Mapping of multiple muscles with transcranial magnetic stimulation: absolute and relative test–retest reliability

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Abstract
The spatial accuracy of transcranial magnetic stimulation (TMS) may be as small as a few millimeters. Despite such great potential, navigated TMS (nTMS) mapping is still underused for the assessment of motor plasticity, particularly in clinical settings. Here, we investigate the within-limb somatotopy gradient as well as absolute and relative reliability of three hand muscle cortical representations (MCRs) using a comprehensive grid-based sulcus-informed nTMS motor mapping. We enrolled 22 young healthy male volunteers. Two nTMS mapping sessions were separated by 5–10 days. Motor evoked potentials were obtained from abductor pollicis brevis (APB), abductor digiti minimi, and extensor digitorum communis. In addition to individual MRI-based analysis, we studied normalized MNI MCRs. For the reliability assessment, we calculated intraclass correlation and the smallest detectable change. Our results revealed a somatotopy gradient reflected by APB MCR having the most lateral location. Reliability analysis showed that the commonly used metrics of MCRs, such as areas, volumes, centers of gravity (COGs), and hotspots had a high relative and low absolute reliability for all three muscles. For within-limb TMS somatotopy, the most common metrics such as the shifts between MCR COGs and hotspots had poor relative reliability. However, overlaps between different muscle MCRs were highly reliable. We, thus, provide novel evidence that inter-muscle MCR interaction can be reliably traced using MCR overlaps while shifts between the COGs and hotspots of different MCRs are not suitable for this purpose. Our results have implications for the interpretation of nTMS motor mapping results in healthy subjects and patients with neurological conditions.

KEYWORDS
cortical mapping, motor cortex, muscle cortical representation, overlap, reliability, transcranial magnetic stimulation
INTRODUCTION

Transcranial magnetic stimulation (TMS) is one of the main techniques for the non-invasive investigation of the motor-cortex in humans. Measuring the effects of the multiple spot stimulation is a powerful approach for brain mapping, especially when it is combined with MRI navigation (called navigated TMS or nTMS)—a procedure which has been FDA approved for the presurgical brain mapping since 2009 (Krieg, 2017). Recently, nTMS motor mapping has been demonstrated to be even more accurate than functional MRI (fMRI) motor mapping, when compared with the direct cortical stimulation results. Most commonly, during TMS motor mapping one investigates motor evoked potentials (MEPs) using surface (Rossini et al., 2015) or needle EMG (Massé-Alarie, Bergin, Schneider, Schabrun, & Hodges, 2017). When MEPs from the stimulation of many cortical points are acquired, the resulting output is referred to as muscle cortical representation (MCR; Bashir, Perez, Horvath, & Pascual-Leone, 2013; de Carvalho, Miranda, Luis, & Ducla-Soares, 1999), also known as TMS cortical motor map (Kraus & Garabagh, 2015; Novikov, Nazarova, & Nikulin, 2018). There are numerous studies showing that the MCR parameters such as excitability, size and topography reflect functionally relevant features of the motor cortex organization in healthy people (Beaulieu, Massé-Alarie, Ribot-Ciscar, & Schneider, 2017; Gentner & Classen, 2006; Nazarova, Novikov, Nikulin, & Ivanova, 2020; Tyč & Boyadjian, 2011) and in patients with motor pathology such as stroke (Lüdemann-Podubecká & Nowak, 2016; Yarossi et al., 2019), amyotrophic lateral sclerosis (Chervyakov et al., 2015; de Carvalho et al., 1999), dystonia (Schabrun, Stinear, Byblow, & Ridding, 2009), and so forth. Considering the TMS mapping of multiple muscles, the existence of the somatotopy gradient for MCRs, along with their extensive overlap, has been discussed from the earliest TMS studies (Gentner & Classen, 2006; Metman, Bellevich, Jones, Barber, & Streletz, 1993). The spatial accuracy of TMS, depending on stimulation parameters, based on animal data may be as small as 2 mm (Romero, Davare, Amendariz, & Janssen, 2019), thus allowing the tracing of within-limb MCR interactions, especially when taking into account individual sulcus anatomy (Raffin, Pellegrino, Di Lazzaro, Thielser, & Siebner, 2015). These multiple-muscle MCR interactions are believed to reflect the basic neuroanatomical prerequisites for the modular organization of movements at the cortical level (Dubbioso, Raffin, Karabanov, Thielser, & Siebner, 2017; Gentner & Classen, 2006).

