick-tufted L5 pyramidal neurons (ca. 300 boutons each from VPM and POM); 3) L3, L4, and L5 pyramidal neurons have multiple (2–4) subcellular TC innervation domains that match the dendritic compartments of pyramidal cells; and 4) a subtype of thick-tufted L5 pyramidal neurons has an additional VPM innervation domain in L4. The multiple subcellular TC innervation domains of L5 pyramidal neurons may partly explain their specific action potential patterns observed in vivo. We conclude that the substantial potential TC innervation of all excitatory neuron types in a cortical column constitutes an anatomical basis for the initial near-simultaneous representation of a sensory stimulus in different neuron types.

Keywords: barrel cortex, dendrite domains, POM, thalamocortical projection, VPM

Introduction

The primary somatosensory (barrel) cortex is effectively driven by weak but near-synchronous thalamocortical (TC) input to the granular layer 4 (Bruno and Sakmann 2006). Anatomically, the TC input to layer 4 (L4) was first described as “specific” TC afferents arising from thalamic relay nuclei, which ascend undivided through L5 and L6 and form a dense plexus of axons in cytoarchitectonic granular L4 and a less dense plexus in L3 (Lorente de No 1922, 1938, 1992; see also Jones and Powell 1970; Killackey 1973). “Unspecific” TC afferents were found to

be sparsely distributed in all cortical layers and to arborize predominantly in L1 and L6 (Lorente de No 1938). This observation has been taken as evidence for the view that TC excitation of the cortex is effected mainly, if not exclusively, by the initial excitation of L4 neurons. In this view of a “hierarchical processing scheme,” as it was first proposed for the visual cortex by Hubel and Wiesel (1962, 1968), sensory excitation then spreads from L4 to supra- and infragranular layers (Simons 1978, 1985; Armstrong-James et al. 1992; Binzegger et al. 2004; Douglas and Martin 2004).

However, dendrites of excitatory neurons from supra- and infragranular layers are also contacted by TC synapses within L4 (White 1978; Peters 1979). It was concluded that TC synapses were determined mainly by the overlap of TC axons with the dendrites of cortical cells and that TC axons make synapses with virtually all neurons that have their somata in L2–L6 (Peters 1979; White 1979). This prediction was further supported by the report of significant TC projections from both the ventroposterior and the posterior thalamic nuclei to supra- and infragranular layers (Wise and Jones 1978; Herkenham 1980; Chmielowska et al. 1989; Lu and Lin 1993).

As functional evidence of whole-column TC excitation, whole-cell, juxtасomal, and extracellular recordings from different cell types indicated that the minimal latency of excitatory postsynaptic potentials and action potential (AP) spiking, evoked by a whisker deflection, is comparable within only a few milliseconds in virtually all layers (Ahissar et al. 2000; Brecht and Sakmann 2002; Brecht et al. 2003; Manns et al. 2004; de Kock et al. 2007). These observations argue for an initial near-simultaneous representation of a tactile stimulus in multiple cortical layers rather than for a sequential representation radiating from the granular layer.

In the sequence of studies reported in this and the 2 preceding articles, we aimed at quantifying the anatomical determinants for near-simultaneous TC activation in the somatosensory cortex. First, we measured the distribution of TC boutons originating in the ventral posterior medial (VPM) nucleus and the medial part of the posterior group nuclei (posteromedial, POM) of the thalamus (Wimmer et al. 2010) using virus-mediated expression of fluorescent proteins in thalamic boutons (Wimmer et al. 2004, 2010). Next, we measured the distribution

of neuronal cell bodies in a cortical column defined by the thalamic projections (Meyer et al. 2010). In the present study, we measured the dendrite distributions of 7 types of cortical excitatory neurons in L2–L6 of a cortical column by *in vitro* whole-cell filling of 82 cortical excitatory neurons. Together with the data on the prevalence of each of these neuron types from the preceding study, this yielded the distribution of putative postsynaptic dendrites of excitatory neurons in a cortical column. Finally, we computed the anatomical overlap between the TC bouton profiles and the distribution of putatively postsynaptic dendrites. The calculation is based on the simplifying assumption that the actual synaptic connectivity is proportional to the potential connectivity quantified as the product of bouton density and dendritic density (i.e., the connectivity is locally random; this notion was first proposed by Peters [1979] and later extended by White [1979, 1989]; for a more recent discussion, see Shepherd et al. [2005]; Stepanyants and Chklovskii [2005]). The resulting estimates of the amount of potential TC innervation of cortical neurons provided quantitative constraints on TC innervation domains.

We found that all excitatory neuron types in a cortical column of somatosensory cortex are potentially innervated by at least 90 TC boutons. This can be regarded as the anatomical basis for the initial near-simultaneous representation of a sensory stimulus measured in different neuron types *in vivo*. In addition, we found that excitatory neurons in L3, L5, and L6 have multiple subcellular TC innervation domains. A subtype of L5 thick-tufted pyramidal neurons had an additional TC innervation domain in L4. Strikingly, L5 thick-tufted pyramidal neurons are potentially innervated by both VPM and POM with comparable numbers of potential contacts per neuron, which may constitute the anatomical basis of their specific AP discharge patterns upon whisker stimulation as observed *in vivo*.

Materials and Methods

We 1) measured the profile of VPM and POM thalamic bouton densities along the longitudinal column axis in somatosensory cortex (using virus-mediated expression of fluorescent proteins in TC boutons as reported in the first article of this series of reports; Wimmer et al. 2010), 2) measured the distribution of neuronal somata stained for the neuron-specific nuclear protein “Neuronal Nuclei” (NeuN; Mullen et al. 1992) in separate experiments (as reported in the second article of this series of reports; Meyer et al. 2010), 3) measured the dendritic density distributions along the longitudinal column axis of neurons that were reconstructed after *in vitro* single-cell recordings in separate experiments ($n = 82$), 4) aligned the bouton and dendritic density profiles using the pia and the white matter (WM) as landmarks in cortex, 5) divided the bouton profiles by a profile representing the distribution of total dendritic length in a cortical column to obtain a profile of normalized TC bouton density per unit dendrite length in a cortical column, and 6) multiplied the aligned TC bouton density profile with dendritic profiles to obtain the distributions of thalamic contacts on individual neurons of different types.

Measurement of Thalamic Bouton Densities

Experiments were conducted in accordance with the German Animal Welfare Act. The methods used for virus delivery and the histological procedures were the same as described before (Wimmer et al. 2004, 2010). Briefly, P28 Wistar rats were anesthetized by intraperitoneal injection of 5 $\mu\text{g}/\text{kg}$ fentanyl, 2 mg/kg midazolam, and 150 $\mu\text{g}/\text{kg}$ medetomidine. Adeno-associated virus (AAV) - synaptophysin - enhanced green fluorescent protein (EGFP) (100–200 nL; 3×10^6 infectious particles/mL) was stereotaxically injected bilaterally into VPM or POM using calibrated injection capillaries. The injection

coordinates were as follows (in mm): 2.85 posterior of bregma, 3.2 lateral of the midline, and 5.05 deep from the pia for VPM and 3.25 posterior, 2.1 lateral, and 5.2 deep for POM. After virus delivery (12–14 days), rats were euthanized with a lethal dose of urethane and transcardially perfused with 15 mL phosphate buffer (PB) followed by 15 mL PB with 4% paraformaldehyde (PFA). The brain was extracted and incubated overnight in PB containing 4% PFA at 4°C. TC vibratome sections (45° from midline) at 70–100 μm thickness were prepared (Bernardo and Woolsey 1987; Agmon and Connors 1991; Fleidervish et al. 1998).

In order to align the virus-labeled bouton profiles with the data on soma distributions and dendritic labelings from different experiments, we defined landmarks in each of the experiments (the details of alignment are described in a separate section, see below). To calibrate the L4-to-L5A border along the longitudinal column axis (as a control landmark) in the virus-labeled slices, these slices were subsequently immunolabeled against NeuN using a monoclonal antibody at 1:1000 (for a detailed protocol, see Meyer et al. 2010).

Then, slices were imaged using confocal mosaic scanning with TCS SP2 (Leica Microsystems) or FV1000 (Olympus) laser scanning microscopes. Either a $\times 63$ Leica (HCX PL APO CS; numerical aperture [NA]: 1.32, oil immersion) or a $\times 40$ Olympus (UPLFLN; NA: 1.3, oil immersion) objective was used. Each mosaic tile measured 180–225 μm in x and y . To obtain the average thalamic bouton fluorescence intensity along the longitudinal axis of a cortical column (“ z -axis”), fluorescence intensity histograms (bin size: 0.4 μm) were calculated from maximum intensity projections of confocal image stacks using ImageJ (<http://rsb.info.nih.gov/ij/>; “bouton z -profiles”). To correct for potential background fluorescence, a linear offset gradient based on the mean of the first 50 (POM) or 100 (VPM) and the last 100 bins was subtracted from the histograms. Finally, histograms were scaled to the standard TC projection column height (Wimmer et al. 2010) of 1840 μm , yielding an effective bin size of 46 μm .

Labeling of Single Excitatory Neurons and Calculation of Dendritic Length Density Profiles

In separate experiments, TC slices of adolescent (mean age: 28.5 ± 2 days, range: 26–42) Wistar rat barrel cortex were cut at 300–350 μm thickness as described previously (Feldmeyer et al. 1999). Using a $\times 2.5$ Olympus objective (Plan Neofluar; NA: 0.075) and brightfield illumination, barrel-related columns in the region of the posteromedial barrel subfield were identified as bright spots in L4 outlined by dark bands. Neurons were visualized using a $\times 40$ Zeiss objective (Plan Achromat; NA: 0.8, water immersion) and infrared differential interference contrast (IR-DIC; Dodt and Ziegglansberger 1990; Stuart et al. 1993)-or infrared gradient contrast (IR-GC; Dodt et al. 1998, 1999)-videomicroscopy. Whole-cell recordings of putative excitatory neurons in barrel columns were made for 10–30 min to fill neurons with biocytin (Sigma). The intracellular solution consisted of (in mM) 135 K-gluconate, 10 HEPES (4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid), 10 phosphocreatine- Na_2 , 4 KCl, 4 ATP-Mg, and 0.3 GTP (adjusted to pH 7.2 with KOH) and contained 0.3–0.6% biocytin. Slices were fixed in 4% PFA for 24 h, labeled using the diaminobenzidine (DAB)-horseradish peroxidase protocol (Horikawa and Armstrong 1988), and embedded in Mowiol (Kuraray Specialities Europe GmbH). Dendritic reconstructions of labeled neurons were made ($n = 82$) with a NeuroLucida system (MicroBrightfield) equipped with a $\times 100$ (NA: 1.25, oil immersion) objective. Slice edges were also traced to approximate the contours of the pia.

