

## ORIGINAL ARTICLE

# Different Patterns of Cortical Inputs to Subregions of the Primary Motor Cortex Hand Representation in *Cebus apella*

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## Abstract

The primary motor cortex (M1) plays an essential role in the control of hand movements in primates and is part of a complex cortical sensorimotor network involving multiple premotor and parietal areas. In a previous study in squirrel monkeys, we found that the ventral premotor cortex (PMv) projected mainly to 3 regions within the M1 forearm representation [rostro-medial (RM), rostro-lateral (RL), and caudo-lateral (CL)] with very few caudo-medial (CM) projections. These results suggest that projections from premotor areas to M1 are not uniform, but rather segregated into subregions. The goal of the present work was to study how inputs from diverse areas of the ipsilateral cortical network are organized within the M1 hand representation. In *Cebus apella*, different retrograde neuroanatomical tracers were injected in 4 subregions of the hand area of M1 (RM, RL, CM, and CL). We found a different pattern of input to each subregion of M1. RM receives inputs predominantly from dorsal premotor cortex, RL from PMv, CM from area 5, and CL from area 2. These results support that the M1 hand representation is composed of several subregions, each part of a unique cortical network.

**Key words:** connections, hand, motor cortex, network, parietal cortex

## Introduction

One common finding across neuroanatomical and functional studies is that the primary motor cortex (M1) is not uniform, but may be comprised of subregions with distinct characteristics. For example in owl monkeys, the caudal part of M1 has larger pyramidal cells than the rostral part (Stepniewska et al. 1993; Preuss et al. 1996) and, whereas the hand representation extends across both regions, current intensity to evoke movements with intracortical microstimulations (ICMS) tends to be lower in

caudal M1 (Preuss et al. 1996). In macaque monkeys, it is also in the caudal part of M1 that most neurons with monosynaptic projections onto motoneurons controlling hand and arm muscles are located (Rathelot and Strick 2006, 2009).

The hand representation of M1 is part of an extensive network of cortical areas that interact for the production of arm movements. Several premotor as well as somatosensory areas send projections to M1 (Stepniewska et al. 1993; Dum and Strick 2005), providing a substrate by which they can modulate the

activity and outputs of M1 neurons (Tokuno and Nambu 2000; Davare et al. 2008; Prabhu et al. 2009; Ziluk et al. 2010). Neuroanatomical tracing studies in New World monkeys also often report that the distribution of these cortical inputs in M1 is not homogeneous. Projections from somatosensory areas are denser in the caudal portion of the M1 hand representation and sparser in rostral M1 (Stepniewska et al. 1993, 2006). Interhemispheric projections seem to follow an inverse pattern. They are located primarily in the rostral part of M1, and are virtually absent in the caudal part of the hand representation (Gould et al. 1986; Dancause et al. 2007). Inputs from ipsilateral premotor areas have also been proposed to preferentially target rostral M1 (Stepniewska et al. 1993, 2006; Dum and Strick 2005; Dancause, Barbay, Frost, Plautz, Stowe, et al. 2006).

In a previous study in squirrel monkeys, we analyzed the topographic distribution of projections from the ipsilateral ventral premotor cortex (PMv) in relation to the M1 arm representation and found a more complex pattern than expected (Dancause, Barbay, Frost, Plautz, Popescu, et al. 2006). PMv's projections were not simply primarily targeting the rostral M1 but rather formed 3 clusters [rostro-medial (RM), rostro-lateral (RL), and caudo-lateral (CL)], leaving only the caudo-medial (CM) region of M1 receiving few projections. These results extend the concept of M1 subdivisions beyond the rostro-caudal axis and suggest that PMv has a very specific pattern of connections with 4 regions within M1. One possibility is that other premotor and sensory areas also preferentially project to specific regions within M1, altogether forming multiple specialized modules, each with a distinct pattern of cortical connections. This idea was recently supported in a study in New World monkeys using long-trains of stimulation to identify "functional zones" (i.e., reach, defense, and grasp) within the frontal cortex and the posterior parietal cortex (Gharbawie, Stepniewska, Kaas 2011). Injections of tracers in each zone revealed a unique pattern of connections. For example, the M1 grasp zone had few connections with S1 and was densely connected with the lateral portion of the posterior parietal cortex, where the parietal grasp zone is located. In contrast, the M1 defense zone had dense connections with S1 and had broader connections in the parietal cortex, with both the defense and grasp zones.

In the present set of experiments, our goal was to study the topographic specificity of cortical projections to M1 in *Cebus apella*. In particular, we wished to directly compare the pattern of ipsilateral inputs to the 4 potential subregions (RM, RL, CM, and CL), within the M1 hand representation highlighted in our previous studies (Dancause, Barbay, Frost, Plautz, Popescu, et al. 2006). We found that the proportion of input from each distant area of the sensorimotor cortex substantially varied across M1. These results show that subregions within the M1 hand representation have a unique pattern of cortical inputs from premotor and somatosensory cortical areas. We propose that the M1 hand area is composed of multiple integration modules that process and integrate largely different cortical inputs and could support specialized aspects of motor functions.

## Materials and Methods

### Surgical Procedures

Three naïve young adult female capuchin monkeys (*C. apella*), with weight ranging from 1.2 to 3.5 kg, were used in the present study. Our experimental protocol conformed to the guidelines of the Canadian Council on Animal Care and was approved by the Comité de Déontologie de l'Expérimentation sur les Animaux of the Université de Montréal.

All surgical procedures were conducted on the left hemisphere under aseptic conditions. Anesthesia was induced with an intramuscular injection of Ketamine (Ketaset®; 15 mg/kg) and transitioned to isoflurane (~2% in 100% O<sub>2</sub>). The animal was placed in a stereotaxic frame, premedicated with an intramuscular injection of Dexamethasone 2 (Vetoquinol®; 0.5 mg/kg). Lactated ringer's solution was delivered intravenously (10 mL/kg/h), and pulse rate, respiration rate, arterial oxygen saturation, and body temperature were monitored and documented throughout the surgery. A dose of Mannitol 20% (1500 mg/kg) was given intravenously prior to the craniotomy to prevent swelling of the brain.

A craniotomy and durectomy exposed the forelimb areas of M1, PMv, dorsal premotor cortex (PMd), supplementary motor area (SMA), as well as the primary somatosensory cortex (S1) and the lateral portion of area 5. The cortex was covered with warm silicone oil and digital photographs of the exposed brain surface were taken. Digital pictures were imported in a technical illustration software (Canvas, ADC systems). A grid (250 µm resolution) was overlaid on the photomicrograph and used to guide electrode penetrations during electrophysiological mapping. For the collection of electrophysiological data, isoflurane was withdrawn and anesthesia maintained with intravenous injections of Ketamine (20 mg/kg/h) and Diazepam (Valium; 0.01 mg/kg/h). ICMS mapping techniques were used to identify the location of motor representations (Nudo et al. 1992; Dancause, Barbay, Frost, Zoubina, et al. 2006; Dancause et al. 2007). Multiunit recording techniques were used to locate S1 (areas 1 and 2) and area 5 (Dancause, Barbay, Frost, Plautz, Stowe, et al. 2006; Padberg et al. 2007, 2010). After physiological data collection, neuronal tracers were injected in M1. At the end of the surgery, monkeys received a subcutaneous injection of Carprofen (Rimadyl®; 10 mg/kg) and Enrofloxacin (Baytril®; 5 mg/kg) and intramuscular injection of buprenorphine (Temgesic®, 0.01 mg/kg). The recovery of the monkey was closely supervised and the drug regiment ensued for 3 days. At the end of the tracer incubation period of 21 days, the animal was killed with a lethal dose of pentobarbital and perfused, and the brain was processed for tissue sectioning.

### Electrophysiological Motor Mapping

In all 3 animals, we identified the location of the hand representation in M1, PMd, and PMv with ICMS. SMA's hand representation was also located with ICMS in 2 animals (CB-6 and CB-8). A glass-coated tungsten microelectrode (impedance ~1 MΩ; FHC Bowdoin, ME, USA) mounted on a micromanipulator (David Kopf Instruments, model 2662, Tujunga, CA, USA) was lowered perpendicular to the cortex. Cortical maps were derived using standard ICMS trains (Mansoori et al. 2014; Touvykine et al. 2015). A stimulation train consisted of 13 monophasic cathodic square pulses (0.2 ms duration; 3.3 ms interpulse interval) and trains were delivered at 1 Hz. At each cortical site, electrical stimulation was applied at different depths (1750–6000 µm) to identify the regions at which responses were evoked with the lowest current intensity. For M1, PMd, and PMv, the maximal responses were found between 1750 and 2500 µm and for SMA between 3000 and 4000 µm. Microelectrode interpenetration distances were 1000 µm in M1 and 500 µm in PMd, PMv, and SMA. At each penetration site, once the depth with a maximal response was identified, the movement evoked at threshold current intensity (the current at which movements were evoked by 50% of the stimulation trains) was identified by visual inspection and by palpation of the arm muscles by one experimenter and confirmed by another. This movement was used to construct the motor maps. If no movement was evoked at a maximum

current intensity of 100  $\mu$ A, the site was defined as unresponsive. The evoked movements were categorized using conventional terminology (Gould et al. 1986; Nudo, Milliken, et al. 1996; Dancause, Barbay, Frost, Zoubina, et al. 2006). Sites evoking movements of the digits, wrist, and forearm (pronation and supination) were grouped into the hand representation. As defined, the hand representation in M1 typically forms a contiguous area that is surrounded by nonresponsive sites at the caudal border of M1 and sites evoking movements of the elbow, shoulder, and face at the other borders (Nudo and Milliken 1996; Nudo, Milliken, et al. 1996; Nudo, Wise, et al. 1996; Dancause, Barbay, Frost, Zoubina, et al. 2006). In each monkey, a few cortical sites evoked movements of 2 body parts at threshold current intensity (average = 6 sites per monkey). These cortical sites were included in the category of the most distal movement evoked. For example, if both digit and shoulder movements were evoked, the site was included in the category “digits”.