However, despite the great potential of TMS mapping, it is still underused for the assessment of motor plasticity, particularly in clinical settings. In order to use this approach in clinic, its reliability should be first established, both relative and absolute. Relative reliability reflects the degree of variation in position among participants over repeated measurements and, thus, allowing subjects/patients stratifying, while absolute reliability reflects the degree of variation in measurements for individuals, and, thus, allowing the interpretation of changes at the individual level (Atkinson & Nevill, 1998). Articles dedicated to TMS motor mapping reliability are mostly focused on the relative reliability (Cavaleri, Schabrun, & Chipchase, 2018; Forster, Limbart, Seifert, & Senft, 2014; Jonker et al., 2019; Kraus & Garabagh, 2016; Malcolm et al., 2006; McGregor et al., 2012; Ngomo, Leonard, Moffet, & Mercier, 2012; Pitkänen et al., 2017; Plowman-Prine, Triggs, Malcolm, & Rosenbek, 2008; Sankarasubramanian et al., 2015; Sinitsyn et al., 2019; Sollmann et al., 2013; van de Ruit, Perenboom, & Grey, 2015; Weiss et al., 2013; Wolfa et al., 2004; Zdunczyk, Fleischmann, Schulz, Vajkoczy, & Picht, 2013), while absolute reliability has been investigated much less frequently (Jonker et al., 2019; Ngomo et al., 2012; Sankarasubramanian et al., 2015; van de Ruit et al., 2015). As for the multiple-muscle TMS mapping, to the best of our knowledge, there are fewer reports in the literature (Cavaleri et al., 2018; Forster et al., 2014; Plowman-Prine et al., 2008; Weiss et al., 2013), and the reliability of inter-muscle interactions have not yet been addressed. Furthermore, another limitation, considering TMS MCR reliability, is the lack of an agreement on which MCR parameters should actually be studied. Apart from the MCR parameters commonly used and recommended in the TMS clinical guidelines, such as areas, volumes, centers of gravity (COGs) and hotspots (Rossini et al., 2015), there is a range of more sophisticated parameters, such as MCR shape (Pitkänen et al., 2017), discrete peaks per MCR (Beaulieu, Flamand, Massé-Alarie, & Schneider, 2017; Cavaleri, Schabrun, & Chipchase, 2017; Schabrun, Hodges, Vicenzino, Jones, & Chipchase, 2015), MCR excitability profile (EP) (Novikov et al., 2018; Raffin et al., 2015) etc., but the validity of these parameters has not yet been fully established.

In this study we conducted a systematic investigation of both the absolute and relative reliability of three hand muscle MCRs and their interactions in a homogenous group of healthy male volunteers using a comprehensive (≥170 points) individual brain nTMS mapping. Additionally, we studied the somatotopy gradient among the investigated hand muscle MCRs. Moreover, to complement standard MCR metrics, we used overlaps between MCRs, and differences of MCR EPs with the idea that it would be possible to identify muscle-specific MCR features suitable for probing in longitudinal multiple-muscle TMS mapping studies.

MATERIALS AND METHODS

Data supporting the findings of this study are available from the corresponding author upon reasonable request.

2.1 | Participants

Twenty-two young (19–33 years old) healthy male volunteers were enrolled in the study. Exclusion criteria were: any history of neurological/psychiatric disorders including fainting spells, any medications intake, cardiac implants, metallic implants in the head, implanted pumps, stimulators and shunts or MRI incompatibility for any reasons. We also strictly excluded subjects with special motor skills, for example, professional or trainee sportsmen, musicians, surgeons, painters and people with special hobbies requiring high manual dexterity. Participants self-reported for the usual 6–9 hr of sleep before the TMS
procedure, no alcohol intake 24 hr before, and the usual amount of coffee intake. Subjects self-reported their handedness. All subjects gave written informed consent in accordance with the Declaration of Helsinki. All subjects were screened for contraindications to TMS (Rossi, Hallett, Rossini, & Pascual-Leone, 2009) before the consent process. Experiments were approved by the IRB of the Research Center of Neurology N312 and the local Ethics Committee of HSE University, Moscow.

2.2 Overall experimental approach of test–retest nTMS mapping

Two nTMS mapping sessions were performed on each subject (Day 1 and Day 2) separated by 5–10 days. Both TMS mapping sessions were conducted for each volunteer at the same day time (morning/afternoon/evening). During the period between Days 1 and 2 volunteers were asked to keep their lifestyle unchanged, in particular, not to change their usual motor activity (e.g., their usual fitness schedule), not to start learning new motor skills etc. Because of the violations of some of these requirements, we excluded four volunteers; therefore the data of 18 volunteers were used for further analysis.

2.3 Electromyography

MEPs were obtained from three upper limb muscles: two intrinsic hand muscles – abductor pollicis brevis (APB) and abductor digitii minimi (ADM) and one extrinsic hand muscle – extensor digitorum communis (EDC). Surface EMG was recorded by the integrated EMG device of the eXimia system (3 kHz sampling rate, band-pass filter of 10–500 Hz). Bipolar surface 0.6-cm² ECG Ag-AgCl electrodes (3 M Red Dot) were placed using belly-tendon montage: the active electrode was located over the muscle belly, the referent 1 and 2 cm distally, and the ground one—on the wrist of the same hand. APB and EDC muscles were chosen because they are usually associated with hand motor deficits, for instance, in stroke patients (thumb abduction and finger extension; Nazarova et al., 2021), also APB is one of the most commonly investigated muscles in TMS studies (Rossini et al., 2015). ADM was added to trace the within-hand cortical somatotopy. The MEP peak-to-peak amplitudes were calculated online using the eXimia software. During the initial preprocessing, EMG data were visually inspected; trials were rejected if noise or the preactivation of any muscle was higher than 20 μV (peak-to-peak EMG amplitude).