All reconstructions were rotated so that the pial surface above the soma was oriented horizontally. For alignment, reconstructions were scaled uniformly so that the distance between pial surface and the border between cortex and the WM was 2000 μm using the custom-made software NeuronRegistrar 1D (written in Matlab, The MathWorks). Three-dimensional dendritic length density maps at a bin size of (50 μm)³ were computed from the scaled reconstructions using the custom-made software Rembrandt 3D (written in Matlab). These 3D maps were projected onto the slice plane (“ x - z plane”) by summing all bins sharing equal x and z coordinates. The resulting 2D map was

projected onto the longitudinal column axis (z -axis; perpendicular to the pial surface) by summing all bins at the same depth from the pial surface, yielding a histogram of dendritic length density along the longitudinal axis of a cortical column ("dendritic z -profile"). All dendritic z -profiles were scaled uniformly to the standard projection column height (1840 μm , yielding a bin size of 46 μm).

TC bouton z -profiles and dendritic z -profiles were visualized as line plots and grayscale color plots after linear interpolation and smoothing with a sliding mean filter using the custom-made software "Rothko" (available at <http://www.mpimf-heidelberg.mpg.de/~mhelmsta/tools/>) written in Matlab.

Comparison of Cytoarchitectonic Landmarks in Virus-Labeled Slices and Whole-Cell Recording Experiments

For alignment of bouton z -profiles and dendritic z -profiles from separate experiments, 3 landmarks in cortex were used: 1) the pial surface, 2) the border between L4 and L5A, and 3) the border between L6 and the subcortical WM. The pia and L6-WM border were identified by visual inspection in both virus-labeled slices and DAB-stained slices from whole-cell recording experiments. The pial surface was clearly identifiable as the border between pial blood vessels and cortical tissue. In DAB-stained slices, pia outlines were traced both at the "upper" and at the "lower" z -edges of the tissue and the average outline was used to correct for the curvature of the cortex. The L6-WM border was defined on maximum intensity projection images of confocal NeuN fluorescence image stacks for virus-labeled slices (see above). Other than L6, the WM contained none or only very few neurons (cf. Supplementary Fig. 1B). In DAB-stained slices, the L6-WM border was defined as the transition to a dark horizontal band representing high density of myelinated axons in the WM (cf. Supplementary Fig. 1A).

For virus-labeled slices, the border between L4 and L5A was defined in the NeuN epifluorescence images as the sharp transition from a layer of small neuron somata at high density representing L4 to a layer of larger neuron somata at low density representing L5A (cf. Supplementary Fig. 1B, see below). In DAB-stained slices from whole-cell recording experiments, the border between L4 and L5A was identified as the transition from the dark horizontal band representing L4 to a less stained region marking the beginning of L5A (cf. Supplementary Fig. 1A).

To compare measurements made in virus-labeled slices with those made in slices from whole-cell recording experiments, we confirmed the correspondence of the L4-to-L5A border derived from DAB staining to the L4-to-L5A border derived from the NeuN-positive soma density (Supplementary Fig. 1, $n = 4$). First, a TC slice was cut at 300 μm thickness and selected for whole-cell recording. Then, the subsequent slice was cut at 200 μm thickness and immediately fixed in 4% PFA. Then, the 200- μm slice was immunolabeled for NeuN, and a maximum intensity projection of confocal NeuN fluorescence image stacks was acquired from the upper 15 μm of the slice (measured perpendicular to the slice plane, see Supplementary Fig. 1B). The L4-L5A border was then visually identified on the maximum intensity projection image as described above for virus-labeled slices. The 300- μm slice was stained using the biocytin-horseradish peroxidase-DAB protocol as described above. Then, we compared the L4-L5A border identified in the DAB-stained slice with that identified in the NeuN-labeled slice (Supplementary Fig. 1A). The ratio of the L4-L5A border depth to the cortical thickness above the hippocampal pole with the pia rotated horizontally was measured in both slices. The mean difference of the ratio was $0.64 \pm 0.37\%$. We concluded that the measurement of the L4-L5A border by visual inspection of NeuN-labeled slices is consistent with that by visual inspection of DAB-stained slices and that we could, as a basic prerequisite for alignment of dendritic z -profiles with bouton z -profiles, compare cytoarchitectonic landmark measurements from whole-cell recording experiments with measurements from virus-labeled slices.

Estimation of the Distribution of Dendritic Length of All Excitatory Neurons in a Cortical Column

In the preceding article, we found a total number of approximately 19 000 neurons ($19\,109 \pm 444$, $n = 3$) in a cortical column as defined by the TC axonal projections (Meyer et al. 2010). To correct for nonexcitatory neurons, we assumed a (GABAergic) interneuron ratio

of 15% (Beaulieu 1993), yielding approximately 16 150 excitatory neurons per cortical column (for a comparison of this simplified assumption to other error sources, see Discussion; for a clarification of the nomenclature "excitatory neuron" and "interneuron", see below).

We calculated the distribution of the dendritic path length of excitatory neurons in a cortical column from single-cell dendritic morphologies and whole-column neuron densities. The 82 neurons in our sample were grouped according to their soma depth from the pial surface (cf. Fig. 5; range of soma depth in a standard column: 124-1617 μm , yielding 17 groups; average group size: 5 neurons, effective bin size: 92 μm). Then, average dendritic z -profiles for each group were calculated by averaging the single-cell dendritic z -profiles within a group. Next, the average dendritic profiles were convolved with the neuron density profile obtained from whole-column counts of NeuN-positive somata as reported in the preceding article (Meyer et al. 2010). This yielded a z -profile of the distribution of the total dendritic length of excitatory neurons along the longitudinal axis of a cortical column (Fig. 4A).

Definition of "Excitatory" Neurons

Neurons were selected for whole-cell recording if the soma had a pyramidal shape when visualized under the IR-DIC or IR-GC optics during the experiment (layers 2/3, 5, and 6); they were selected for recording if the soma had a spheroid shape in L4. The type of the recorded neuron was, however, definitively determined only after staining and reconstruction (as described above). Neurons in the supra- and infragranular layers were included in the sample if they had the morphological features of pyramidal neurons. In the granular layer 4, all spiny neurons were included in the sample. Thus, the sample of 82 neurons analyzed here can be summarized for all layers as "spiny neurons" anatomically. However, since there is ample evidence that the neurons we recorded from are glutamatergic (Markram et al. 1997; Feldmeyer et al. 1999, 2002, 2005, 2006; Mercer et al. 2005; West et al. 2006; Krieger et al. 2007; Helmstaedter, Staiger, et al. 2008), we refer to these neurons as "excitatory" throughout the text.

The somatic staining for NeuN as reported in the preceding article (Meyer et al. 2010) is assumed to label all neurons. We have corrected the total number of neurons for the approximate fraction of GABAergic neurons (15%; Beaulieu 1993; see above; cf. the Discussion for a quantification of the effect of this simplification). We therefore use the term "excitatory neuron" also for the NeuN-positive soma density when corrected for the expected ratio of GABAergic interneurons.

Conversion of Bouton Fluorescence z -Profiles to Absolute Bouton Density z -Profiles

The VPM bouton fluorescence density z -profile can be converted to a z -profile of absolute VPM bouton densities by scaling the integral of the profile to the total number of VPM boutons per column. We assumed a total number of 250 VPM neurons projecting to a cortical column (range: 200-300; Land et al. 1995; Varga et al. 2002). For a single VPM neuron, the mean total axonal path length was taken to be approximately $60 \times 10^3 \mu\text{m}$ (Oberlaender, Bruno, and Sakmann, unpublished observations), of which approximately $35 \times 10^3 \mu\text{m}$ was estimated to be within the home column (Oberlaender, Bruno, and Sakmann, unpublished observations). Bouton density was assumed to be $0.33 \pm 0.02/\mu\text{m}$ axon length (Oberlaender, Bruno, and Sakmann, unpublished observations). Together, this yields an estimated total of 2.9×10^6 VPM boutons in the home column ($250 \times 35 \times 10^3 \times 0.33$, range: 2.3 - 3.5×10^6 given 200-300 VPM neurons). Normalizing the integral of the profile (Fig. 1A) to the total estimated number of VPM boutons then yields a peak density of $4.04 \times 10^7/\text{mm}^3$ (range: 3.22 - $4.83 \times 10^7/\text{mm}^3$) in a cortical column. Due to the lack of anatomical data on POM innervation of a cortical column, we did not estimate the total number of POM boutons projecting to a cortical column. Instead, we measured the difference between VPM bouton density in L4 and POM bouton density in L5A, that is, the difference between the respective peak bouton densities. Boutons marked by synaptophysin-EGFP (see above) were visually identified in confocal image stacks. We counted 2861 (VPM) and 1959 (POM) boutons in 6 (VPM) and 4 (POM) volumes sized $11 \times 10^3 \mu\text{m}^3$ each. VPM peak density ($4.4 \pm 0.6 \times 10^7/\text{mm}^3$) matched the peak density based on anatomical assumptions

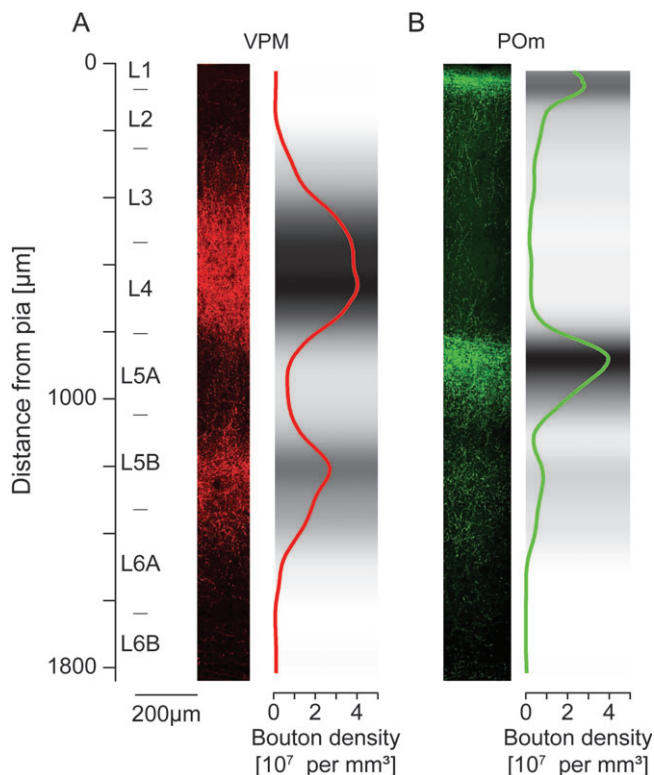


Figure 1. Thalamocortical bouton density profiles in a column. (A) Maximum intensity projection image of a confocal mosaic scan of a thalamocortical slice showing thalamic boutons from the VPM nucleus in a cortical column visualized by virus-expressed synaptophysin-coupled EGFP (left panel). A barrel column in the region above the hippocampal CA3 pole/fornix was selected for confocal scanning. Shown is a 170- μm -wide strip of the column center. The cortical thickness was scaled to a standard column height (distance from pia to the border between cortex and white matter) of 1840 μm . Right panel: VPM bouton fluorescence intensity histogram along the vertical (pia to white matter) column axis (“bouton z-profile”) obtained from the image shown on the left panel. The profile was interpolated and smoothed for visualization. (B) Image showing thalamic boutons from the POm nucleus in a cortical column. The same cortical region as in (A) was selected for confocal scanning. Shown is a 220- μm -wide strip of the column center (the cortical thickness was scaled to 1840 μm). Right panel: POm bouton z-profile obtained from the image shown on the left panel. The profile was interpolated and smoothed for visualization. Bouton densities were calibrated by manual counting of boutons in confocal image stacks. VPM and POm peak bouton densities were found to be equal. Assuming a total of 2.9×10^6 VPM boutons per column (see Materials and Methods), the underlying fluorescence intensities can be converted to bouton densities, yielding a peak density of 4.04×10^7 boutons/ mm^3 for both POm and VPM and 1.8×10^6 total POm boutons per column.