The surface area for each movement category was calculated with a custom-made program in Matlab (MathWorks, MA, USA; Touvykine et al. 2015). The algorithm expanded each colored dot on the map to neighboring pixels using nearest-neighbor interpolation until all pixels were assigned a color. The total number of pixels with the same color was scaled to the real cortical surface area according to a calibration ruler that was placed on the brain for the digital picture of the cortex.

### Electrophysiological Somatosensory Mapping

To get some insights on the location of the hand representation and borders between areas in the parietal cortex, we conducted somatosensory mapping using multiunit recording techniques (Dancause, Barbay, Frost, Plautz, Stowe, et al. 2006). A glass-coated tungsten microelectrode (impedance  $\sim$ 1 M $\Omega$ ), similar to that used for ICMS, was lowered perpendicular to the exposed cortex at depths varying from 500 to 1000  $\mu$ m. The depth of the electrode was adjusted to obtain the most robust response to stimulation of the receptive field. Signals were filtered, amplified, and played through loud speakers for monitoring of the neuronal activity. Light touch, brushing, tapping, squeezing, and joint manipulation were used to determine the extent of the receptive fields and the modality of the stimulus driving the most vigorous cortical responses (Sur et al. 1982; Dancause, Barbay, Frost, Plautz, Stowe, et al. 2006). We made 1 or 2 medio-lateral row(s) of microelectrode penetrations with 500  $\mu$ m interpenetration distance to identify all 5 digits in area 1 and the medial and lateral boundaries of the hand representation. An additional 1 or 2 rostro-caudal row(s) were made to identify the borders between area

1, area 2, and area 5. We established these borders based on the modality of the stimulus driving the neural responses most effectively, the size and location of receptive fields, and the capacity to drive vigorous neural responses with stimulation of the receptive fields (Padberg et al. 2007, 2010). Because of time restrictions caused by motor mapping, we chose not to conduct extensive documentation of the sensory cortex in our animals. In particular, we did not explore the walls of the central sulcus for the identification of area 3a and 3b. We did not include connections to these areas in our quantitative analyses (see below). In addition, for area 5, we only collected information about the rostral border. The limited physiological sensory mapping data should be kept in mind when drawing conclusions about proportions of projections to M1 from these areas.

### Neuronal Tracer Injections

Once the mapping procedures were completed, the animal was put back on isoflurane anesthesia ( $\sim$ 2% in 100% O<sub>2</sub>) for the injection of 4 neuronal tracers all within the M1 hand area. Based on previous results (Dancause, Barbay, Frost, Plautz, Popescu, et al. 2006), the M1 hand representation was arbitrarily subdivided into 4 quadrants (RM, RL, CM, and CL). One of 4 neuroanatomical tracers [biotinylated dextran amine (BDA; 3000 MW; Invitrogen), Fluoro-ruby (FR; 3000 MW; Invitrogen), Fast blue (FB; Dr ILLIG Plastics GmbH), or Fluoro-emerald (FE; 3000 MW; Invitrogen)] was injected in each quadrant. We alternated the tracer injected in the 4 quadrants of the M1 hand area among the 3 monkeys to minimize any bias that may be caused by the properties of the tracers. A 1- $\mu$ L Hamilton syringe coupled to a microsyringe pump controller (Harvard Apparatus, Holliston, MA, USA) was used for all injections. To label the cortex through all layers of the gray matter, injections were made in 3 boluses at pre-determined depths (1800, 1500, and 1200  $\mu$ m), creating a column labeling all layers of the gray matter. Following each bolus of injection, the syringe was kept in place for 5 min to reduce potential backflow. The amounts of tracer injected at each injection site are provided in Table 1.

### Histological Procedures

Twenty-one days following neuronal tracer injections, the animals were deeply anesthetized with ketamine (Ketaset<sup>®</sup>; 15 mg/kg; intramuscular) and administered a lethal dose of pentobarbital (Euthansol; 100 mg/kg; intraperitoneal). The animals were perfused transcardially with a solution of 0.2% heparin in 0.1 M phosphate-buffered saline (PBS; pH 7.4), followed by 3%

**Table 1** Neuroanatomical tracers efficacy

	Injection location	Tracer	Volume of injected tracer ( $\mu$ L)	Volume of injection core (mm <sup>3</sup> )	Total number of labeled cells	Injection location	Tracer	Volume of injected tracer ( $\mu$ L)	Volume of injection core (mm <sup>3</sup> )	Total number of labeled cells
	RM					CM				
CB-8		BDA	0.4	0.46	2012		FE	0.5	0.3	2765
CB-6		FB	0.2	14.7	33 070		FR	0.4	1.3	4600
CB-3		FR	0.2	2.5	2539		FB	0.2	16.2	22 453
	RL					CL				
CB-8		FB	0.25	19.7	25 134		FR	0.4	1.0	4833
CB-6		FE	0.4	0.36	1939		BDA	0.4	1.4	1222
CB-3		BDA	0.2	2.9	1955		FE	0.2	0.7	806

BDA, biotinylated dextran amine; FB, Fast blue; FE, Fluoro-emerald; FR, Fluoro-ruby.

paraformaldehyde in 0.1 M PBS (pH 7.4). The brain was removed and the cerebral cortex isolated. The fronto-parietal cortices were then flattened between 2 glass slides (Gould and Kaas 1981; Killackey et al. 1983; Dancause et al. 2005; Dancause, Barbay, Frost, Plautz, Stowe, et al. 2006). The cortical block of tissue was post-fixed for 2 h in a solution of 20% sucrose and 3% paraformaldehyde in 0.1 M PBS, followed by a solution of 20% sucrose with 2% dimethyl sulfoxide in 0.1 M PBS overnight and a solution of 20% sucrose in 0.1 M PBS for 48 h. The cryoprotected tissue was quickly frozen at  $-55^{\circ}\text{C}$  with methyl butane before being stored at  $-80^{\circ}\text{C}$ . Sections were cut tangentially to the cortical surface with a cryostat (thickness 50  $\mu\text{m}$  for CB-3 and CB-6 and 40  $\mu\text{m}$  for CB-8). The sections were collected in series of 6. The first and fourth sections (2/6) were used for the documentation of labeled cell bodies following injection of 1 of the 3 fluorescent tracers (FR, FE, and FB). Those sections were immediately mounted after sectioning, air-dried, and coverslipped the same day. The second and fifth sections (2/6) were processed to examine the cell bodies labeled with BDA. The third section and sixth sections were kept for Nissl staining (1/6) and myelin staining (1/6). Details of our histology protocols for the BDA and myelin staining have been previously published (Dancause et al. 2005; Dancause, Barbay, Frost, Plautz, Stowe, et al. 2006).

### Neuroanatomical Reconstruction and Quantification of Labeled Cells

To study the distribution of labeled cell bodies, we used a neuroanatomical reconstruction system (NeuroLucida, MicroBrightField, Colchester, VT, USA) associated with a microscope (Olympus BX51, Tokyo, Japan). For CB-3 and CB-6 (section thickness 50  $\mu\text{m}$ ), every third section was sampled between depths of 650–1600  $\mu\text{m}$ . In CB-8 (section thickness 40  $\mu\text{m}$ ), every fourth section was sampled between depths of 640–1640  $\mu\text{m}$ . For each injection, a total of 7 sections were reconstructed. In tangential sectioning, the first sections are often incomplete and the cortical gray matter is not of equal thickness across the hemisphere. Thus, the depth of sections provides limited information about cortical layers in which labeled cells are located. However, the systematic sampling across the range of sections that had consistent labeling in all 3 monkeys should reflect the overall pattern of connections for each injection site.

Tangential sectioning facilitated the co-registration of anatomical and neurophysiological data (Xiao and Felleman 2004; Dancause et al. 2005). In reconstructed sections, the location of large blood vessels, corresponding to large circular holes, was marked as well as other landmarks, such as sulci and tracer injection sites. On the photomicrograph used to construct the physiological maps, the points of entry in the cortex of large blood vessels and the same landmarks were identified. Reconstructed sections and the photomicrograph along with physiological maps were imported into Adobe Illustrator CS5 (Adobe Systems, San Jose, CA, USA) and were aligned using the pattern of blood vessels and landmarks. The co-registered data allowed us to identify the cortical areas that were physiologically defined on the anatomical sections. As only information about the rostral border of area 5 was available, our contour extended caudal and medial from that border along the intraparietal sulcus. The number of labeled cell bodies in each area was calculated with Neuroexplorer, an analysis program linked to NeuroLucida (Dancause, Barbay, Frost, Plautz, Popescu, et al. 2006; Dancause, Barbay, Frost, Plautz, Stowe, et al. 2006). In the present study, we chose not to unfold the central sulcus to retain maximal co-registration of physiological and anatomical data for M1,

area 1, area 2, and area 5. Consequently, areas 3a and 3b were below the deepest sections reconstructed (1600  $\mu\text{m}$ ) and were not analyzed.