2.4 nTMS cortical mapping

All TMS mapping procedures were carried out in accordance with the TMS safety guidelines (Rossi et al., 2009). Prior to the mapping, individual anatomical T1-weighted magnetic resonance images were acquired by a 1.5 T MR-scanner Siemens Magnetom Avanto (structural T1-weighted images, MPRAGE, 1 mm isotropic voxel, acquisition matrix 256x256). Single pulse nTMS investigation was performed using a co-planar figure-of-eight induction coil (outer winding diameter 70 mm) for biphasic stimulation, connected to an eXimia magnetic stimulator (Nexstim Ltd., Helsinki, Finland, version 3.2.2), eXimia NBS navigation system (Nexstim Ltd.) was used for MRI-guided navigation.

As a first step a “rough TMS mapping” (Krieg, 2017) procedure, where the intensity was chosen to induce MEP with the amplitudes range around 500–800 μV, starting from the “hand knob” region in the precentral gyrus (Youssry et al., 1999) was performed to find a “technical hotspot” for the APB muscle for further resting motor threshold (RMT) probing. The “technical hotspot” was defined as the coil position (considering the electric field [EF] direction) resulting in the highest MEP amplitudes from the APB muscle. Then RMT for the given “technical hotspot” was determined as a minimal stimulator output (MSO), producing contralateral APB MEPs with minimal peak-to-peak amplitude of 50 μV in a resting muscle, in 5 out of 10 stimuli using a Rossini-Rothwell method of RMT hunting (Rossini et al., 2015). We have always started with the intensity used for the “rough TMS mapping”. For the Day 2 nTMS mapping session, we looked for the APB hotspot and RMT again.

The intensity of the stimulation during nTMS mapping was 110% of the RMT for APB. The time lag between the stimuli varied randomly between 3 and 10 s. Each of the two mapping sessions included five sub-sessions, separated by 2–10 min. Each sub-session consisted of 49–79 TMS pulses (a constant number of points for each volunteer). The stimulation nodes were pre-set using a virtual MRI-based grid in the navigation software with 5 × 5 mm² squares. We kept the coil perpendicular to the closest segment of the central sulcus (Bashir et al., 2013; Krieg et al., 2017; Raffin et al., 2015).

The outer margin of a given MCR was determined if no MEP could be elicited in two consecutive stimulation trials. During nTMS mapping sub-sessions, the same cortical targets were stimulated in the forward and reverse order five times (sub-session 1 “forward”—sub-session 2 “reverse”—sub-session 3 “forward”—sub-session 4 “reverse”—sub-session 5 “forward”) to avoid the effect of the repeated stimulation of the same target (Figure 1). On Day 2, the points were stimulated in exactly the same order as on Day 1. The precision of nTMS, linked to the coil localization error of the navigation system (Ruohonen & Karhu, 2010) was kept below 2 mm for each cortical spot and the coil tilt was constant, according to the navigation system feedback. In the end, the whole analyzed stimulation session consisted of 177–395 stimuli per day depending on the volunteer (see an example of APB MCR in Figure 1).

2.4.1 Along central sulcus nTMS mapping

Considering that the coil in our study was kept perpendicular to the central sulcus, we also aimed at comparing our results with those, obtained using the so-called “along-sulcus” TMS mapping (Raffin et al., 2015; Raffin & Siebner, 2019), although it was not our original
intention. As using this “along-sulcus” TMS mapping approach, a clear within-limb somatotopy gradient between the first dorsal interosseus and ADM MCRs, was recently reported (Raffin et al., 2015; Raffin & Siebner, 2019). We manually identified stimulation points located in the “hand knob” along the central sulcus for Day 1. We refer to these data as “along-sulcus” mapping, while for mapping data based on all points as “whole” MCRs.

2.4.2 | Single-muscle MCR size and topography parameter calculation

MCRs were constructed and analyzed using TMSmap—a freely available software for the quantitative assessment of TMS cortical mapping data (Novikov et al., 2018). As indicated in (Novikov et al., 2018), the locations of the EF maxima, provided by the navigation software, were projected on the sphere tangential to the cortex surface. We calculated commonly used size MCR parameters such as areas, volumes ($\sum_{N=1}^{\text{MEP}_1 > 50 \mu V \text{ peak-to-peak amplitude}} \Delta S_i$, where $N$—the number of squares with area $\Delta S_i$, where $\text{MEP}_1$ peak-to-peak amplitude $\geq 50\mu V$), mean MEP per MCR, and topography parameters such as hotspot and COG locations (the COG formula was the same as elsewhere, including the TMSmap software description [for details please see Novikov et al., 2018]). We merged stimulation points located closer than 2 mm (see Figure 1) (Novikov et al., 2018) and used mean MEP amplitudes after merging for MCR construction. The threshold MEP amplitude for MCR construction was 50 $\mu V$ peak-to-peak for a merged point. For the purpose of MCR comparison we redefined the hotspot: instead of using the “technical hotspot” where RMT was defined, we considered the whole TMS mapping data per day. Thus, we again defined the “final hotspot” as the location over which averaged MEPs with the highest peak-to-peak amplitude were evoked in each of the target muscles, so like that we determined hotspots for all three muscles. Additionally, we compared the EPs of different MCRs using earth mover’s distance (EMD) metrics (see the next subsection).