($4.04 \times 10^7/\text{mm}^3$). Since the infection rate of thalamic neurons upon virus injection was estimated to be only 85% (Wimmer et al. 2010), this indicates that our estimate of VPM boutons per home column based on anatomical assumptions may be slightly too low, possibly because it does not account for VPM projections from other than the somatotopically aligned barreloid.

We used the direct bouton density measurements only to compare POm and VPM peak bouton densities. POm peak bouton density ($4.7 \pm 0.6 \times 10^7/\text{mm}^3$) was not significantly different from VPM ($P = 0.54$, paired *t*-test, 2 tailed). We thus scaled the POm bouton fluorescence density profile so that the peak density was equal to that of the VPM bouton density profile, obtaining a profile of absolute POm bouton densities, and a total number of 1.8×10^6 POm boutons projecting to a cortical column. For further calculations, the numbers of VPM and POm boutons were reduced by 15% to account for boutons contacting interneurons based on the ratio of interneurons in a cortical column (see above).

For estimating the contribution of septal neurons to the total dendritic path length in cortex, we took the volume of the septum related to a particular column as approximately 0.13 mm^3 (Table 2 in Meyer et al. 2010, i.e., $0.13/0.24 = 55\%$ of the column volume) with a septal neuron density of $68.6 \times 10^3/\text{mm}^3$ (Table 1 in Meyer et al. 2010, i.e., $68.6/78.9 = 87\%$ of the density inside a column); together, this means that including septal neurons adds approximately 48% to the intracolumnar dendritic length. TC input from other than the somatotopically aligned barreloid increases the TC axonal path length by approximately 67%; together, this yields a $1.67/1.48 = 13\%$ increase under these assumptions. If septal neurons receive less or no direct VPM input, these numbers can substantially change; we therefore estimate a 10–30% increase in the numbers on TC boutons if these effects are included.

Statistical Analyses

All statistical analyses in this study, such as Student’s *t*-tests, one-way analyses of variance (ANOVAs), and multiple comparison tests (Scheffé procedure), were performed with the statistical toolbox of Matlab.

The cluster analysis for identifying the subtype of L5B pyramidal neurons with an additional VPM innervation domain in L4 was made in the following 2D parameter space: 1) the number of oblique dendrites within 300 μm above the L4-to-L5A border and 2) the total dendritic path length within 506–782 μm depth from the pia (corresponding to bins 11–17). The cluster analysis was made as described previously (Cauli et al. 2000; Helmstaedter et al. 2009): Euclidean distances, Ward’s method (Ward 1963), and Thorndike procedure (Thorndike 1953) for linkage distance cutoff (see Fig. 7B, lower panel), rejecting 1-item clusters (all in Matlab).

The statistical test for the predicted number of VPM boutons has to be interpreted cautiously; because the innervation in L4 is highly correlated with the total dendritic density in L4, and because the latter parameter was used for separating the clusters, nonparametric tests are expected to report arbitrary levels of significance. We used a parametric test (here: one-sided *t*-test assuming normally distributed data with unequal variances) that can be interpreted as an indication of the amount of separation between the clusters.

Results

VPM and POm Bouton Profiles

Figure 1 shows dorsomedial portions of barrel cortex in the section plane of the TC slice after injection of VPM or POm with AAV expressing EGFP-coupled synaptophysin. Figure 1A (left panel) shows the staining of boutons of axons originating in the VPM. Figure 1B (left panel) shows another TC section with staining of boutons of axons originating in the POm. Both sections illustrate the differential vertical density distributions of VPM and POm boutons as quantified in the bouton fluorescence intensity histograms (“bouton z-profiles”, right panels of Fig. 1A,B). Each profile represents the respective thalamic bouton fluorescence intensity as a function of the distance from the pial surface along the longitudinal column axis (for calculation of bouton z-profiles, see Materials and Methods). VPM and POm bouton density peaks were well separated, with a local overlap at 780–830 μm depth below the pia if scaled to a standard column height of 1840 μm (cf. Fig. 1A,B). The bouton fluorescence z-profiles were then converted to absolute bouton density z-profiles. Manual bouton counts yielded peak TC bouton densities of $4.4 \pm 0.6 \times 10^7/\text{mm}^3$ and $4.7 \pm 0.6 \times 10^7/\text{mm}^3$ for VPM and POm, respectively. These densities matched the anatomical estimate of VPM innervation based on assumptions of the number of thalamic neurons in VPM, the axonal path length of VPM projections, and the bouton density on VPM axons ($4.04 \times 10^7/\text{mm}^3$ peak density and a total of 2.9×10^6 boutons in a cortical column). The total number of

Table 1

Predicted number of thalamocortical boutons per excitatory neuron

| Type of excitatory neuron | VPM boutons per neuron | P0m boutons per neuron | TC boutons per neuron (VPM + P0m) | Total dendritic length per neuron (μm) | Spines per neuron | Ratio of spines occupied by VPM (%) | Mean distance between VPM boutons along dendrites (μm) | Ratio of spines occupied by P0m (%) | Mean distance between P0m boutons along dendrite (μm) |
|--------------------------------|------------------------|------------------------|-----------------------------------|---|-------------------|-------------------------------------|---|-------------------------------------|--|
| L2 pyramid ($n = 7$) | 16 \pm 14 | 75 \pm 33 | 92 \pm 36 | 4309 \pm 1716 | 2155 \pm 858 | 1 | 265 | 3 | 57 |
| L3 pyramid ($n = 9$) | 222 \pm 131 | 63 \pm 16 | 285 \pm 132 | 5947 \pm 1147 | 2973 \pm 574 | 7 | 27 | 2 | 95 |
| L4 star pyramid ($n = 6$) | 197 \pm 46 | 57 \pm 46 | 254 \pm 65 | 3553 \pm 732 | 1777 \pm 366 | 11 | 18 | 3 | 62 |
| L4 spiny stellate ($n = 7$) | 188 \pm 60 | 22 \pm 10 | 210 \pm 61 | 2549 \pm 648 | 1275 \pm 324 | 15 | 14 | 2 | 116 |
| L5 slender tufted ($n = 20$) | 150 \pm 50 | 269 \pm 121 | 420 \pm 131 | 5774 \pm 1749 | 2887 \pm 874 | 5 | 38 | 9 | 21 |
| L5 thick tufted ($n = 22$) | 299 \pm 99 | 277 \pm 67 | 576 \pm 120 | 10 356 \pm 1917 | 5178 \pm 959 | 6 | 35 | 5 | 37 |
| L6A pyramid ($n = 11$) | 131 \pm 66 | 66 \pm 42 | 197 \pm 79 | 3908 \pm 794 | 1954 \pm 397 | 7 | 30 | 3 | 59 |

Note: The predicted number of boutons calculated for each neuron as the integral of the VPM or P0m innervation density profile (see inset in Fig. 5B). The number of spines per neuron was calculated assuming a spine density of $0.5/\mu\text{m}$ dendritic path length. Mean distance of VPM and P0m boutons: total dendritic length per neuron divided by the number of boutons per neuron. All values reported as mean \pm standard deviation. Note that the predicted absolute number of thalamocortical boutons per neuron is based on the axonal path length of VPM neurons extended in the somatotopically aligned home column, only, and thus represents a lower bound estimate. All values have an additional estimated uncertainty of $\pm 30\%$ due to the uncertainty in assumptions on the total number of thalamocortical boutons in a column, see Materials and Methods and Discussion for details.

P0m boutons in a cortical column was estimated to be 1.8×10^6 (for details of the calculations, see Materials and Methods).

Distribution of the Dendrites of Excitatory Neurons in a Cortical Column

To quantify the overlap of TC bouton density distributions with dendritic length density distributions of excitatory neurons in a barrel-related column, we labeled neurons in acute brain slices of barrel cortex during whole-cell recording. Their soma and dendritic arbors were reconstructed. Figure 2 shows one reconstruction of an L5 thick-tufted neuron superimposed on a brightfield image of the acute slice (a brightfield image of the same slice after DAB staining is shown in Supplementary Fig. 1A; for a quantification of dendritic length ratio outside the home column, see Supplementary Fig. 2). Dendritic length density histograms (“dendritic z -profiles”) were calculated from the reconstructions, representing the dendritic length as a function of the distance to the pial surface along the longitudinal column axis (right panel in Fig. 2; for details, see Materials and Methods). Both the thalamic bouton z -profiles and the reconstructed neurons were obtained from barrel columns located above the lateral ventricle/hippocampal fornix region, where the large barrels of macrovibrissae are located.

To confirm that the cytoarchitecture of the barrel columns analyzed for thalamic bouton distributions (Fig. 1) was similar to that of the barrel columns containing biocytin-filled neurons (Fig. 2), we compared cytoarchitectonic registration landmarks (for details of the comparability of landmarks obtained from DAB-stained slices with landmarks found in virus-labeled slices, see Materials and Methods; Supplementary Fig. 1). The cortical thickness (distance from pia to L6-WM border) was $1860 \mu\text{m}$ in the VPM bouton-labeled slice (Fig. 1A), $2050 \mu\text{m}$ in the P0m bouton-labeled slice (Fig. 1B), and $1994 \pm 105 \mu\text{m}$ ($n = 82$) in the slices containing DAB-stained biocytin-filled neurons. All measurements were then scaled uniformly to a standard projection column height of $1840 \mu\text{m}$ (the dimensions of the “standard” projection column were reported in the first article of this series; Wimmer et al. 2010). The ratio of the L4–L5A border to the total cortical thickness was comparable (43.8% and 44.1% in the virus-labeled slices vs. $43.7 \pm 2.4\%$ in DAB-stained slices). We concluded that bouton z -profiles based on the virus-labeled slices and dendritic z -profiles based on the DAB-stained slices could be superimposed after registration along the z -axis.