For each injection site in a monkey, the number of labeled cell bodies in the various ipsilateral cortical areas was counted as a percentage of the total number of neurons labeled with this tracer in the reconstructed hemisphere. Because the count of labeled cells in M1 was affected by the multiple injection cores, we excluded all cells located in M1 from this calculation. The percentages thus only represent “extrinsic” ipsilateral inputs to M1 (Dancause, Barbay, Frost, Plautz, Stowe, et al. 2006). Results from the 3 monkeys were combined to obtain the final distribution of cells following injections in the RM, RL, CM, and CL subregions within M1. An ANOVA, with monkeys as the statistical units, was conducted on each of the 4 subregions to determine whether or not the percentages differed between areas. Post hoc analyses were conducted to determine which areas significantly differed from the others, using Tukey’s test to correct for multiple testing. In addition, for each monkey, a  $\chi^2$  test was conducted to determine if the distribution of labeled neurons across cortical areas were the same for the 4 subregions of M1. Because only 3 monkeys were used in the present study, we did not report the main effects of the ANOVA but only used post hoc results to guide the description of the pattern of labeling and classify the strength of connections. Nevertheless, the low number of animals should be kept in mind when considering statistical results.

Finally, to compare the pattern of projections of each distant cortical area with the others, we normalized the percentage values according to the following formula:

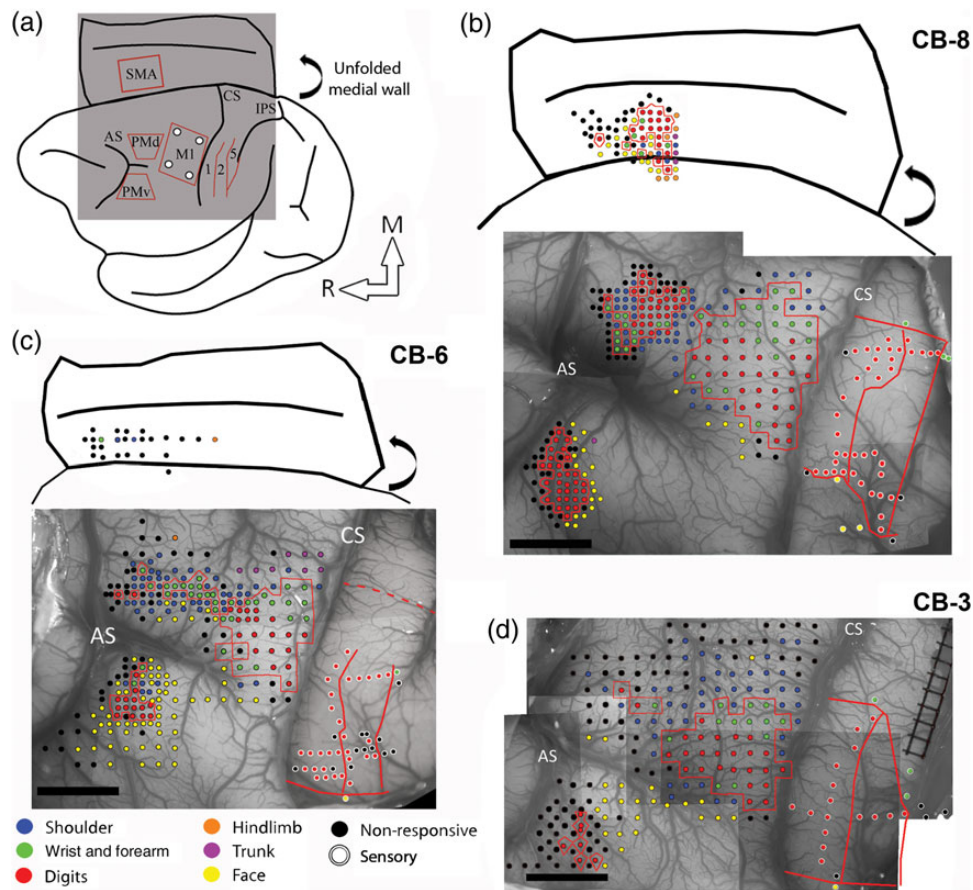
$$\text{Normalized projections to an M1 subregion} = \frac{\% \text{ of labeled cells in area } n}{\% \text{ of labeled cells in area } H}$$

where “area n” is the distant area studied (i.e., PMv, PMd, SMA, area 1, area 2, and area 5) and “area H” is the area with the highest proportion of labeled cells following injections in a given subregion of M1 (RM, RL, CM, and CL). The values obtained vary between 1, for area H with the highest proportion of labeled cells, and potentially 0 in the case of an area with no labeled cells following injections in a subregion of M1. The numbers reflect the relative strength of the projections from area n to the subregion in comparison with the other cortical areas in the sensorimotor network we studied.

## Results

### Hand Representations in Primary Motor and Premotor Cortex of Cebus Monkeys

In the present study using ICMS techniques, we provide the most complete documentation of the hand representations in motor and premotor cortex in Cebus monkeys to date. Figure 1 shows the electrophysiological mapping data from all 3 monkeys. Sites evoking digits, wrist, and forearm formed a contiguous area in M1 that was surrounded by movements of the elbow, shoulder, orofacial, or nonresponsive sites. The M1 hand representation extended over a large cortical area, and comprised many sites evoking digit or wrist movements (Fig. 2). In all 3 monkeys, proximal and nonresponsive sites were typically found at the medial and rostral borders of the hand representation. At the lateral border, orofacial responses were the most frequent. In one animal (CB-6), we also evoked trunk movements at the medial border of the M1 hand area. In the caudal part of M1, responses were



**Figure 1.** Electrophysiological mapping. (a) Cartoon showing a lateral view of the brain and the approximate location of the craniotomy (gray square) and the areas of the frontal and parietal cortex studied with electrophysiological mapping methods. The medial wall is shown unfolded (curved arrow). (b) Motor and sensory mapping data in CB-8. Each colored dot on the digital photograph of the cortex represents a microelectrode penetration site. The evoked movement (frontal cortex) or somatosensory receptive field location (parietal cortex) is color-coded (legend). In the parietal cortex, the same color-code is used for the receptive field location on the body, but the dots have white contours. The hand representations of M1, PMv, PMd, and SMA are outlined with red contours. In the parietal cortex, the medio-lateral extent of the hand representation in S1 and the location of rostro-caudal borders between area 1, area 2, and area 5 are shown with red lines. Note that the location of stimulation sites for SMA are approximated based on the distance from PMd and the depth of the electrode. They are shown over a cartoon representation of the unfolded medial wall to provide an estimated location of the sites in relation to the medial wall's convexity and the cingulate sulcus. The location of these 2 landmarks is based on the anatomical reconstructions of each monkey. (c) Motor and sensory mapping data for CB-6. In this animal, we were able to locate SMA but got limited data from it. As for CB-8, the location of the stimulation sites in SMA is estimated. In S1, we did not physiologically define the medial border of the hand representation. Based on other animals, we estimated the location of this border and show it with a dotted red line. (d) Motor and sensory mapping data for CB-3. In this animal, we did not locate SMA. AS: arcuate sulcus; CS: central sulcus; IPS: intraparietal sulcus; M1: primary motor cortex; PMd: dorsal premotor cortex; PMv: ventral premotor cortex; SMA: supplementary motor area; 1: area 1; 2: area 2; 5: area 5; M: medial; R: rostral. Scale bar = 5 mm.

evoked with low current intensities on the convexity of the rostral bank of the central sulcus. At each tested site next to the central sulcus, we unsuccessfully attempted to evoke movements along the rostral wall (depths of ~2000 to ~4000  $\mu\text{m}$ ; stimulating at steps of 500  $\mu\text{m}$ ), supporting that in Cebus monkeys M1 is largely located on the convexity and that area 3a is situated in the anterior bank of the central sulcus (Felleman et al. 1983; Dum and Strick 2005). The accessibility of the M1 hand representation facilitated motor mapping and allowed us to target our tracer injections to consistent locations in relation to the organization of M1, as defined with ICMS techniques.