2.4.3 | The topography parameters of multiple muscle MCR interactions

As a reflection of multiple muscle MCR interactions we used:

1. Geodesic shifts between different muscles MCR COGs’ and hotspots’ $x$ and $y$ coordinates (reflecting shifts in the medio-lateral and anterior–posterior directions, respectively);
2. Area and volume overlap between two MCRs. For MCR overlap calculation we used an approach similar to the Jaccard index, implemented in TMSmap (Novikov et al., 2018). Briefly, area and

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**FIGURE 1**  Navigator TMS (nTMS) of the left primary motor cortex for a representative subject (visualization in the TMSmap software). Stimulation points are shown for both days (upper raw—Day 1, bottom raw—Day 2). In the “Sub-sessions 1–5” column the order of the stimulation is shown. Magenta filled arrow indicates the first stimulation point, magenta non-filled arrow—the last stimulation point in an nTMS mapping sub-session. White dotted line indicates the central sulcus. One can notice that the location of the stimulation points for Day 1 and Day 2 are almost identical. “Stimulation points” column represents all stimulation points per Days 1 and 2, while in the “Merged stimulation points” column, points after spatial filtering are shown. Stimulation was done using 35% of the maximal stimulator output (110% of the resting motor threshold). $X$-axis corresponds to the lateral to medial direction; $Y$-axis corresponds to the posterior to anterior direction. The color scale represents motor evoked potential peak-to-peak amplitude in microvolts ($\mu V$).
volume normalized overlaps between MCRs were calculated using the following formulas:

\[ a_{12} \% = \frac{a_{12}}{a_1 + a_2 - a_{12}} \times 100 \%, \quad v_{12} \% = \frac{v_{12}}{v_1 + v_2 - v_{12}} \times 100 \% , \]

where \( a_1, a_2 \) are the areas of MCR 1 and 2, respectively; \( v_1, v_2 \) are the volumes of MCR 1 and 2, respectively; \( a_{12}, v_{12} \) are the common area and volume for the MCR 1 and 2, respectively. So, the normalized overlaps between MCRs can be interpreted in such a way that the value of 100% indicates completely identical MCRs, while the value of 0% indicates no overlap between MCRs. Further in the text we will use only normalized overlaps and refer to them as overlaps.

3. We conceptualize EP as a way to refer to complex 3D-profile of MCR. EP for the along sulcus TMS mapping has been previously reported to differ between distal upper limb muscles MCRs (Raffin et al., 2015). We have previously proposed EP to serve as a parameter reflecting the convergence phenomenon of the motor cortex organization for the whole cortex nTMS mapping (Novikov et al., 2018). EPs of two MCRs of different muscles on 1 day were compared using the earth mover’s distance (EMD), also called the Wasserstein metric (Rubner, Tomasi, & Guibas, 1998). EMD characterizes a minimum cost of turning one distribution into the other (Rubner et al., 1998; see a detailed description of EMD in Novikov et al., 2018). We have earlier implemented EMD calculation for quantitative comparison of EPs in the TMSmap software (Novikov et al., 2018), but it has not yet been validated in a test–retest study. The same approach of EMD calculation was used for the reliability assessment of MCR EPs between days.

2.6 | Statistical analysis

Apart from the analysis in TMSmap, further statistical analysis was performed using the SPSS package for Windows (IBM) and Matlab (Natick). The level of significance was defined as \( \alpha = .05 \). Family-wise error rate (FWER) correction was performed to control for multiple comparisons.

2.6.1 | MCR within-limb somatotopy gradient assessment

We used one-way repeated measures ANOVA (rmANOVA) to confirm the within-limb gradient among hand muscle MCRs (for this we again used 17 right-handed subjects, excluding one left-handed participant). We compared COGs of the different muscle MCRs to probe the mediolateral shift among them for (a) the individual “whole” MCRs, (b) the “along-sulcus” mapping data, and (c) the MNI version of MCRs. Additionally, using two-way rmANOVA we compared geodesic shifts between different muscle MCR COGs inside 1 day for the “whole” and the “along-sulcus” mapping approaches (factors: muscle, TMS mapping approach). Finally, we used one-way rmANOVA to compare the areas of different muscle MNI MCRs for each day. Shapiro–Wilk test for normality (Ghasemi & Zahediasl, 2012) and Mauchly’s test for sphericity assessment (Beddo & Kreuter, 2004) were applied.