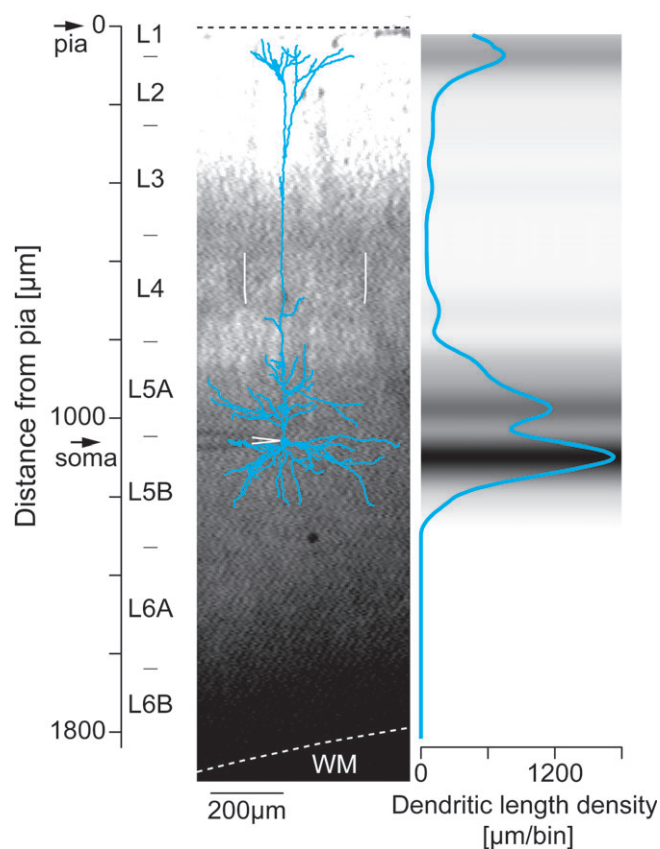


Figure 2. Dendritic length density profile of an L5 thick-tufted pyramidal neuron. Brightfield image of an acute thalamocortical slice of barrel cortex (left panel). Brightness/contrast was adjusted for better visualization of the barrel borders (lateral borders indicated by white lines; see Supplementary Fig. 1 for an image of the same slice after diaminobenzidine labeling for biocytin). WM: white matter. Soma location of the whole-cell recorded and biocytin-filled neuron indicated by white pipette tip. Blue: soma-dendritic reconstruction. For visualization, the reconstruction was scaled uniformly to a standard column height of $1840 \mu\text{m}$. Right panel: Dendritic length density histogram along the vertical (pia to WM) column axis (“dendritic z -profile”) of the neuron shown on the left panel. The profile was interpolated and smoothed for visualization.

Our sample comprised 82 excitatory neurons. These were subclassified qualitatively into 7 types based on their dendritic shape and soma-to-pia distance (Fig. 3; for a depiction of all dendritic reconstructions, cf. Supplementary Fig. 3). L2 pyramidal neurons mostly had a broad apical tuft with an early

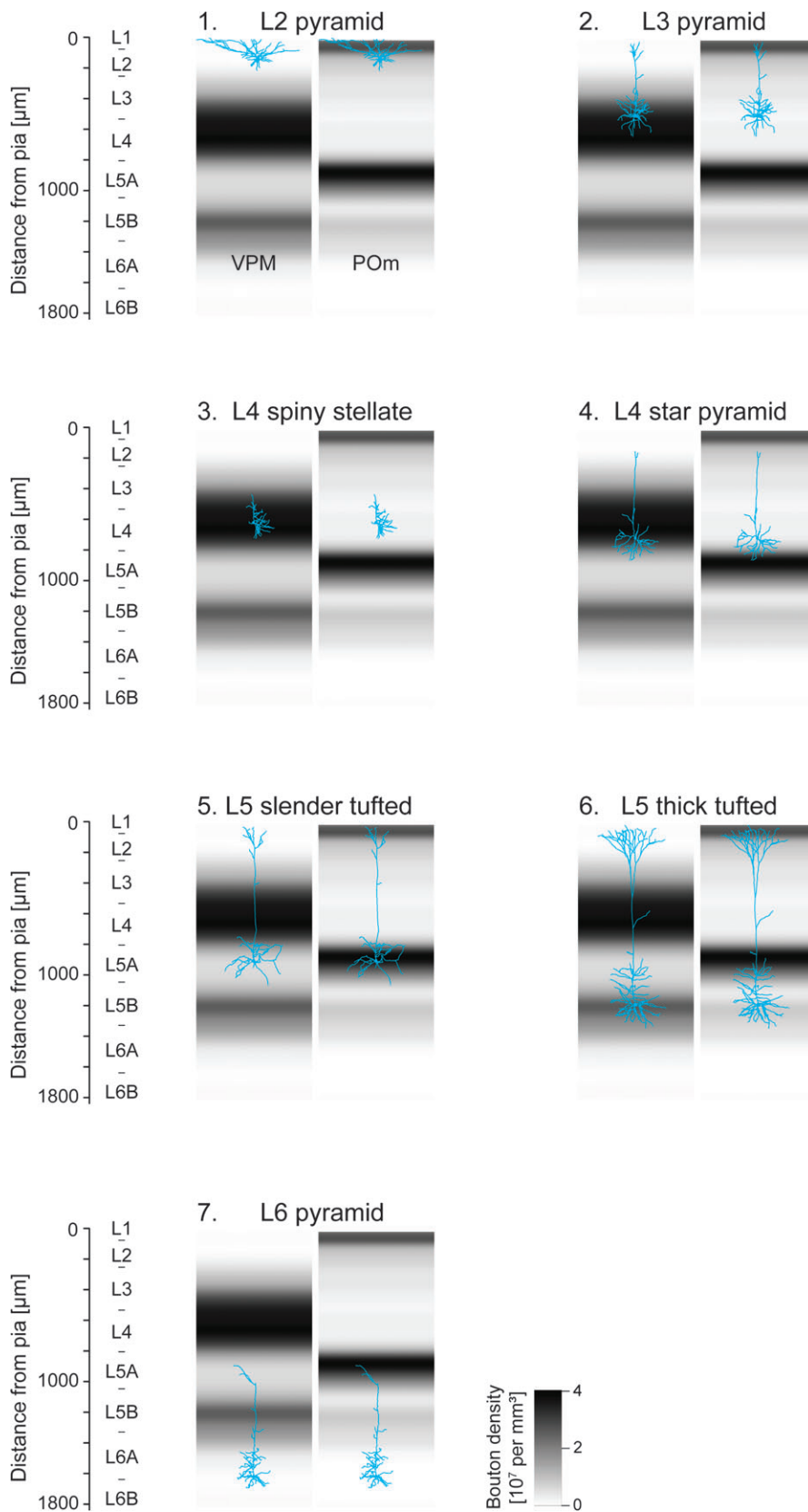


Figure 3. Examples of the overlap between single soma-dendritic neuron reconstructions of 7 types of excitatory cortical neurons and the VPM and POm bouton z-profiles. Representative single-cell reconstructions (blue) superimposed on the VPM bouton intensity profile (gray values, left panels) and the POm bouton intensity profile (right panels). Both VPM and POm bouton z-profiles were normalized to the peak fluorescence intensity (for conversion to bouton density, cf. Fig. 1 and Materials and Methods). Color bar applies to all panels.

primary bifurcation ($n = 7$, soma distance from pia: 124–275 μm); L3 pyramidal neurons had longer apical dendrites with narrower apical tufts and longer basal dendrites ($n = 9$, soma-to-pia distance: 295–529 μm); L4 star pyramidal neurons had an apical dendrite that did not reach L1 and had no prominent tuft ($n = 6$, soma-to-pia distance: 614–750 μm); L4 spiny stellate neurons had short mostly asymmetric primary dendrites and did not show a prominent apical dendrite ($n = 7$, soma-to-pia distance: 607–802 μm); L5 slender-tufted pyramidal neurons had a small apical tuft and few apical oblique dendrites ($n = 20$, soma-to-pia distance: 772–975 μm); L5 thick-tufted pyramidal neurons had a broad apical tuft and more apical oblique dendrites ($n = 22$, soma-to-pia distance: 997–1251 μm); L6 pyramidal neurons had apical dendrites that did not reach beyond L4 ($n = 11$, soma-to-pia distance: 1269–1617 μm , cf. Brumberg et al. 2003; Andjelic et al. 2009). According to their soma depth, the L6 neurons in our sample may be regarded as L6A neurons (Torres-Reveron and Friedlander 2007). We did not differentiate different dendritic subtypes of L6 neurons (Kumar and Ohana 2008; Andjelic et al. 2009; Chen et al. 2009).

Figure 3 shows examples of individual somatic and dendritic reconstructions representative of the 7 excitatory cell types, superimposed on VPM (left panels) and POm (right panels) bouton z -profiles. Any overlap of domains of high thalamic bouton density and dendritic arbors of cortical neurons indicates that these neurons could receive direct thalamic input after a sensory stimulus.

Quantification of TC Innervation of Individual Neurons

In order to compute the probability of TC innervation for the different types of cortical neurons, TC bouton z -profiles (normalized to represent the absolute number of thalamic contacts; see Fig. 1 and Materials and Methods) were multiplied with dendritic z -profiles (Fig. 2). As a simplification, at a given cortical depth, the number of thalamic boutons was taken to be uniformly “distributed” between all dendrites (Peters 1979; White 1979, 1989; Shepherd et al. 2005; Stepanyants and Chklovskii 2005). An individual dendrite will then receive the fraction of the boutons that corresponds to the ratio of its length to the total dendritic length present at that location.

We therefore estimated the total dendritic length distribution of excitatory neurons in a cortical column. The average dendritic z -profiles of excitatory neurons at a given soma depth were convolved with a neuron density z -profile that represented the number of excitatory neurons at that depth within an average TC projection column as defined in the 2 preceding articles (Meyer et al. 2010; Wimmer et al. 2010). This yielded a z -profile of the total dendritic path length of excitatory neurons at a given depth in a cortical column (“total dendritic z -profile”; Fig. 4A; for details, see Materials and Methods). The total dendritic length density was highest in the upper 150 μm of the cortex, with a peak density of 812 m/mm^3 at 70 μm depth, corresponding to the apical tufts of all pyramidal neurons in a cortical column. The total dendritic length of all excitatory neurons in a cortical column was 75 m.

We then divided the VPM and POm bouton z -profiles (Fig. 1) by the total dendritic z -profile (Fig. 4A). The resulting z -profiles show the depth dependence of VPM (red line in Fig. 4B) and POm (green line in Fig. 4B) bouton density per unit dendritic path length in a cortical column (“normalized” VPM and POm

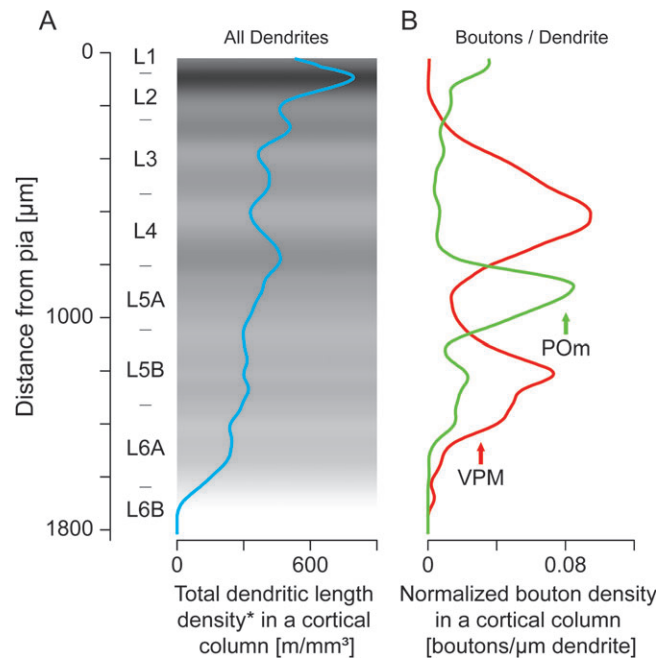


Figure 4. Total dendritic length distribution and normalized thalamocortical innervation density in a cortical column. (A) Dendritic z -profile of all excitatory neurons in a cortical column. All excitatory neurons were grouped based on soma depth from the pia surface. Average dendritic z -profiles were calculated for all groups. Each profile was then multiplied by the total number of excitatory neurons in a cortical column within the respective bin and the multiplied profiles were summated (i.e., the average dendritic z -profiles were convolved with a soma density profile). This yielded a profile representing the total excitatory neuron dendritic length at a given depth, which was finally normalized to the total cortical column volume to obtain an estimate of the dendritic path length per volume density (m/mm^3). *Corrected for the expected fraction of excitatory neurons (85%, see Materials and Methods). (B) Normalized VPM (red line) and normalized POm (green line) bouton density profiles in units of boutons per micrometer dendrite obtained by dividing the bouton z -profiles (Fig. 1) by the total dendritic length profile as shown in (A). Profiles were interpolated and smoothed for visualization (see Materials and Methods).

bouton density profiles). Normalized VPM bouton density was highest between 400 and 800 μm depth from the pia (maximum: 0.10 boutons/ μm dendrite). Normalized POm bouton density was highest between 900 and 1100 μm depth (maximum: 0.09 boutons/ μm dendrite). Note that the normalization to dendritic path length modified the VPM and POm bouton profiles as the dendritic length density in a cortical column varied with depth from the pia (Fig. 4A). The normalized POm bouton density peak close to the pia was relatively lower compared with the raw POm bouton density peak (cf. green line in Figs 1B and 4B), reflecting the high dendritic density below the pia due to the apical tufts.