Additional ICMS mapping was conducted to locate the hand representations in PMv, PMd, and SMA. The hand representation of PMv was found lateral to the spur of the arcuate sulcus. On each motor map, we calculated the shortest straight line between the borders of the hand representation of M1 and the premotor areas. The average distance between the M1 and PMv hand representations for the 3 monkeys was 6.5 mm. Almost exclusively digit movements could be evoked in PMv and the hand representation

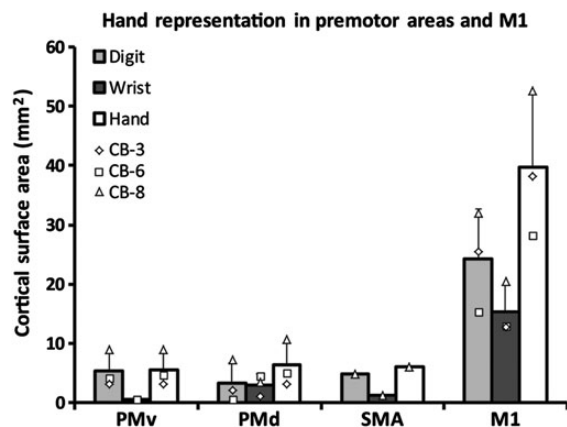
was much smaller than in M1 (Fig. 2). Caudal and medial to the hand representation of PMv, cortical sites evoking orofacial movements were typically found. In some cases, we also evoked trunk, shoulder, or elbow movements. Lateral and rostral to the hand representation of PMv, cortical sites were typically nonresponsive and we did not evoke movements along the caudal wall of the sulcus (depths of ~2000 to ~4000  $\mu\text{m}$ ). Thus, as for M1, it appears that the hand representation of PMv in cebus monkeys is located on the cortical surface. The PMd hand area was located medial to the spur of the arcuate sulcus. It was in close proximity, but separated from the M1 hand representation (average distance 0.7 mm) and thus more difficult to identify. We based our estimation of the border between PMd and M1 on a combination of histological data and ICMS topography and thresholds. In Nissl-stained sections, PMd lacks giant pyramidal cells (Preuss and Goldman-Rakic 1991). In physiological maps, we find cortical sites evoking proximal or orofacial responses that separate cortical sites evoking hand movements in M1 and PMd, and the stimulus intensity to evoke movements in PMd are greater than in M1. The size of the hand

representation in PMd was comparable to PMv. However, a greater proportion of wrist movements were evoked in PMd than PMv (Fig. 2). Nonresponsive and proximal sites were found at the medial, rostral, and lateral borders of the PMd hand area. In addition, some sites eliciting orofacial movements were found at the lateral border. In one case (CB-6), the PMd hand representation was not contiguous, but was rather formed of several isolated clusters separated by nonresponsive sites and sites evoking proximal movements. We were able to locate the SMA hand area in 2 monkeys (CB-8 and CB-6). In comparison with PMv and PMd, responses were elicited at greater depths (~2500 to ~4000  $\mu\text{m}$ ), suggesting that most of the SMA hand representation is located along the medial wall. Based on these 2 monkeys, the distance between the M1 and SMA hand representations was 8.2 mm. We collected enough data to construct a motor map in only one monkey (CB-8). Although this would have to be confirmed in additional animals, it appears that the size of the SMA hand representation is comparable to other premotor areas (Fig. 2). Ventrally, nonresponsive sites surrounded the hand representation. Rostral to the hand area of SMA, in addition to nonresponsive sites, we found several orofacial responses. At both the dorsal and caudal borders of the

SMA hand representation, we found sites from which we could evoke hindlimb, trunk, shoulder, and elbow movements.

### Hand Representations in the Parietal Cortex

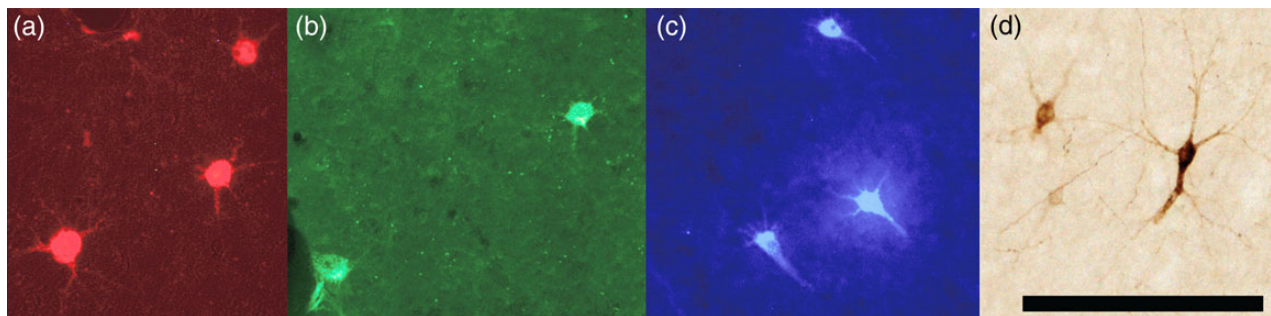
In the parietal cortex, we conducted electrophysiological mapping using multiunit recording techniques (Felleman et al. 1983; Dancause, Barbay, Frost, Plautz, Stowe, et al. 2006; Padberg et al. 2007). One or 2 medio-lateral row(s) of electrode penetrations was done in area 1. The receptive fields of digits were organized somatotopically, with digit 5 found most medial and digit 1 most lateral. Lateral to the hand representation, we found receptive fields on the face and medial to the hand representation, the receptive fields were on the wrist and forearm. Due to time constraints, we could not identify the medial border of hand representation in CB-6 and thus, the location of this border was estimated based on somatosensory mapping data of other Cebus monkeys from our laboratory ( $n = 8$ ). One or 2 rostro-caudal row(s) of electrode penetrations provided insights on the location of borders between area 1, area 2, and area 5. In area 1, neurons reacted to light cutaneous stimulations. In the rostral aspect of area 1, receptive fields were typically on the palm of the hand and were progressively found closer to the tip of digits as we moved the recording electrode more caudal. The transition to area 2 was marked by the occurrence of larger receptive fields that covered several digits and often extended to the palm. Neurons in area 2 also often responded to movement of joints. In area 5, numerous sites were nonresponsive or had activity that was weakly driven by manipulation of the hand and arm. Responsive sites typically reacted to joint manipulation, squeezing of multiple fingers or even the entire hand and forearm of the monkey (Padberg et al. 2007). Based on these results, we were able to approximate the location of medial and lateral borders of the hand representation in areas 1 and 2 and the borders between area 1, area 2, and area 5 for the quantification of labeled cells.



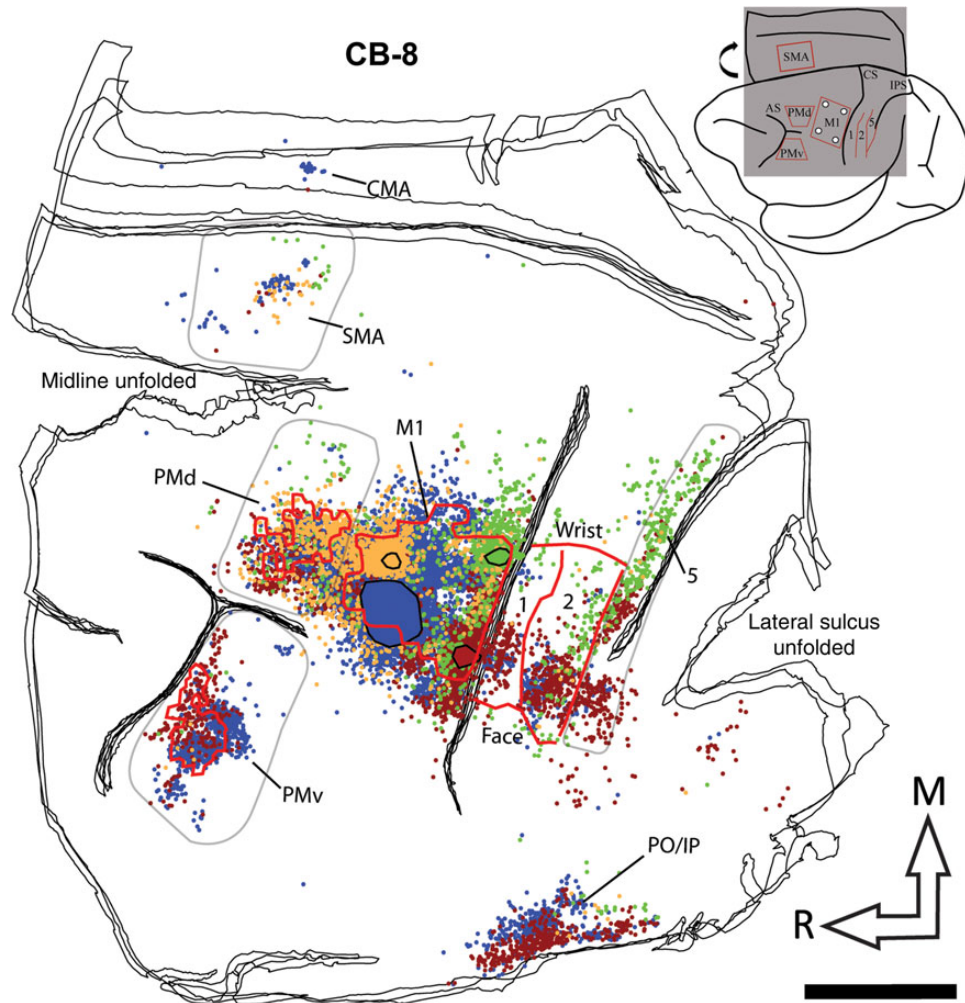
**Figure 2.** Cortical surface area of the hand representation in the primary (M1) and premotor areas of Cebus monkeys. The graph shows the cortical area from which movements of the digits and wrist could be evoked in the different motor areas (mean  $\pm$  SD). The hand representation in PMv ( $n = 3$ ), PMd ( $n = 3$ ), and SMA ( $n = 1$ ) was of comparable size and much smaller than in M1 ( $n = 3$ ). Data from individual monkeys are shown with different symbols displayed on the top of the bars. In all 4 cortical areas, there was a larger territory devoted to movements of the digits in comparison with wrist. However, this difference was smaller in PMd and greater in PMv.