2.6.2 | Reliability assessment

Before the reliability analyses, we also assessed normality by Shapiro–Wilk test (Ghasemi & Zahediasl, 2012) and homoscedasticity, using Levene’s test (Beddo & Kreuter, 2004; Weir, 2005). When non-normality or heteroscedasticity were found for more than one muscle, a natural logarithmic transformation was applied before reliability analysis for this parameter for all three muscles (Atkinson & Nevill, 1998; Beaulieu, Flamand, et al., 2017). In case when after logarithmic transformation the normality still was not achieved we decided not to do bootstrapping, considering that in case of low amount of data it is difficult to estimate the distribution’s tails (Orloff & Bloom, 2014), and in this case, we did not estimate smallest detectable change (SDC). The general pipeline of MCR reliability assessment is presented in Figure 2.

2.6.3 | Relative reliability assessment (ICC)

We used a two-way mixed model absolute agreement intraclass correlation coefficient (ICC) to assess relative reliability for single-muscle MCR parameters (Beaulieu, Massé-Alarie, et al., 2017; Koo & Li, 2016). We calculated ICC between days for RMTs, map areas, volumes, mean MEP, hotspots’ and COGs’ x and y coordinates for each MCR. To check the reliability of multiple muscle MCR relationship we calculated (a) the shifts between different muscles MCRs’ COGs’ and
hotspots’ coordinates, (b) the extents of the different muscle MCR overlaps, and (c) EMDs between the different muscle MCR EPs. For relative reliability description, we used the following ranges: below 0.50—poor; between 0.50 and 0.75—moderate; between 0.75 and 0.90—good; above 0.90—excellent (Koo & Li, 2016).

2.6.4 | Muscle-specificity assessment

In addition to relative reliability, we investigated whether MCR parameters were muscle-specific. We defined “muscle-specificity” of the TMS MCR parameter, as a possibility to predict a given muscle among other investigated muscles studying mapping parameters across subjects. For this, in addition to ICC, we assessed the Spearman rank correlation for the same MCR parameters. We considered a parameter to be muscle-specific when its values for a given muscle between days, agreed (ICC) or correlated (Spearman correlation coefficient) significantly more for the same-muscle MCR, compared to when this parameter was evaluated for the MCRs of other muscles (either on the same day or between days). In other words, muscle-specificity provides the possibility of identifying the correct muscle, among other muscles.

2.6.5 | MCR EP comparison using EMD

To investigate the validity of different MCR EPs we compared:

1. Normalized EMD for the same muscle MCRs between days, calculated according to the formula (an example for APB MCR):

$$nEMD_{d1APB, vs. d2APB} = \frac{EMD_{d1APB, vs. d2APB}}{EMD_{mean}}$$

where

$$EMD_{mean} = \frac{(EMD_{d1APB, vs. d2APB} + EMD_{d1APB, vs. d2ADM} + EMD_{d1APB, vs. d2EDC})}{3}$$

2. And normalized EMD across-muscle MCRs between days, calculated according to the formula (an example for APB MCR versus two other muscles’ MCRs):

$$nEMD_{d1APB, vs. d2ADM} = \frac{EMD_{d1APB, vs. d2ADM}}{EMD_{mean}}$$

$$nEMD_{d1APB, vs. d2EDC} = \frac{EMD_{d1APB, vs. d2EDC}}{EMD_{mean}}$$

To compare normalized EMDs between same-muscle MCR EPs across days versus different muscle MCR EPs across days we used the non-parametric Mann–Whitney U test as the distribution of the residuals in one of the samples was not normal according to Shapiro–Wilk test.

2.6.6 | Absolute reliability (SDC) assessment

For the assessment of the absolute reliability of the MCR parameters we used the SDC (a.k.a. minimal detectable change) calculation. SDC represents the minimal change that a subject must show on the scale to ensure that the observed change is real and not just a measurement error (Beaulieu, Flamand, et al., 2017; Weir, 2005). We estimated SDC using the standard error of measurements (SEMeas):

$$SDC = C_C \cdot \sqrt{2 \cdot SEMeas},$$

where $C_C$ is a coefficient which is equal to 1.00, 1.65 or 1.96 associated with a 68, 90, and 95% confidence intervals (CI), respectively. The 68% CI was included because we assume that it may be adequate for change interpretation at the individual level. $SEMeas = \sqrt{MSE}$, where MSE is the mean squared error obtained from the rmANOVA applied on test and retest measurements (Weir, 2005). When SDC values were obtained for log-transformed parameters, we then performed the antilog transformation as in (Beaulieu, Massé-Alarie, et al., 2017).
3 | RESULTS

Figure 3 provides an example of a representative subject’s test-retest single-muscle MCRs and their overlaps for both days. The figure shows the distribution of MEP amplitudes depending on the stimulation location relative to the mapping grid. One can clearly see that all MCRs are different but it is also hard to categorize this difference visually. In this example, the APB MCR is shifted laterally compared to two other MCRs (COG x is smaller), it also has the biggest size; and the overlap between APB and ADM MCRs is the most prominent. We provide the data for all the participants’ single-muscle MCR parameters from TMSmap software (areas, volumes, mean MEPs, hotspot, and COG location) and the inter-muscle MCR interaction parameters (areas and volumes overlaps and EMDs between the different muscle MCRs) (Tables S7–S9 in the Supporting Information).