TC Innervation Depends on Neuron Type and Soma Depth

The normalized bouton density profiles (Fig. 4B) were then multiplied by the dendritic z -profiles of individual neurons (Fig. 2), yielding z -profiles of TC innervation density that represent the distribution of the absolute number of VPM or POm boutons contacting a given neuron at a given depth (insets in Fig. 5B for the example neuron shown in Fig. 2).

To compare the potential TC input of different neuron types, the average number of POm and VPM boutons per type was calculated from the integrals of these individual thalamic contact density z -profiles (see insets in Fig. 5). The cell types

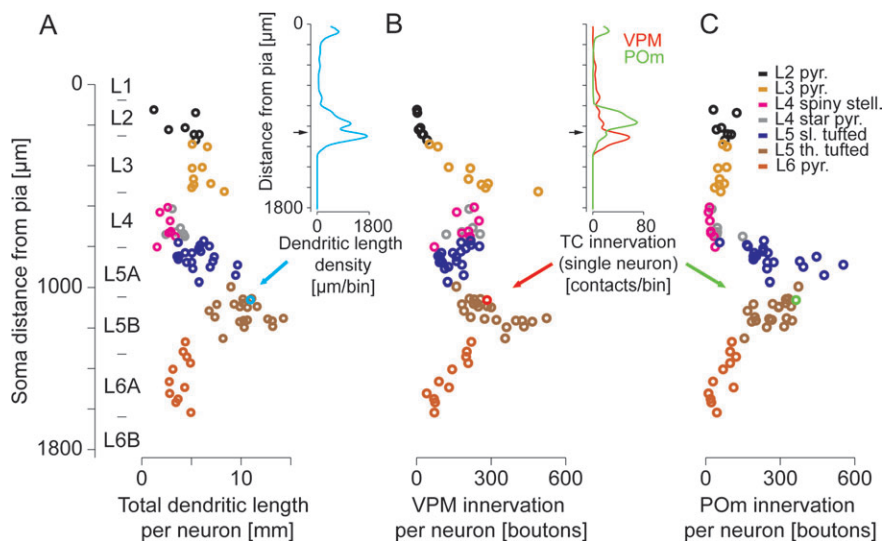


Figure 5. Predicted number of thalamocortical boutons per neuron in dependence of soma location along the vertical column axis. (A) Soma depth dependence of total dendritic length per neuron quantified for all 82 excitatory neurons in the sample as the integral of the dendritic z-profiles. The dendritic length per neuron increased with soma depth in L2/L3 and L5 but not within groups of neurons of the same type. Colors: types of excitatory cortical neurons as defined by their somatodendritic morphologies (legend in (C)). For a depiction of all reconstructions, see Supplementary Figure 3. Blue arrow: neuron shown in the inset. Inset: example of the dendritic length density distribution of one neuron (same neuron as shown in Fig. 2); black arrow: soma position. (B) Soma depth dependence of total putative VPM innervation per neuron quantified as the integral of the single-neuron VPM innervation z-profile (inset, red line). Note that the total innervation of L3 and L5 thick-tufted pyramidal neurons increased significantly with soma depth, while L5 slender-tufted and L6 pyramidal innervation decreased. Inset: Potential VPM (red line) and POm (green line) innervation z-profile of the L5 thick-tufted pyramidal neuron shown in Figure 2. Multiplication of the normalized bouton density profiles (Fig. 4B) with dendritic z-profiles of individual neurons (Fig. 2, inset in (A)) yields profiles representing the absolute number of putative thalamocortical contacts. The integrals of the innervation z-profiles are an estimate of the absolute number of boutons contacting a neuron. Vertical axis as inset in (A). (C) Soma depth dependence of total POm innervation per neuron. Quantification as in (B). Innervation of L5 slender-tufted pyramidal neurons increased significantly with soma depth, while L5 thick-tufted and L6 pyramidal neuron innervation decreased.

were significantly different in average VPM innervation ($P < 0.0001$, one-way ANOVA; Table 1 and Fig. 5), with a peak for L5 thick-tufted pyramidal neurons (299 ± 99 boutons per neuron). The mean VPM innervation of L5 thick-tufted pyramidal neurons was significantly higher than VPM innervation of L2 (16 ± 14), L5 slender-tufted (150 ± 50), and L6 (131 ± 66) pyramidal neurons ($P < 0.05$, multiple comparison test, Scheffé procedure). There was no significant difference between L3 (222 ± 131), L4 (star pyramidal neurons: 197 ± 46 ; spiny stellates: 188 ± 60), L5 slender-tufted, and L6 neurons. This shows that, on average, excitatory neurons in layers 3, 4, 5, and 6 can receive significant VPM innervation. Potential POm innervation was highest for L5 pyramidal neurons (slender tufted: 269 ± 121 ; thick tufted: 277 ± 67) and showed higher variability for the other neuron types (see Table 1). All excitatory neuron types with the exception of L4 spiny stellates potentially receive at least 60 POm boutons.

We then analyzed the correlation of soma depth and TC innervation for different neurons of the same type. Total dendritic length (i.e., the integrals of dendritic z-profiles, cf. Fig. 2) was not found to be significantly depth dependent for any neuron type (open circles in Fig. 5A). When total VPM innervation per neuron was plotted as a function of pia-soma distance (open circles in Fig. 5B), however, L3, L5, and L6 pyramidal neurons showed a dependence on soma depth, and 2 patterns emerged: While the innervation of L3 ($r = 0.87$, $P < 0.01$) and L5 thick-tufted ($r = 0.70$, $P < 0.001$) pyramidal neurons increased significantly with increasing soma depth, VPM innervation decreased for L5 slender-tufted ($r = -0.46$, $P < 0.05$) and L6 ($r = -0.92$, $P < 0.0001$) pyramidal neurons. For L5 and L6 neurons, POm innervation also depended on soma depth (open circles in Fig. 5C): In contrast to VPM innervation, POm innervation increased with soma depth for L5 slender-

tufted neurons ($r = 0.57$, $P < 0.01$), while it decreased for L5 thick-tufted neurons ($r = -0.57$, $P < 0.01$). For L6 pyramidal neurons, POm innervation, like VPM innervation, decreased with soma depth ($r = -0.75$, $P < 0.01$). These correlations indicate that, anatomically, there is a high variability in potential TC input of L3, L5, and L6 neurons, which can be, in part, explained by the vertical position of the individual neurons in the cortical column that determines the axodendritic overlap.

Subcellular TC Innervation Domains for 7 Excitatory Neuron Types

We finally quantified potential subcellular TC innervation domains for each type of excitatory neuron. Figure 6 shows averaged soma-centered dendritic z-profiles (left panels) and thalamic contact density z-profiles (VPM: middle panel, POm: right panel) of all 7 types of excitatory neurons (for an illustration of the underlying anatomical overlap, cf. Fig. 3; for calculation of contact density z-profiles, cf. Fig. 4). Each contact z-profile represents the vertical distribution of potential thalamic input along the dendritic compartments of a particular type. The number of potential TC contacts for each cell type, specified for the basal dendrites, proximal apical oblique dendrites, distal apical oblique dendrites, and apical tuft dendrites is reported in Table 2.

Figure 6 shows that pyramidal neurons in L2 are likely to receive POm input at their apical tufts and probably lack VPM input. Pyramidal neurons in L3 are likely to receive strong VPM input at their basal and apical oblique dendrites, and, to a smaller degree, distributed POm input at their apical oblique dendrites and apical tuft. L4 spiny stellate neurons are likely to receive almost exclusively VPM input. L4 star pyramidal neurons are likely to receive VPM input at their basal dendrites and apical oblique dendrites and POm input at their basal