### Distribution of Labeled Cell Bodies in the Ipsilateral Hemisphere

Following identification of the hand area in M1, 4 different neuroanatomical tracers were injected in subregions (RM, RL, CM, and CL). Figure 3 shows examples of neurons labeled by the different tracers. The use of multiple tracers injected in the same animal at physiologically identified locations allowed us to directly compare the pattern of inputs of these subregions of M1 within each animal. Figure 4 shows the anatomical reconstruction of the flattened cortex and labeled cell bodies coregistered



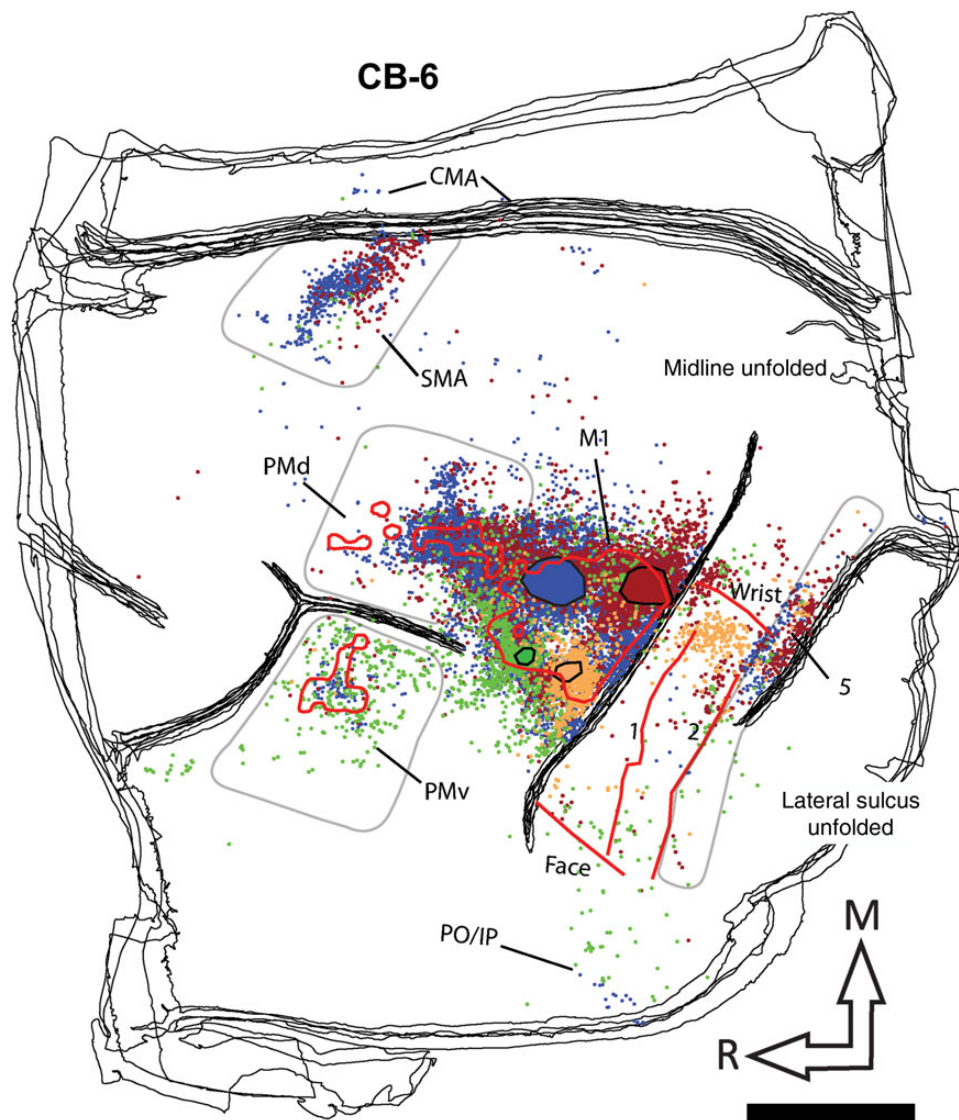
**Figure 3.** Photomicrographs of retrogradely labeled cells in the hemisphere ipsilateral to the injection of neuronal tracer. Photomicrographs of retrogradely labeled cells with (a) FR, (b) FE, (c) FB, and (d) BDA in the ipsilateral hemisphere. Only cell bodies displaying a fully round soma and a minimum of 2 protuberances (considered being either dendrites and/or axon) were considered labeled neurons. A marker was overlaid onto the cell in NeuroLucida (Microbright-Field, Inc.) in the reconstruction of the section. Brightness and contrast were adjusted. Scale bars = 100  $\mu\text{m}$ .



**Figure 4.** Reconstruction of the labeled cell bodies in CB-8. The cartoon (upper right corner) illustrates the approximate extent and location of the sections in relation to the entire brain. The black lines show the outlines of tangential sections of the flattened hemisphere. In the frontal cortex, the location of PMv, PMd, and M1 hand representations are outlined with red contours. Because SMA was mainly located along the medial wall, we could not precisely align it with the anatomical data and thus do not show it on the reconstruction. In the parietal cortex, the medio-lateral extent of the hand representation and the location of the borders between area 1, area 2, and area 5 are shown with red lines. The location and extent of the injection cores in M1 are outlined with large color-coded blobs with black contours (BDA = orange; FE = green; FR = red, and FB = blue). Labeled cells from all 4 tracers are shown throughout the hemisphere with small dots that are colored according to the same colors used for the injection cores. The areas chosen to quantify cell bodies in PMv, PMd, SMA, and area 5 are shown with pale gray contours. Note that the contours are used to show cells that were counted in each area, and do not necessarily represent the extent of the areas. For areas 1 and 2, the contours physiologically defined for the hand representation (red lines) were used to quantify cells. In addition to the physiological data, these areas were defined using myelin and Nissl-stained sections and anatomical landmarks. In myelin sections, area 1 is more lightly myelinated than area 2 (Padberg et al. 2007). In Nissl-stained sections, the premotor cortex lacks giant pyramidal cells (Preuss and Goldman-Rakic 1991). We also used Nissl sections to help define borders between areas in the parietal cortex. Layer IV in area 2 is more densely packed than in area 1 and layers IV and VI in area 5 is more densely packed and darker than area 2 (Padberg et al. 2007). AS: arcuate sulcus; CMA: cingulate motor areas; CS: central sulcus; IPS: intraparietal sulcus; M1: primary motor cortex; PMd: dorsal premotor cortex; PMv: ventral premotor cortex; PO/IPC: posterior operculum/inferior parietal cortex; SMA: supplementary motor area; 1: area 1; 2: area 2; 5: area 5; M: medial; R: rostral. Scale bar = 5 mm.

with the physiological mapping data of CB-8. Figures 5 and 6 show reconstructions of CB-6 and CB-3, respectively. Within M1, the location and extent of the injection cores resulting from each injection are outlined. Each injection labeled many cells within M1 and several cells in all premotor and sensory areas for which we collected electrophysiological mapping data (i.e., PMv, PMd, SMA, area 1, area 2, and area 5). The injection cores of the tracers prevented accurate counts in M1, and we excluded these data from our quantitative analyses. The percentages of labeled cells thus reflect the distribution of “extrinsic” inputs from PMv, PMd, SMA, area 1, area 2, and area 5 to the injected subregions of M1 (Dancause, Barbay, Frost, Plautz, Stowe, et al. 2006). Cells in these areas represented from 78 to 99% (mean = 89.2%) of

the total number of labeled cells in the reconstructed sections following each tracer injection. Additional cells were located along the cingulate sulcus in areas corresponding to the cingulate motor areas (CMAs) and the posterior operculum and inferior parietal cortex (PO/IPC), in areas corresponding to the secondary somatosensory cortex (S2) and parietal ventral cortex. However, the tissue in these regions was incomplete in several sections for each animal, and we thus excluded these areas from our quantitative analyses. A few cells were also inconsistently found in other areas, for example rostral to PMv, between PMd and SMA, or caudal to area 5. Finally, because the central sulcus was not unfolded, labeling in area 3a and 3b was not analyzed in the present study.