The results are described in the following manner. First, we report the within-limb somatotopy gradient for the individual MRI MCRs and then for MNI data. Second, we show relative and absolute reliability and the muscle-specificity of different MCR parameters, assessed for all single-muscle MCRs. Finally, we present the inter-muscle MCR interaction reliability in the same order as for a single-muscle MCRs (see the general pipeline of the reliability assessment in Figure 2 in Section 2).

3.1 | Within-limb somatotopy gradient

3.1.1 | Somatotopy gradient for individual MRI data

Despite the great overlap between the muscle MCRs, we observed a mediolateral shift between the APB MCR and the two other muscle MCRs, based on COGs’ x coordinates. This shift was significant for Day 1 (1 way rmANOVA: Day 1: \( p < 10^{-6} \) (pair-wise comparison after FWER APB-ADM: \( p < 10^{-5} \), ADM-EDC: \( p = 0.227 \), APB-EDC: \( p < 10^{-3} \)). For the second day, this shift was observed only as a tendency (\( p = 0.021 \) [pair-wise comparison after FWER APB-ADM: \( p = 0.064 \), ADM-EDC: \( p = 0.539 \), APB-EDC: \( p = 0.071 \)].

3.1.2 | Along-sulcus TMS mapping

We compared our results with those obtained using the ”along-sulcus” mapping (Raffin et al., 2015; Raffin & Siebner, 2019) because previously for this approach within-limb somatotopy gradient between the first dorsal interosseus and ADM muscle MCRs was reported (Raffin et al., 2015; Raffin & Siebner, 2019). We were able to identify 28 to 63 points per participant in the precentral gyrus along the central sulcus (see an example in Figure 4). In two participants, all...
three muscle MCRs were shifted rostrally beyond the central gyrus, and, thus, there were no MEPs ≥50 μV peak-to-peak in the points along the central sulcus (see an example in Figure 5 in the Supporting Information). When comparing the mediolateral shift of the “along-sulcus” MCRs, based on Day 1 COG x-coordinates in 15 participants with no null along-sulcus MCRs, we observed a significant difference between the APB MCRs versus the ADM and EDC MCRs COG x-coordinates (one-way rmANOVA \( p < 10^{-4} \), pair-wise comparison after FWER—APB-ADM: \( p = .001 \), ADM-EDC: \( p = 1 \), APB-EDC: \( p < 10^{-3} \)).

To check whether the mediolateral somatotopy gradient was more or less pronounced in the “along-sulcus” in contrast to the “whole” MCRs, we compared the differences between the COG x-coordinates between the different muscle MCRs for these two mapping approaches. We found that the mediolateral somatotopy gradient was greater for the “whole” MCRs compared to the “along-sulcus” MCRs (rmANOVA with two factors (“muscle-combination” and “TMS mapping type”) showed a significant factors interaction (\( p = .005 \)). While looking at the example in Figure 4, one can see that the size of different muscle MCRs for “whole” and the “along-sulcus” mapping does not correspond. While the “whole” APB MCR is bigger than the “whole” ADM MCR, for the “along-sulcus” MCRs this interrelation is opposite. Considering (a) this mismatch, (b) the fact that it was impossible to obtain ≥70 stimulation points along the central sulcus (in analogy with 10 repetitions of 7 targets as in [Raffin & Siebner, 2019]) in any participant, and (c) that in two of our participants we obtained no MEPs along the central sulcus at all (see example in Figure 5), we have not further analyzed the “along-sulcus” MCRs quantitatively.

### 3.1.3 MNI data somatotopy gradient

In addition to individual data assessment, we also probed the somatotopy gradient of MCRs across subjects, using data co-registered to the MNI space. For MNI assessment we used 17 right-handed subjects. We observed that APB MCR were more lateral compared to the ADM and EDC MCRs for both days based on COG x-coordinates (1-way rmANOVA, pair-wise comparison after FWER, APB-ADM: Day 1: \( p = .001 \), Day 2: \( p = .016 \), APB-EDC day1: \( p < 10^{-3} \), Day 2: \( p = .004 \)). We also found the difference between ADM and EDC MCR areas: ADM MCR area was significantly smaller for both days (1-way rmANOVA, pair-wise comparison after FWER, day1: \( p = .005 \), Day 2: \( p = .041 \)).

In addition, we created weighted normalized MCRs for both days (Figure 6). Normalized MEPs amplitudes (averaged across subjects) varied from 0 to 48%, meaning that not more than half of the subjects had their maximum MEPs in the same spot. Consistently with the previously observed somatotopy gradient, we again found a lateral shift of the COG of the APB MCR compared to ADM and EDC MCRs.
**FIGURE 5** An example of a TMS mapping dataset where there are no motor evoked potentials (MEPs) ≥ 50 μV along the central sulcus (white dotted line). Stimulation was done using 39% of the maximal stimulator output (110% of the resting motor threshold). In the upper row along-sulcus mapping data for all three muscles are shown, no MEP with amplitude ≥50 μV can be seen. The lower row represents the “whole” muscle cortical representations (MCRs) for all three muscles. Centers of gravity (COGs) are shown with white crosses, hotspots—with white triangles. The color scale represents MEP amplitude in microvolts. Magenta squares indicate the same cortical area. X-axis corresponds to the lateral to medial direction; Y-axis corresponds to the posterior to anterior direction.