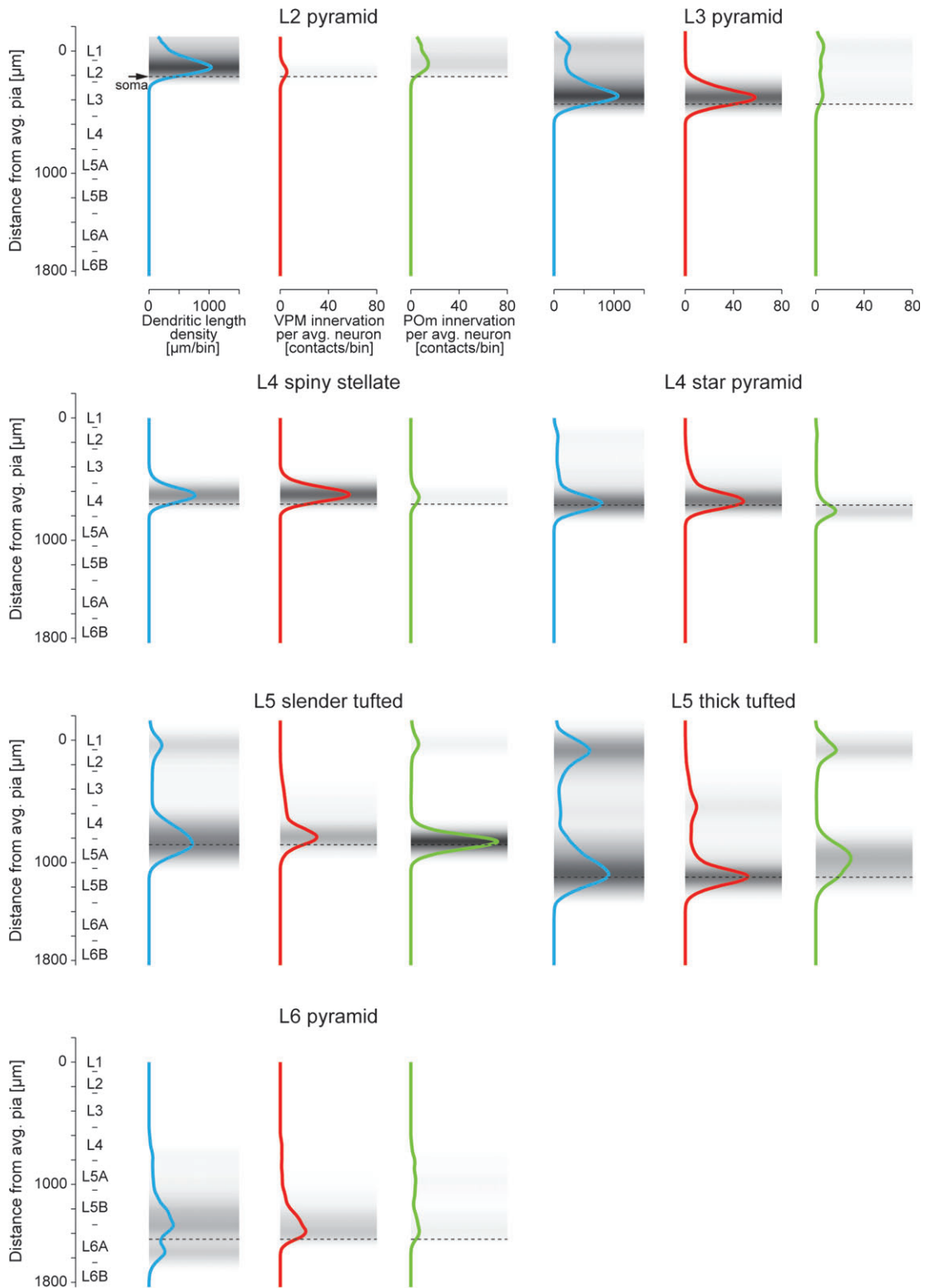


Figure 6. Average dendritic z-profiles and thalamocortical innervation density profiles of 7 excitatory neuron types. Left panels: Soma-aligned dendritic z-profiles (cf. Fig. 2) averaged for all neurons of a given type (see Supplementary Fig. 3 for all reconstructions). Average soma positions per type are indicated (dashed lines). Middle and right panels: Average VPM (middle panels) and POm (right panels) innervation density z-profiles representing the vertical distribution of potential thalamic contacts received by the 7 excitatory neuron types in a cortical column. Dashed lines indicate average soma positions. Note that because the profiles were aligned to the soma of the neurons, the average profiles can extend beyond the average pia (at $z = 0$).

Table 2
Predicted number of thalamocortical boutons in the subcellular innervation domains

| Domain | L2 | | L3 | | L4sp | | L5sl | | L5th | | L6 | |
|-----------------------------------|-----|-----|-----|-----|------|-----|------|-----|------|-----|-----|-----|
| | VPM | POm | VPM | POm | VPM | POm | VPM | POm | VPM | POm | VPM | POm |
| Tuft | 1 | 44 | 0 | 22 | 0 | 3 | 1 | 25 | 12 | 68 | 5 | 4 |
| Distal apical oblique dendrites | 9 | 28 | 7 | 15 | 9 | 2 | 17 | 2 | 54 | 6 | 14 | 25 |
| Proximal apical oblique dendrites | | | 163 | 21 | 119 | 9 | 101 | 118 | 116 | 165 | 97 | 33 |
| basal dendrites | 6 | 3 | 52 | 5 | 69 | 43 | 31 | 124 | 118 | 37 | 15 | 5 |
| Sum | 16 | 75 | 222 | 63 | 197 | 57 | 150 | 269 | 299 | 277 | 131 | 66 |

Note: Integrals of the thalamocortical innervation density profiles (as in Table 1), differentiated by the dendritic domains of each excitatory neuron type. Dendritic domains were defined using the dendritic length density profiles (see Fig. 2, cf. Supplementary Fig. 3): basal dendrites: below the soma; apical tuft dendrites: above the first bin of the apical tuft dendritic density peak; the remaining part of the dendritic density profile was split in half to approximately define proximal and distal apical oblique dendrites. For data on the subgroups of L5 thick-tufted pyramidal cells, see Supplementary Table 2.

dendrites. L5 slender-tufted pyramidal neurons are likely to receive VPM input at their apical oblique dendrites. They are also likely to receive POm input at their basal and apical oblique dendrites and, to a smaller degree, at their apical tuft. L5 thick-tufted pyramidal neurons are likely to receive strong VPM input at their basal dendrites and at their proximal and distal apical obliques and in addition POm input at the proximal apical oblique dendrites and at their apical tufts. Finally, a fraction of L6 neurons is likely to receive VPM input at their apical oblique dendrites.

In summary, dendritic fields of all excitatory neuron types overlap with thalamic boutons. L3 pyramidal neurons, L4 star pyramidal neurons, and L5 slender-tufted and L5 thick-tufted pyramidal neurons can all receive direct thalamic input originating from 2 different thalamic subnuclei.

Subcellular TC Innervation Domains Define a Subtype of Thick-Tufted L5B Neurons

We found a small fraction of thick-tufted L5B pyramidal neurons ($n = 4$ of 22) that extended a substantial part of their apical oblique dendrites to L4 (Fig. 7A, right panel for an example), while the majority of thick-tufted neurons had virtually no dendrites in L4 (e.g., Fig. 7A, left panel; see Supplementary Fig. 3 for all L5B reconstructions). Quantitatively, these neurons were identified by a cluster analysis with the number of apical oblique dendrites in L4 and the total dendritic path length in L4 as parameters (Fig. 7B, right panel, see Materials and Methods). They had more than 4 apical oblique dendrites and more than 1 mm dendritic path length in L4 (Fig. 7B, left panel). Because this specific branching was well aligned with the VPM projections terminating in L4, this subtype of L5B neurons had a fourth subcellular TC innervation domain, which the majority of L5B cells did not show (Fig. 7C vs. D, Supplementary Table 2; predicted number of TC boutons per neuron in the L4 innervation domain: 132 ± 72 [$n = 4$] vs. 29 ± 5.8 [$n = 18$], $P < 0.03$; see Materials and Methods for details and validity of this statistical test).

Discussion

The aim of the present work was to make quantitative predictions about the innervation of excitatory cortical neurons by TC boutons in a column of primary somatosensory

barrel cortex. For this purpose, the dendritic domains of the neuron types were superimposed with TC bouton domains, and absolute numbers of TC innervation were estimated. In previous work, these neurons were shown to respond to a simple sensory stimulus (whisker deflection) in a type-specific manner (Brecht and Sakmann 2002; Brecht et al. 2003; Manns et al. 2004; de Kock et al. 2007). The present results suggest a possible anatomical basis for at least some of the cell type-specific responses.

Neuron Type-Specific Subcellular Innervation Domains

The analysis of subcellular TC innervation domains on excitatory cortical neurons (Figs 6–8 and Table 2) revealed highly cell type-specific innervation patterns. These can be interpreted as follows (see Figs 6 and 8): TC innervation was “focal” and “unique” (one subcellular innervation domain, innervation by one thalamic nucleus) for pyramidal neurons in L2 (POm innervation domain at the apical tuft), L4 spiny stellate neurons (VPM innervation domain throughout their entire dendritic field), and lower L6 pyramidal neurons (VPM innervation domain at their apical oblique dendrites, cf. also Fig. 5B). Pyramidal neurons in L3, L4, L5, and upper L6 on the other hand showed “multiple” and “dual” TC innervation (2, 3, or 4 subcellular innervation domains, innervation by 2 thalamic nuclei). For L2 pyramidal neurons, TC innervation was restricted to a rather broad innervation domain by POm at their apical tuft, whereas their remaining dendritic compartments were potentially only sparsely contacted by TC projections at all. Basal and apical oblique dendrites of L2 pyramidal neurons may be targeted mainly by intracortical connections (Bannister 2005; Bureau et al. 2006).

Clustered versus Distributed Subcellular TC Innervation Domains

We quantified both the total expected TC innervation per neuron (Fig. 5 and Table 1) and the spatial distribution of TC innervation along the dendrites of a given neuron (Figs 6–8 and Table 2). While the first method accurately describes the expected total amount of TC input per neuron, the latter is better suited to detect clustered innervation domains along the dendrites of a given neuron type. However, it underestimates distributed innervation. It is therefore questionable whether the weak average innervation of L4 spiny stellates by POm (Figs 6 and 8; cf. Fig. 5C; 22 ± 10 POm boutons total, see Table 1) and the innervation of L2 pyramidal neurons by VPM (Figs 6 and 8; cf. Fig. 5B; 16 ± 14 VPM boutons total, see Table 1) in fact represent significant thalamic input to these neurons.

Some L6A pyramidal neurons extend their apical tufts to L5A where they could receive significant POm input (cf. Figs 3, 5C, 6, and 8). This is only evident in the analysis of the total expected innervation per neuron (Fig. 5C and Table 1; 66 ± 42 POm boutons total), especially for upper L6A neurons (Fig. 5C). Note that all other types of pyramidal neurons (located in L2–L5) can receive POm input within L1 where all of these types of pyramidal neurons extend their apical tufts; it is possible that upper L6 pyramidal neurons, although they do not extend an apical dendrite to L1 (see also, e.g., Chen et al. 2009), can potentially still integrate POm and VPM input.

Definition of Excitatory Cell Types

In this study, established morphological features of dendrites and the cortical depth of the soma location were used to