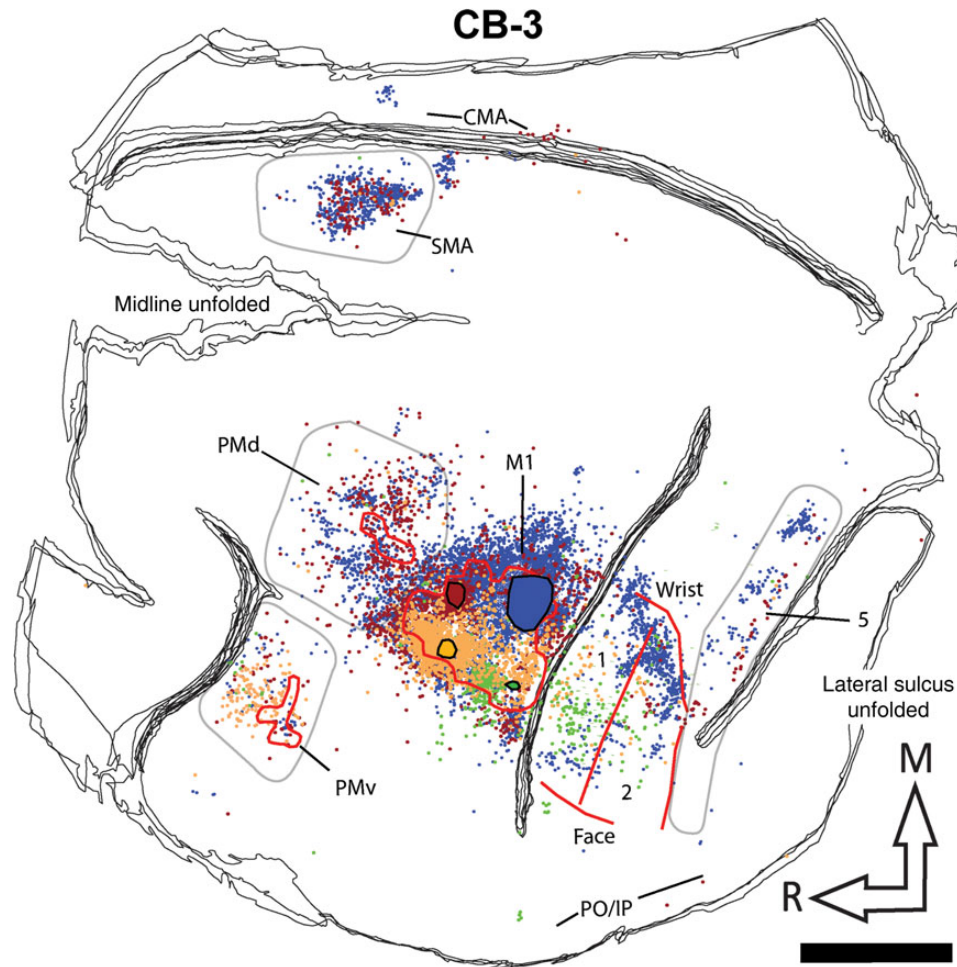
**Figure 5.** Reconstruction of the labeled cell bodies in CB-6. Color codes, contours, and abbreviations are as described in Figure 4. However, note that the location of the different tracers within M1 varies from one animal to another. Furthermore, in this animal, the location of the medial border of the hand representation was estimated. Scale bar = 5 mm.

We first quantified the general pattern of inputs to M1. When pooling the data from all M1 injections ( $n = 12$ ), we found only slightly more labeled cells in premotor (48.4%) than in the somatosensory areas (40.7%). PMd had the highest proportion of labeled cells (23.1%), followed by area 5 (16.5%), PMv (15.8%), area 2 (15.2%), SMA (9.5%), and area 1 (9.0%). Figure 7 shows separately the distribution of labeled cells that resulted from injections in each subregion of M1 [mean  $\pm$  standard deviation (SD)]. As can be observed in the reconstructions (Figs 4–6), there was variability across cases, which may have been caused by the efficacy of the tracers, or different sizes of the effective injection cores or intrinsic variability of connections across monkeys. The normalization of the quantity of cells in each area to the total number of labeled cells reduced this variability and a relatively consistent pattern of labeling was observed across animals. A first striking feature of the distribution of labeled cells bodies after the different injections is that both rostral injections (RM and RL) labeled many more cells in the premotor areas (78.4

and 68.4%, respectively) than in parietal areas. After injections in RM ( $n = 3$ ), most of these cells were in PMd (55.8%), followed by SMA (14.7%). When comparing the proportion of labeled cells across areas, we found that only the proportion of labeled cells in PMd was significantly greater than in all other cortical areas ( $P < 0.05$ ). Following injections in RL ( $n = 3$ ), most labeled cells were found in PMv (44.8%), followed by PMd (15.5%), and only the proportion of labeled cells in PMv was significantly greater than in any other cortical area ( $P < 0.05$ ).

In contrast to rostral injections, both caudal injections (CM and CL) labeled more cells in the parietal cortex (58.4 and 73.7% respectively). After injections in CM, the greatest proportion of labeled cells was in area 5 (32.0%), followed by PMd (15.9%). In addition, many labeled cells were also found in SMA (14.5%), area 2 (14.4%), area 1 (12.0%), and fewer in PMv (3.6%). These proportions were not significantly different across areas with the exception of the proportion of labeled cells in area 5, which was greater than in PMv. Thus, injections in the CM resulted in





**Figure 6.** Reconstruction of the labeled cell bodies in CB-3. Color codes, contours, and abbreviations are as described in Figure 4. Scale bar = 5 mm.

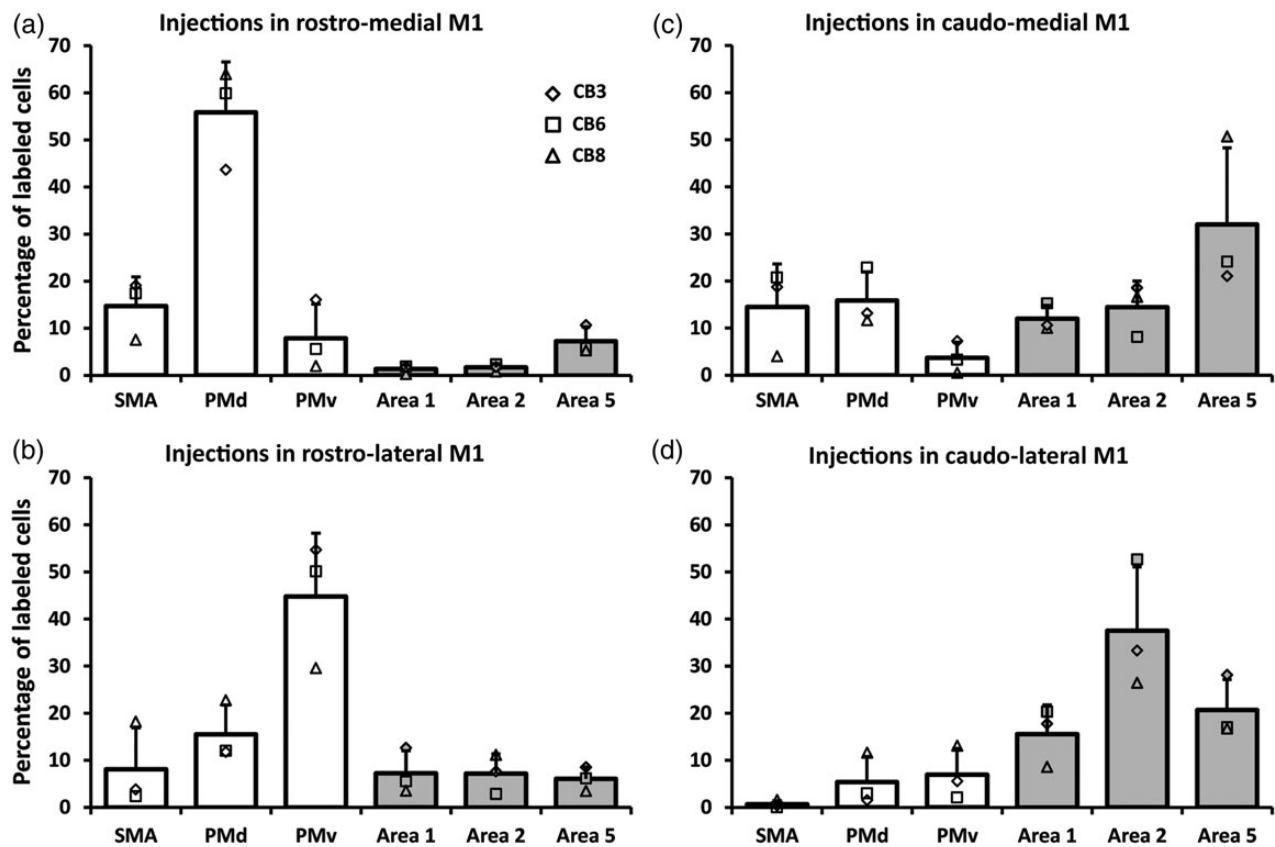
a more even distribution of labeled cells across cortical areas. Finally, after injections in CL, most of the labeled cells were in area 2 (37.5%), followed by area 5 (20.6%) and area 1 (15.6%). The proportion of cells in area 2 was significantly greater than all other areas, except area 5. The proportions of labeled cells in all other areas were not significantly different. For each monkey, a  $\chi^2$  test confirmed that the distribution of labeled neurons across cortical areas was different following the injections in the 4 subregions of M1 ( $P < 0.0001$ ).

Next, we wished to get better insights on the relative projections from each distant cortical area to the hand representation of M1. To do so, for each injection site, the percentage of labeled cells found in a sensorimotor area was normalized to the predominant connection following this injection (Fig. 8a). We separated the strength of projections into 3 categories: Major (normalized projection value  $>0.66$ ), Moderate (between 0.66 and 0.34), and Minor ( $<0.34$ ). This analysis shows that PMd had the broadest pattern of projections, sending major projections to the RM hand representation of M1, but also moderate projections to RL and CM. PMv's pattern of projection was more segregated, sending major projections to RL and minor projections to other subregions of M1. Among premotor areas, the relative strength of SMA projections across M1 subregions was the weakest. It only sent moderate projections to CM and minor projections to other regions of M1. All 3 parietal areas only sent minor

projections to rostral M1 (both RM and RL). Area 1 had the weakest projections to the M1 hand area. It sent moderate projections to CL and CM. Area 2 sent major projections to CL and moderate projections to CM. In contrast, area 5 sent major projections to CM and moderate projections to CL. Figure 8b shows the pattern of projections from PMd, PMv, SMA, area 1, area 2, and area 5 to different subregions in the M1 hand representation.