**FIGURE 6** Weighted MNI muscle cortical representations (MCRs) of APB, ADM and EDC, Day 1. Centers of gravity (COGs) are shown with white crosses. Weighted MCR areas (in cm²) and maximum of the overlapped motor evoked potentials (MEPs) (in %) are shown. Magenta squares indicate the same cortical area. X-axis corresponds to the lateral to medial direction; Y-axis corresponds to the posterior to anterior direction. The color scale reflects the mean weighted MEP at a stimulation point.
can also be seen that EDC normalized MCR has a bigger red zone, indicating that the optimal areas with high MEPs for EDC MCRs across subjects are located closer to each other compared to the two other muscle MCRs.

3.2 Single-muscle cortical TMS map reliability analysis

3.2.1 Relative reliability

TMS map size parameters
ICC for areas and log-transformed volumes and mean MEPs varied from good to excellent (0.71–0.85), depending on a muscle (Table 1).

TMS map standard topography parameters (hotspots, COGs)
ICC for hotspots and COGs was almost always excellent for individual MRI data (> = 0.9) (Table 2). In the MNI space ICC decreased: for COGs it varied from good to excellent, while for hotspots—from fair to good (Table 2). We suggest that MNI ICC for hotspots and COGs is a more correct because it is not sensitive to high data variance among the individual brains.

3.2.2 Absolute reliability

MCR size parameters
Individual SDC (CI 95%) for areas varied from 2.01 to 2.58 cm². When the CI is decreased to 68%, individual SDC for areas was half the size: 1.03–1.32 cm², depending on the muscle. Evidently, a similar dependency of the SDCs on CI has been also observed for all other parameters (Table 1).

MCR standard topography parameters (COGs and hotspots)
Individual SDC (CI 95%) for the hotspot coordinates was around 1 cm, while for COG coordinates – around 0.5 cm. SDC values for different muscle MCRs were very similar (Table 2).

3.2.3 Muscle-specificity

MCR size parameters
Based on ICC, none of the MCR size parameters was muscle-specific (Figure 7). When using Spearman’s correlation, however, we observed muscle specificity after FWER correction—this can be seen as stronger correlation values on the main diagonal of the matrix compared to off-diagonal values, reflecting a correlation between different muscles’ MCR parameters values (Figure 7). The situation for the MNI data was similar, mean MEP muscle-specificity was significant (Figure 8).

| TABLE 1 | Relative (ICC) and absolute (SDC) reliability values for single-muscle (APB, ADM and EDC) cortical representation (MCR) size parameters in the individual space. The data are shown for different confidence intervals (CIs) |
|------------------|------------------|------------------|------------------|------------------|------------------|
| Parameter, muscle | ICC | SE meas | Individual | 68% CI | 90% CI | 95% CI | Group (n = 18) |
|------------------|------------------|------------------|------------------|------------------|------------------|------------------|
| RMT % | APB | 0.99 | 0.48 | 0.7 | 1.1 | 1.3 | 0.3 |
| Area, cm² | APB | 0.85 | 0.72 | 1.03 | 1.69 | 2.01 | 0.47 |
| | ADM | 0.84 | 0.76 | 1.08 | 1.78 | 2.11 | 0.50 |
| | EDC | 0.70 | 0.93 | 1.32 | 2.17 | 2.58 | 0.61 |
| Ln (mean MEP) | APB | 0.72 | 0.296 (x/±1.34) | 0.419 (x/±1.52) | 0.691 (x/±2.00) | 0.821 (x/±2.27) | 0.193 (x/±1.21) |
| | ADM | 0.79 | - | - | - | - | - |
| | EDC | 0.74 | 0.293 (x/±1.34) | 0.414 (x/±1.51) | 0.684 (x/±1.98) | 0.812 (x/±2.25) | 0.191 (x/±1.21) |
| Ln (volume) | APB | 0.79 | 0.497 (x/±1.64) | 0.703 (x/±2.02) | 1.160 (x/±3.19) | 1.377 (x/±3.96) | 0.325 (x/±1.38) |
| | ADM | 0.85 | - | - | - | - | - |
| | EDC | 0.72 | 0.425 (x/±1.53) | 0.602 (x/±1.82) | 0.993 (x/±2.70) | 1.179 (x/±3.25) | 0.278 (x/±1.32) |

Note: x/±—anti-log value of SE meas or SDC results.
Abbreviations: ICC, intraclass correlation coefficient (a value between 0 and 1); Ln (mean MEP), log-transformed mean motor evoked potentials; Ln (volume), log-transformed MCR volumes; RMT, resting motor threshold; SDC, smallest detectable change; SE meas, standard error of measurement.
Considering that the MNI variant of ICC for COGs and hotspots might be more suitable when using data from many subjects (due to a smaller data variance and because of the co-registration to a common brain template), we demonstrate here also the results in the MNI space (Figure 8). No muscle-specificity is observed for COGs or hotspots, either in the MNI or in the individual data.