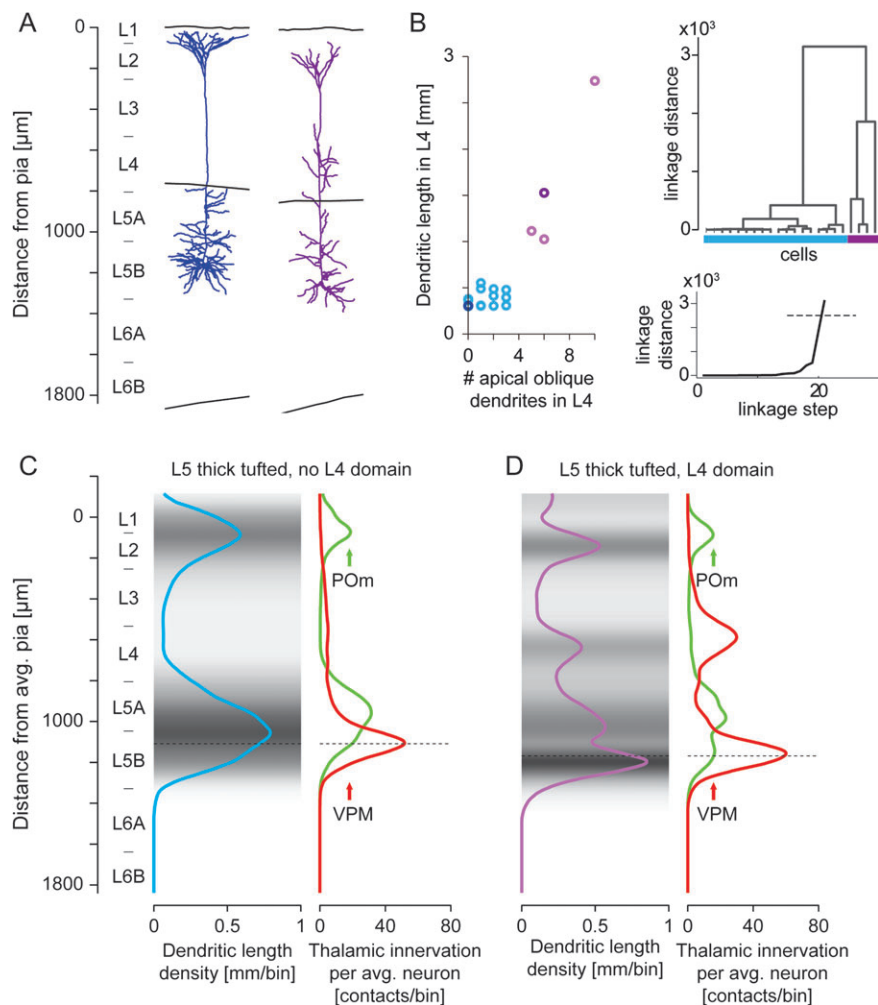


Figure 7. Subgroup of thick-tufted pyramidal neurons with an additional VPM innervation domain in L4. (A) Two examples of L5 thick-tufted pyramidal neurons with different dendritic arborization patterns (blue and purple). Note that the neuron on the left panel has a broad apical tuft and several distal apical oblique dendrites that terminate directly below L4. The neuron on the right panel has a narrower apical tuft and many distal apical oblique dendrites in L4. (B) Quantification of the dendritic arborization in L4 for the whole sample of L5 thick-tufted pyramidal neurons ($n = 22$). In contrast to the majority of the sample, which had no significant dendritic domain in L4 (blue circles, $n = 18$), 4 of the neurons (as defined by a cluster analysis, right panels) had 5 or more apical oblique dendrites of different primary bifurcation origin within 300 μm above the lower L4 border and more than 1000 μm total dendritic length within 500 and 800 μm depth from the pia (purple circles). Dark blue and dark purple circles: neurons shown in (A); for an illustration of all of the L5B neurons, see Supplementary Figure 3. Right panels: linkage dendrogram (top) and linkage distance (bottom) generated in a cluster analysis (Ward's method, Euclidean distances, and Thorndike criterion for cluster cutoff, see Materials and Methods) that was the basis for the clustering shown in the plot on the left. (C, D) Comparison of the dendritic density profiles (left panels) and the resulting VPM and POm innervation profiles (right panels) for the 2 subtypes of thick-tufted L5 neurons as defined in (B). Dashed lines indicate average soma positions. Note the additional VPM innervation domain in L4 for the rare subtype with significant dendritic arborization in L4 (D, right panel; number of predicted VPM boutons in L4: 132 ± 72 [$n = 4$] vs. 29 ± 5.8 [$n = 18$], $P < 0.03$; see Materials and Methods).

distinguish excitatory cell types. We found, however, that within the assumed homogeneous class of excitatory thick-tufted L5 pyramidal neurons, specific branching features of the apical oblique dendrites that were well aligned with the VPM innervation pattern yielded strongly differing subcellular innervation domains for a few of the neurons (see Fig. 7). This could suggest subtypes of thick-tufted L5B neurons. Similarly, L6 pyramidal neurons could be further differentiated by, for example, the extent of their apical dendrites in L5A or L4, possibly predicted by their soma locations (Fig. 5B,C). In both cases, the selectivity of the thalamic projections along the vertical column axis attaches potential functional relevance to the purely morphological finding of variable dendritic branching patterns. Thus, the predicted TC input patterns can contribute to a more precise classification of excitatory neurons within the layers of a cortical column.

Estimation of Absolute TC Bouton Numbers per Neuron

For a quantitative interpretation of the TC input to a cortical column, it was crucial to obtain absolute estimates of the number of potential TC contacts per excitatory neuron in a column. These estimates relied on a few anatomical assumptions, each of which contributes to the uncertainty of the estimate. The total number of boutons projected from one VPM barreloid to one cortical column was estimated using the reported number of neurons in a VPM barreloid (250–300; Land et al. [1995], expected uncertainty range: 200–300, e.g., Varga et al. [2002]), the average axonal path length of one VPM neuron extended in the home column (35 mm, approximate range: 20–50 mm, Oberlaender, Bruno, and Sakmann, unpublished observations), and the average distance between boutons on the TC axons (3 ± 0.2 μm , Oberlaender, Bruno, and Sakmann, unpublished observations; cf. however, 5 μm in

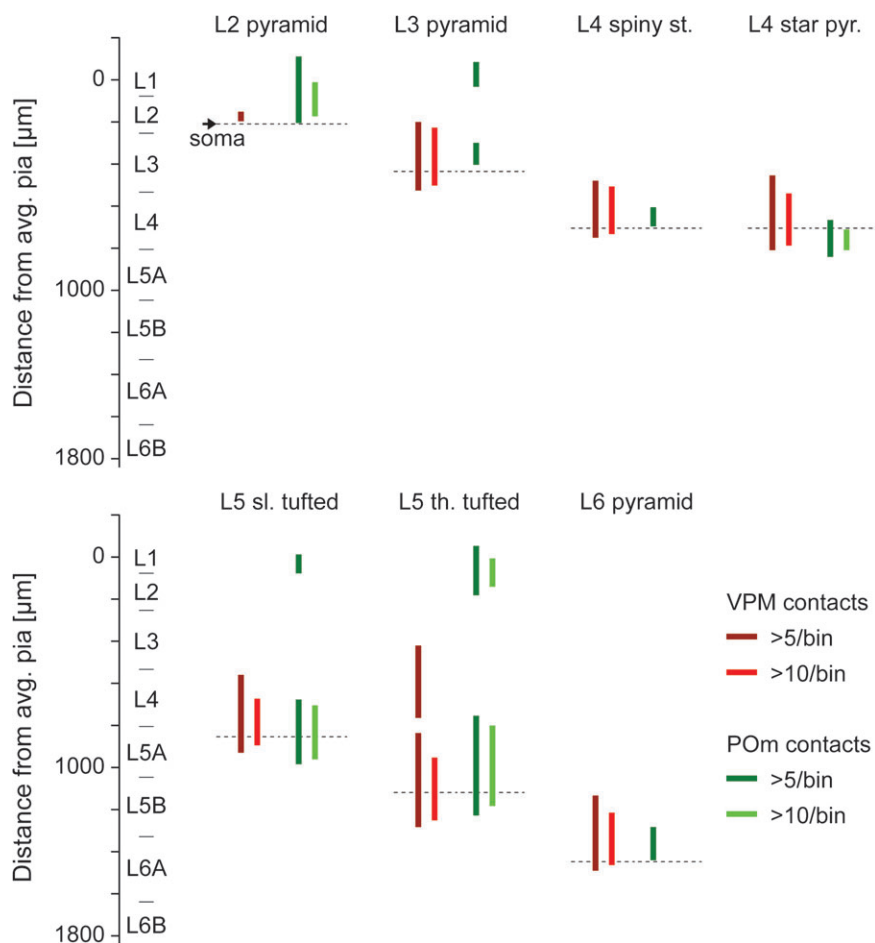


Figure 8. Subcellular thalamocortical innervation domains for different excitatory neuron types in a cortical column. Thalamocortical innervation domains calculated by thresholding the contact z-profiles (cf. Fig. 6) at bouton densities of 5 per bin (effective bin size: 46 μm , cf. Materials and Methods; dark red and dark green for VPM and P0m, respectively) and 10 per bin (light colors). Each neuron type showed a unique combination of subcellular thalamocortical innervation domains (see Discussion).

mouse; White et al. 2004). The ratio of excitatory neurons in a column was assumed to be 85% (Beaulieu 1993; range: 80–90%). All of these assumptions contribute as multiplicative factors to the estimates of absolute bouton numbers; thus, the combined uncertainty for all estimates of absolute bouton numbers is approximately $\pm 30\%$. In addition, the ratio of excitatory neurons is inhomogeneous along the vertical column axis (e.g., Beaulieu 1993). However, compared to the uncertainties in the assumptions about the number of total TC boutons, this variability in the ratio of inhibitory neurons (range: 10–20% for different cortical layers) is of less significance.

We used an estimate of the total axonal path length of the average VPM neuron extended in the somatotopically aligned home column (see above, Materials and Methods). VPM neurons also project to neighboring columns and the septa between barrels in cortex; these projections can amount to as much as 40% of the axonal path length (Oberlaender, Bruno, and Sakmann, unpublished observations). Therefore, the absolute estimates of the number of potential TC contacts per excitatory neuron reported here are lower estimates of the total TC input to a single neuron. To account for TC input from other than the somatotopically aligned barreloid, the reported numbers (Tables 1 and 2) would have to be increased by approximately 10–30% (assuming that dendrites of septal

neurons contribute $\sim 40\text{--}50\%$ to the total dendritic length in barrel cortex, see Meyer et al. [2010] and Materials and Methods). Thus, approximately half of the additional VPM projections could target neurons within the home column.

The Number of VPM Boutons per Neuron in Comparison to Previous Studies

First accounts of TC projections in primary sensory areas concluded that the primary target of TC axons were neurons in layer 4 (Lorente de No 1922, 1938, 1992; Jones and Powell 1970; Killackey 1973). Subsequent ultrastructural analysis using electron microscopy of combined axon degeneration (without differentiation between VPM and P0m) and Golgi staining or anterograde transporter labeling demonstrated, however, that also deep L3 pyramidal neurons and L5 and L6 pyramidal neurons were contacted by TC afferents within L4 (White 1978, 1989; Keller et al. 1985). It was concluded (Peters 1979; White 1979) that all neurons extending their dendrites to L4 received thalamic input. Furthermore, the fraction of putatively excitatory (asymmetric) synapses that was made by TC afferents was estimated to be as high as 18–23% in mouse barrel cortex (Hersch and White 1981, 1982; White and Hersch 1981; Keller et al. 1985; White 1989), while much lower

numbers were obtained for visual cortex (e.g., Garey and Powell 1971; da Costa and Martin 2009).

Our findings quantitatively confirm that 1) all neurons in a column potentially receive substantial TC input (90–580 boutons per neuron) and that 2) the fraction of potential TC synapses per neuron is at least 17% for spiny stellates in L4 (range: 4–17% for all excitatory neuron types, see Table 1; note that this is a lower bound estimate because we did not calculate TC input from other than the somatotopically aligned barreloids, see above). However, our data suggest that a substantial fraction of these potential TC synapses is not made within L4 but in the supra- and infragranular parts of the TC projections. For example, we estimate only 20–30% of all VPM boutons on an L5 thick-tufted pyramidal neuron to be made in L4 (~55 of ~300; Table 2 and Figs 6 and 8, or ~90 of ~300, Supplementary Table 2), while the remaining contacts would be established mainly in L5. Our data do not support significant monosynaptic VPM input to L1 (reported by Oda et al. [2004], see, however, Rubio-Garrido et al. [2009]).