## Discussion

Our previous work highlighted that PMv projections to M1 are not uniformly distributed but rather specifically target subregions (Dancause, Barbay, Frost, Plautz, Popescu, et al. 2006). In the present study, our objective was to further understand the organization of ipsilateral cortical inputs in the M1 hand area. In each monkey, we injected various retrograde neuroanatomical tracers in 4 regions of the M1 hand representation and quantified the labeled cell bodies across several ipsilateral premotor and somatosensory areas. Injections in each subregion resulted in a unique pattern of cortical labeled cells. These data support a view that M1 is composed of several modules each receiving inputs of unequal strength from diverse motor and sensory cortical areas of the ipsilateral hemisphere. Neurons in these M1 subregions are thus part of specific networks in which they can primarily interact



**Figure 7.** Proportion of labeled cell bodies in ipsilateral hemisphere. Each panel shows the average proportion of labeled cells in the different premotor and parietal areas (areas 1, 2, and 5) in the reconstructed sections (mean  $\pm$  SD;  $n = 3$ ). Data from individual monkeys are shown with different symbols displayed on the top of the bars. (a) Following injections in the RM, most labeled cells were in the premotor areas and we found the greatest number of cells in PMd. (b) Injections in the rostro-lateral (RL) M1 also labeled more cells in the premotor cortex but the area with most labeled cells was PMv. (c) Injections in the CM labeled more cells in area 5. However, the proportion of cells in other areas was more evenly distributed. (d) Finally, injections in the CL labeled more cells in area 2, followed by area 5.

with distinct interlocutor(s) for the production of motor outputs involved in hand movements.

### Motor and Somatosensory Areas in the Capuchin Monkey

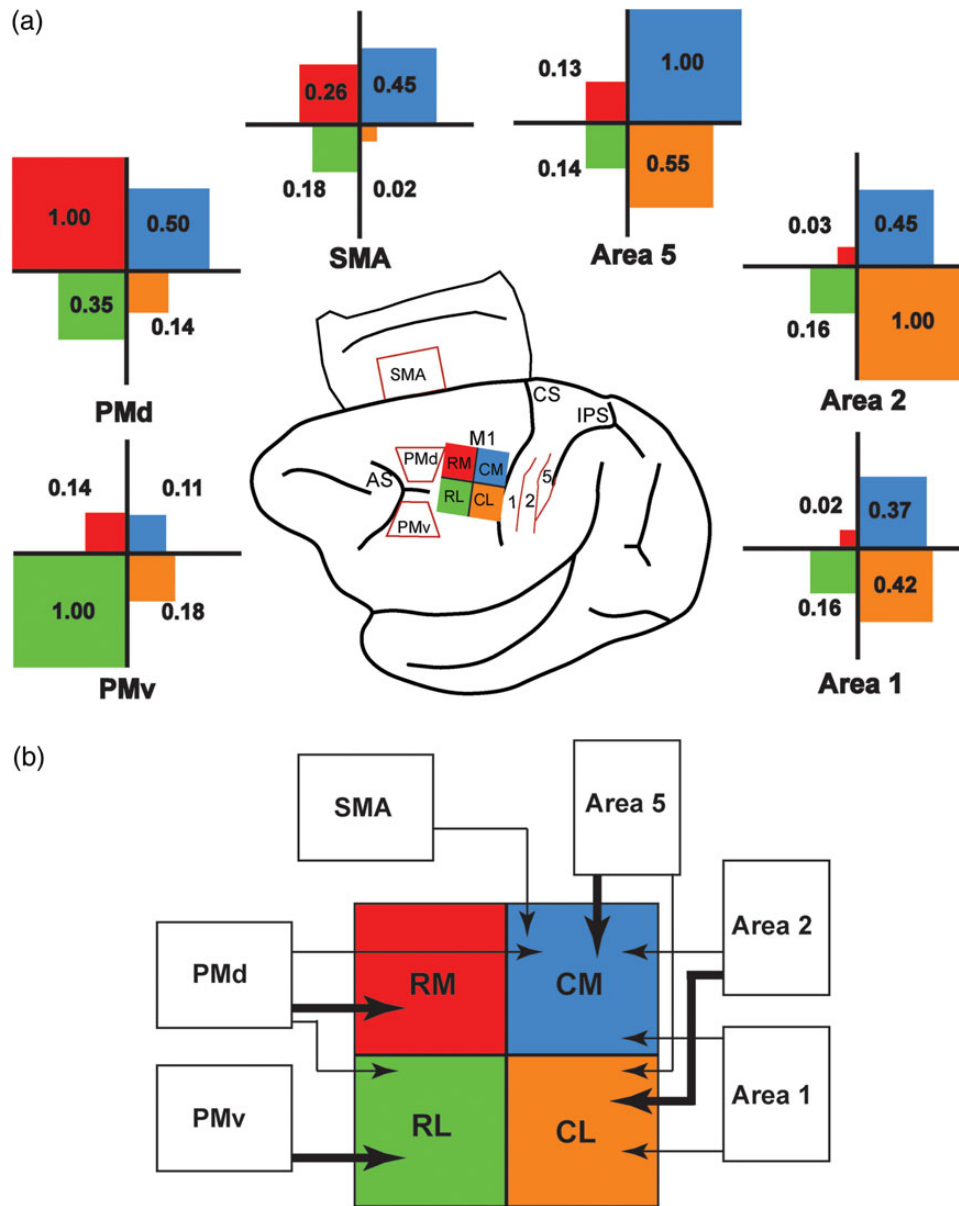
We chose to conduct our study in *Cebus* monkeys, a New World species capable of pseudo-opposition of the thumb and index (Christel and Frigaszy 2000) that can use tools such as branches and stones, approximating the function of macaque monkeys or even some great apes (Visalberghi et al. 1995; Moura and Lee 2004; Evans and Westergaard 2006). These specialized hand tasks could be sustained by the numerous corticomotoneuronal projections from M1 that are present in capuchin and lacking in less dexterous New World monkeys (Bortoff and Strick 1993; Lemon 2008). As suggested from previous studies (Gould and Kaas 1981; Dum and Strick 2005), we found that the hand representations of M1, PMv, and PMd are located on the cortical surface and are thus easily accessible for cortical mapping. We found that the hand representation in M1 was large ( $\sim 40 \text{ mm}^2$ ) and more than 6 times the size of premotor areas, which were all of comparable sizes. This organization is quite different than what has been reported in squirrel monkeys, another New World species, where PMv is much larger than other premotor areas (Dancause, Barbay, Frost, Zoubina, et al. 2006; Dancause et al. 2007; Eisner-Janowicz et al. 2008; Dancause 2013). It is not clear why PMd and SMA (although we have limited data on SMA in the present study) are larger in *Cebus* monkeys in

comparison with other New World primates. Perhaps, the increased repertoire of movements such as the opposition of the thumb rely on corticomotoneuronal connections, while the complexity of motor behaviors such as the use of tools emerged with the development of some premotor areas such as PMd and SMA.

Another appealing aspect of capuchin monkeys is that they have a well-defined area 2 (Padberg et al. 2007). In other New World monkey species, the existence of area 2 is controversial (Kaas 2004). While some studies conclude that the cortex caudal to area 1 has characteristics of area 2 (Merzenich et al. 1978; Liao et al. 2013), others conclude that it does not (Padberg et al. 2005) or leave the question open (Dancause, Barbay, Frost, Plautz, Stowe, et al. 2006; Gharbawie, Stepniewska, Kaas 2011; Negyessy et al. 2013; Ashaber et al. 2014). In the present study, we did find receptive fields that are characteristic of area 2 and a transition that marked the rostral border of area 5. Altogether, the presence of these somatosensory areas with the accessibility of motor representations and behavioral traits of *Cebus* monkeys make them a particularly appealing species to study the cortical network sustaining hand functions.

### Pattern of Projections from Somatosensory Areas to M1

There is a wealth of studies reporting that M1 is interconnected with area 1, area 2, and the posterior parietal cortex (area 5; Jones et al. 1978; Pons and Kaas 1986; Ghosh et al. 1987; Stepniewska et al. 1993). Likewise, we consistently found that



**Figure 8.** Normalized strength of cortical projections to the M1 hand area. (a) To compare the pattern of projections from each distant cortical area to M1, we normalized the proportion of labeled cells. For each injection site (RL: rostro-lateral; CL: caudo-lateral; RM: rostro-medial; CM: caudo-medial), the proportion of labeled cells found in a distant cortical area was divided by that of the area with the most labeled cells. For example, for RM injections, PMd had the highest proportion of labeled cells (normalized value = 1.0). The percentage of labeled cells found in SMA following RM injections was approximately 4 times lower than in PMd (normalized value = 0.26). The cartoon in the middle of the figure shows the color code for the different injection sites (RL = green; CL = orange; RM = red; CM = blue). For each distant cortical area (SMA, PMd, PMv, area 1, area 2, and area 5), a cross diagram shows the normalized strength of its projections to the 4 subregions of the M1 hand area. In each quadrant, the value of the normalized projections is provided and the size of the color square is proportional to the strength of the projections. (b) Cartoon summarizing the pattern of projections to the M1 hand representation from the different areas of the sensorimotor cortex investigated in the present study (SMA, PMd, PMv, area 1, area 2, and area 5). Connections of M1 reported in other studies but not studied here are not shown (e.g., area 3a, S2, etc.). Thick arrows represent “major” projections and thin arrows “moderate” projections, based on the relative strength of connections (a). PMd had the broadest pattern of projections, sending major projections to RM and moderate projections to both RL and CM. In contrast, PMv had a more focal pattern, sending only major projections to RL. SMA had moderate projections to CM. All 3 parietal areas studied were mainly connected with the caudal part of M1. Area 1 had moderate projections to both CM and CL. Area 2 had major projections to CL and moderate projections to CM, and area 5 had major projections to CM and moderate projections to CL.