TC Projections from VPM versus POM

We used the peak bouton density to calibrate VPM and POM projection densities (see Materials and Methods). This allowed to quantitatively compare the amount of thalamic projections converging on cortical neurons from either one or both thalamic nuclei. We found that the estimated number of VPM boutons was remarkably similar for L3, L4, and L5 excitatory neurons (estimated range: 150–299 VPM boutons per neuron; Table 1 and Fig. 5B). L6 pyramidal neurons were still predicted to receive 131 ± 66 VPM boutons per neuron, and only L2 pyramidal neurons are predicted to have virtually no VPM input (16 ± 14). The intertype variability was higher for the number of POM boutons per neuron type, with L5 slender-tufted (269 ± 121) and L5 thick-tufted (277 ± 67) pyramidal neurons potentially receiving about 4–5 times as many synapses as the other pyramidal neurons (range: 57–75). L5 slender-tufted pyramidal neurons were predicted to be the most densely POM-innervated neurons, with a synapse distance of 21 μm and a ratio of POM-occupied spines of 9%. Our data predict that thick-tufted L5 pyramidal neurons receive equal amounts of TC input from VPM and POM (with the exception of the putative subtype that has an L4–VPM domain as described in Fig. 7), while the TC input to all other cortical neurons is either VPM dominated (L3, L4, and L6) or POM dominated (L2 and L5 slender tufted).

Comparison with Estimates Derived from Physiological Measurements

The different patterns of TC innervation reported here are based on calculating bouton–dendrite overlap profiles, making the simplifying assumption of locally random synaptic connectivity between TC axons and the dendrites of cortical neurons (Peters 1979; White 1979, 1989; see Shepherd et al. 2005; Stepanyants and Chklovskii 2005 for a critical account). The results reported here therefore represent “potential” synaptic connectivity and provide first quantitative bounds on the amount of TC innervation of cortical neurons. Whether the overlap regions that we calculated here indeed correspond to actual synaptic contacts needs to be verified by, for example, labeling synaptic contacts simultaneously with markers specific for the pre- and postsynaptic compartments or by electron

microscopic methods suitable to image large volumes like an entire layer of a cortical column (for a review, see Helmstaedter, Briggman, et al. 2008).

Qualitatively, the fact that, in response to a whisker deflection, reliable, short onset latency (8–11 ms) excitatory postsynaptic potentials (EPSPs) are recorded in all cortical layers except in L2 (18 ms) (Brecht and Sakmann 2002; Brecht et al. 2003; Manns et al. 2004) renders it likely that the bouton–dendrite overlap domains described here represent indeed functional TC innervation domains as we found significant VPM innervation for all neuron types except for L2 (Figs 5–8 and Table 1). The onset latency of the earliest EPSP component in L3 pyramidal neurons (9.6 ms), for example, is not significantly different from that of the earliest component of EPSPs in L4 spiny stellate neurons (8.0 ms) (Brecht and Sakmann 2002; Brecht et al. 2003). This argues for significant monosynaptic TC activation of both neuron types, which is in agreement with our anatomical data (222 ± 44 vs. 188 ± 23 [mean \pm standard error of mean] VPM boutons per neuron, respectively). It is, however, important to note that high bouton numbers per neuron or dense innervation domains close to the soma do not directly imply higher activation since release probability may be different for different connections (Markram et al. 1998; Reyes et al. 1998; Reyes and Sakmann 1999) and since intracellular mechanisms may significantly modify the effect of a single synaptic contact depending on the location along the dendritic tree (see below).

Previous quantitative estimates of VPM innervation were based on inferences of VPM convergence onto excitatory L4 neurons (based on the fraction of connected pairs; Bruno and Sakmann 2006) and the average number of release sites for a single VPM–L4 connection (Gil et al. 1999). This suggested that the maximum possible number of VPM release sites that could contact a spiny stellate cell is approximately 600. Our lower bound anatomical estimate of the number of contacts (~200 VPM boutons per L4 neuron, cf. Table 1) lies well below this upper bound estimate for the number of release sites. This could suggest possible multiquantal release at VPM boutons. More recent anatomical data seem to support the order of magnitude of approximately 200 VPM boutons per L4 neuron (Schoonover and Bruno, in preparation).

Recently, the input to cortical excitatory neurons has been studied using the expression of a light-activatable cation channel (Nagel et al. 2003) in populations of putatively presynaptic axons; among these were TC afferences (Petreanu et al. 2009). It was concluded that L5B pyramidal neurons lack significant POM input in disagreement with the anatomical predictions presented in this study (Figs 6–8 and Tables 1 and 2). Several reasons may contribute to this discrepancy. Our data suggest that apical tuft input could be a major factor for the excitation of L5 thick-tufted pyramidal neurons by POM (25% of all POM boutons per L5 thick-tufted pyramid; Table 2, Supplementary Table 2). The detection of apical tuft input is, however, hard to achieve in L5 thick-tufted neurons given their low input resistance, the lack of voltage control of distant dendrites in the setting of a somatic recording, and the strong attenuation of distally evoked spikes along the dendritic tree (Larkum et al. 2009). Even for basal dendrites the attenuation of synaptic input can be substantial (Nevian et al. 2007). In addition, the contribution of active dendritic mechanisms like calcium APs to the somatic current signal (Larkum et al. 2009) is difficult to assess. It remains to be shown by ultrastructural

analysis whether thick-tufted L5B neurons receive significant POm input as predicted by our data.

Functional Implications: Capacity for Coincidence Detection by Pyramidal Neurons

The highest number of potential VPM contacts is received by thick-tufted neurons in the infragranular layer (L5), whereas contact density is highest for spiny cells in the granular layer (Table 1). In vivo recordings of AP responses have shown that L5 thick-tufted pyramidal neurons respond to a whisker deflection as rapidly as L4 spiny stellate neurons do but, on average, with a higher number of APs (de Kock et al. 2007). Thick-tufted pyramidal neurons are sensitive to coincident input to different dendritic compartments and typically respond with AP bursts to multilayer synaptic input (Larkum et al. 1999; Larkum and Zhu 2002). These observations are surprising, given the fact that input resistance of L5 thick-tufted pyramidal neurons is considerably lower than that of spiny stellates and because of the strong attenuation of post-synaptic potentials (PSPs) along their basal dendrites (Nevian et al. 2007). One factor that could contribute to the higher responsiveness of thick-tufted pyramidal neurons is their multiple innervation by VPM on their basal and apical oblique dendrites, respectively (Figs 6–8 and Table 2). Our sample contained a subset of cells that had a dense dendritic domain in L4, putatively enabling in particular those neurons to integrate VPM input at 2 dendritic compartments (Fig. 7).

In addition, coincident input to basal and apical oblique dendrites from VPM and to apical oblique and apical tuft dendrites from POm could generate bursts of APs, as observed in vitro and in vivo (Larkum et al. 2001; de Kock et al. 2007; Schaefer et al. 2007; de Kock and Sakmann 2008). Larkum and colleagues found that, in distal tuft dendrites of L5 thick-tufted pyramidal neurons, attenuation of NMDA-mediated spikes is strong, but they estimated that approximately 100 tuft synapses can be sufficient to trigger a calcium spike at the apical initiation zone (Larkum et al. 2009). This is approximately 25% more than the 68 POm boutons and 12 VPM boutons per tuft predicted as a lower bound by the present study (Table 2), indicating that synchronous POm activity could be close to the threshold of initiating active propagation of distal inputs to the axon initial segment for integration with proximal input (see also Larkum et al. 2004). As discussed above, this nonlinear transmission of apical tuft input to the soma may explain the substantial discrepancy between our predictions and physiological data based on the activation of, presumably, a few POm afferents (Petreanu et al. 2009).

A dual TC innervation pattern is also suggested for pyramidal neurons located in L3. These cells are potentially innervated by VPM at their basal dendrites and potentially receive input from POm at their apical tufts in L1 (Figs 6 and 8). L2/3 pyramidal neurons can generate dendritic spikes and integrate distal and proximal synaptic input, which, however, only results in brief “bursts” of 2 APs in vitro (Larkum et al. 2007). AP bursts from L3 pyramidal neurons are in fact infrequent and consist of only 2 APs in in vivo recordings from the anesthetized cortex (de Kock and Sakmann 2008). In the awake animal, however, they become more frequent and can consist of 3 APs (de Kock and Sakmann 2008). This difference in the awake animal could, for example, result from increased POm activity, which could be mediated by arousal (Trageser and Keller 2004; Trageser

et al. 2006) or from reduced inhibition at the distal dendrites (see Larkum et al. 2007). To investigate these potential effects of TC innervation in vivo, experiments are needed where L1 excitation in awake animals is measured (Kuhn et al. 2008; Murayama and Larkum 2009; Murayama et al. 2009).

A dual TC innervation, distributed over multiple dendritic compartments, is also suggested for slender-tufted pyramidal neurons in L5. Both their basal and their apical dendrites overlap with POm bouton domains. The early compound PSP evoked by a whisker deflection also has, however, a short latency (~10 ms) (Manns et al. 2004). Presumably, the VPM projections to apical oblique dendrites mediate this early-onset PSP (Figs 6 and 8). APs in L5 slender-tufted pyramidal neurons are, however, evoked with a relatively long delay compared with other layers (de Kock et al. 2007), suggesting that the early-onset synaptic input is either weak or attenuated by rapid inhibition. The APs occurring later could then be evoked by the L4 spiny stellate-to-L5 slender-tufted pyramid pathway (Feldmeyer et al. 2005; Schubert et al. 2006), which would represent a disynaptic input from VPM. Slender-tufted pyramidal neurons could then be sensitive to coincident excitation conveyed by this disynaptic VPM pathway (to their basal and proximal apical oblique dendrites) and inputs from the POm pathway that excite their basal and apical oblique dendrites and the apical tufts in L1. Finally, the excitation of slender-tufted cells is increased during active whisking (Kerr et al. 2007; de Kock and Sakmann 2009), suggesting increased input from VPM and potentially direct input from POm in the awake behaving condition.

Conclusions

The anatomical data suggest, in combination with the synchrony of APs in VPM cells (Bruno and Sakmann 2006), 2 limiting cases for an “effective” TC excitation of the cortex. On the one hand, it is based on a focal and dense input to a compact dendritic compartment, as has been quantified for L4 spiny stellate neurons. This neuron type is thought to be the main recipient of sensory input to a column in somatosensory cortex (Armstrong-James et al. 1992; Douglas and Martin 2004). On the other hand, effective excitation in the vibrissal cortex is also based on a spatially distributed input to several anatomically and electrically well-separated dendritic compartments (Larkum and Zhu 2002). This is the case for L5 thick-tufted pyramidal neurons. Here, the effective TC activation relies on the generation of active dendritic excitation by distributed synaptic input (Larkum et al. 2001).

In combination with data on functional geometry of TC connectivity (i.e., paired recordings; Bruno and Sakmann 2006), the quantitative data on thalamic innervation patterns of the major excitatory neuron types will serve as a basis for simulations of the signal flow through a cortical column in silico (Helmstaedter et al. 2007).

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Supplementary Material

Supplementary material can be found at: <http://www.cercor.oxfordjournals.org/>

Notes

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