M1 received projections from these areas in all 3 monkeys. It may be worth noting that although not analyzed in the present study, M1 also has substantial connections with area 3a and fewer connections with area 3b (Krubitzer and Kaas 1990; Stepniewska et al. 1993). Because area 3a and area 3b are buried in the central sulcus of *Cebus* monkeys, the study of connections of these areas would be easier in species with better access to these regions. The

large proportion of ipsilateral inputs to M1 originating from area 1, area 2, and area 5 of the parietal lobe we found (40.7%) contrasts with a previous quantitative study in macaque monkeys that reported much greater cortical connections between premotor areas and M1 (Ghosh et al. 1987). It is difficult to be certain about the cause of these discrepancies, as the 2 studies were conducted in different species. However, we found a greater

proportion of projections from parietal areas to caudal M1, an area buried deep in the central sulcus and more difficult to access in macaque monkeys. One possibility is that we found a much greater proportion of connections from parietal areas because half of our injections specifically targeted caudal M1, resulting in a more complete documentation of M1's inputs.

It is also worth noting that the distribution of labeled cells in area 5 tended to vary along the intraparietal sulcus, depending on the location of the injections in M1. This is particularly obvious in CB-8, where labeled cells from the CL injection tended to be located more rostro-laterally, and from the CM injection tended to be more caudo-medially along the intraparietal sulcus (Fig. 4). The lack of physiological data for area 5 limited our capacity to subdivide it and to conduct separate analyses for each potential region. Nevertheless, there appears to be a similar trend between the qualitative pattern of projections of area 5 in our data with that reported in Old and New World monkeys (Gharbawie, Stepniewska, Kaas 2011; Gharbawie, Stepniewska, Qi, et al. 2011). From these studies as well as in ours, it appears that the rostral portion of area 5, where the "grasp zone" is located, is preferentially connected to CL while more caudal portions of area 5, where the "reach" and "defense zones" are located, are more interconnected with CM.

Whereas quantitative values of connections do not necessarily translate into efficacy, our data might suggest that sensory inputs are as powerful in modulating M1 neurons for the production of motor outputs as premotor areas. Similar to previous studies (Ghosh et al. 1987), we found comparable proportions of inputs from area 1, area 2, and area 5. Balanced proportions of inputs from these areas could support a continuous flow of somatosensory information at various stages of processing (Krubitzer and Kaas 1990). Populations of neurons located in subregions of M1 could use these different sensory signals to affect motor outputs depending on the demands of the task, such as the urgency of the response. Perturbations of arm movements have striking modulatory effects on the activity of M1's neurons (Everts and Tanji 1976; Herter et al. 2009) that can rapidly integrate information about the arm to correct errors (Pruszynski et al. 2011). Possibly, these rapid adaptive processes are supported by the numerous projections from sensory areas to M1. Ecological pressure prioritizing the efficacy of integration of sensory information to the effectors of motor outputs may have resulted in the preferential localization of sensory projections to the caudal portion of M1. This phylogenetically "newer" cortex hosts most pyramidal cells with direct access to motoneurons (Rathelot and Strick 2006, 2009). These pyramidal cells are certainly equipped to integrate complex signals into spinal outputs (Spruston 2008) and have the wiring for the most rapid and effective cortical effects on hand and forearm muscles.

### Pattern of Projections from Premotor Areas to M1

A previous study investigated the cortical network between the digit representations of M1 and premotor areas in Cebus monkeys (Dum and Strick 2005). Normalization of the strength of inputs to M1 showed that the predominant projections came from PMd, followed by PMv and SMA. We found an identical general pattern of inputs to M1. However, the use of multiple smaller tracer injections in subregions of M1 in the present study showed that the distribution of cortical input across the M1 hand area is quite different for each premotor area. PMd sends broader connections across most of the hand area of M1. In contrast, projections from SMA and PMv were more restricted, respectively, to CM and RL. These data suggest that, in comparison with other

premotor areas, PMd may have stronger and more widespread modulatory effects on M1's neurons. Single neuron recording experiments in monkeys have proposed that PMd is involved in the preparation for goal-oriented reaching and online control of movements (Kurata 1993; Lee and van Donkelaar 2006; Pesaran et al. 2006). The specific processing carried out by PMd neurons could be required across a wide range of specialized functions carried out by subregions in M1.

In squirrel monkeys, projections of PMv are found in 3 clusters extending over both hand and proximal representations. The largest and densest cluster is located in the RL portion of M1 arm area, and 2 other smaller clusters in the RM and CL regions (Dancause, Barbay, Frost, Plautz, Popescu, et al. 2006). Results from the present study in Cebus monkeys, where we looked more specifically at the pattern of projections to the hand area, are essentially in accordance with these findings. The greatest proportion of labeled cells in PMv was following RL injections, with fewer labeled cells following injections in others subregions. The finding that PMv sends its most robust projections to RL is thus consistent in different New World monkey species. It is not clear why PMv neurons preferentially project to RL. Studies of cortical interactions have shown that PMv has powerful modulatory effects on M1 outputs (Cerri et al. 2003; Shimazu et al. 2004; Davare et al. 2008), with no indication of weaker or stronger interactions with subregions. This is not so surprising when considering that, in both humans and macaque monkeys, caudal M1 is buried deep within the central sulcus. Thus, studies using transcranial magnetic stimulations (Davare et al. 2008) or microwire arrays on the cortical surface (Cerri et al. 2003; Shimazu et al. 2004) are likely to have largely tested interactions between PMv and rostral M1. Based on our anatomical results, it would be interesting to compare the interactions between PMv, but also PMd, with subregions of M1, something that may be more easily conducted in New World monkeys where the hand representations in these cortical areas are readily available on the cortical surface.

### Support for Subdivisions Within the M1 Hand Area

Many studies have documented subdivisions of the M1 hand area in primates and humans. In particular, it was suggested that the forelimb representation of M1 would be subdivided into a rostral and a caudal aspect, each having different cytoarchitectural, neuronal response properties and patterns of connections with other cortical areas (Jones et al. 1978; Strick and Preston 1982a, 1982b; Humphrey and Reed 1983; Stepniewska et al. 1993, 2006; Geyer et al. 1996; Preuss et al. 1996). For example, corticospinal cells in the rostral portion of M1 mostly make corticospinal connections with upper cervical segments (He et al. 1993). M1 neurons in rostral M1 are typically driven by proprioceptive inputs, whereas neurons with cutaneous receptive fields are more common in caudal M1 (Strick and Preston 1982b). In humans, comparable divisions of M1 into rostral and caudal subregions have also been suggested based on cytoarchitecture and receptor binding and they show different patterns of activation that is task-dependent (Geyer et al. 1996).

In the present study, we made injections of tracers into 4 subregions of the M1 hand representation based on our previous anatomical findings (Dancause, Barbay, Frost, Plautz, Popescu, et al. 2006). This experimental design does not exclude the possibility that more subregions exist within M1 or that these somewhat arbitrary subdivisions are actually functionally relevant. It did, however, allow direct comparison of the pattern of connection of each subregion of the M1 hand representation we chose to study and convincingly showed that they have different patterns

of cortical inputs. These data add onto the already compelling evidence, suggesting the existence of a rostral and caudal subdivision within M1. They also support the concept of a more elaborate parcellation of the hand area of M1 by showing that medio-lateral regions within the rostral and caudal M1 hand representation also have different patterns of cortical inputs.

Cortical mapping experiments using long-train stimulations have shown that cortical sites within given subregions of M1 tend to evoke a specific type of movement (e.g., defense or grasping) that can be associated with the monkey's behavioral repertoire (Graziano, Taylor, Moore 2002). Based on these results, it has been proposed that the M1 forelimb area is organized into multiple subregions, each controlling a category of movements (Graziano, Taylor, Moore, Cooke 2002). Injections of neuronal tracers into different functional zones in M1 identified with long-train stimulations revealed these zones are part of networks that include many of the same cortical areas but with great disparities for the amount of connections. Perhaps, the parietal and premotor areas with same repertoire of movements (Graziano, Taylor, Moore 2002; Gharbawie, Stepniewska, Kaas 2011) would be preferentially interconnected to form networks that carry out subcategories of movements. Overlap between networks would allow coordination of these components to produce integrated movements. Obviously, the functional implication of the fragmentation of M1 into functional subregions will be best settled using neural recording in awake behaving monkeys. Nevertheless, our data contribute to accumulating neuroanatomical evidence, from various species and using different motor mapping techniques (Stepniewska et al. 2009; Gharbawie, Stepniewska, Kaas 2011; Gharbawie, Stepniewska, Qi, et al. 2011), that converge on the same conclusion: Subregions within M1 are parts of largely distinct cortical networks. These recent anatomical data renew long-standing questions regarding internal organization of M1 hand area (Schieber and Hibbard 1993) that are yet awaiting answers.

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