**TABLE 2** Relative (ICC) and absolute (SDC) reliability values for hotspots and centers of gravity (COGs) for single-muscle (APB, ADM, and EDC) cortical representations (MCRs) in the individual space for different confidence intervals (CIs)

<table>
<thead>
<tr>
<th>Parameter, muscle</th>
<th>ICC</th>
<th>SEMeas</th>
<th>SDC Individual</th>
<th>68% CI</th>
<th>90% CI</th>
<th>95% CI</th>
<th>95% CI</th>
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<tbody>
<tr>
<td>COG x, mm</td>
<td></td>
<td></td>
<td></td>
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<td></td>
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<td></td>
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<tr>
<td>APB</td>
<td>0.99</td>
<td>1.79</td>
<td>2.53</td>
<td>4.17</td>
<td>4.96</td>
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<tr>
<td>ADM</td>
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<tr>
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<td>2.17</td>
<td>3.58</td>
<td>4.26</td>
<td>1.00</td>
<td></td>
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<tr>
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<td></td>
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<td></td>
</tr>
<tr>
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<td>3.97</td>
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<td>7.79</td>
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</table>

Note: x-axis corresponds to the lateral to medial direction, y-axis corresponds to the posterior to anterior direction.

Abbreviations: ICC, intraclass correlation coefficient (a value between 0 and 1); SDC, smallest detectable change; SEMeas, standard error of measurement.

**FIGURE 7** Correlation matrices for the size parameters of a single-muscle cortical representations (MCRs) in the individual space (1—APB, 2—ADM, 3—EDC). Diagonal elements in the matrix indicate within-muscle comparison, while off-diagonal elements indicate between-muscle comparisons. FWER, family-wise error rate; MEP, motor evoked potential.
3.3 | Multiple-muscle MCRs interaction (within-limb TMS somatotopy) reliability

3.3.1 | Relative reliability

Reliability of the shifts between either COGs, or hotspots of the different muscle MCRs was poor (Table 3). At the same time, relative reliability of the areas and volumes overlaps of MCRs of different muscles was good to excellent (ICC from 0.8 to 0.9), except for the APB-ADM area and volume overlaps (ICC of which still corresponds to moderate reliability) (Table 3). ICC of EMDs between the different muscle MCR EPs was smaller (from 0.4 to 0.56), but still higher than that for COGs and hotspots shifts (Table 3).

3.3.2 | Absolute reliability

SDC for area and volume overlaps is in the range 0.17–0.34 (CI 95%), while SDC for the inter-muscle EMDs was around 7.5% (CI 95%; Table 3).

3.3.3 | Muscle-specificity

Based on ICC before FWER, muscle-specificity can be seen for EMDs between different muscle MCR EPs (Figure 9). Based on Spearman’s correlation, areas, and volume overlaps, and inter-muscle EMDs are muscle specific and after FWER such specificity remains for the MCR areas overlaps (Figure 9).

![Correlation matrices for topography parameters of single-muscle cortical representations (MCRs) in the MNI space (1—APB, 2—ADM, 3—EDC). The color bar indicates the strength of the correlation. Diagonal elements in the matrix indicate within-muscle MCR comparison, while off-diagonal elements indicate between-muscle MCR comparison. COG, center of gravity; FWER, family-wise error rate; MEP, motor evoked potential.](image-url)
DISCUSSION

In this study we investigated the within-limb somatotopy gradient and the reliability of MCRs of three hand muscles in a homogenous group of healthy male volunteers, using a comprehensive grid-based sulcus-informed nTMS motor mapping. The main findings of the study are listed below.

We observed a somatotopy gradient for the hand muscle MCRs reflected by APB MCR COGs being more lateral compared to other muscles’ MCR COGs (this result was significant for the first day). We demonstrated that the most commonly used metrics of MCRs, such as areas, volumes, COGs, and hotspots, while having overall high relative reliability, are generally characterized by low absolute reliability.

For within-limb TMS somatotopy, the most common metrics such as the shifts between different muscles’ MCR COGs and hotspots have poor relative reliability and are not muscle-specific. While overlaps and, to a lesser extent, EMDs between different muscle MCR EPs, are more reliable and tend to demonstrate muscle-specificity.

Further, we discuss each of these points in detail.

<table>
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<tr>
<th>Parameter, muscle</th>
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<td></td>
<td></td>
<td></td>
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<td>90% CI</td>
<td>95% CI</td>
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<td>19.0</td>
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<td>8.9</td>
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<td>3.89</td>
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<td>7.52</td>
</tr>
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<td></td>
</tr>
<tr>
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<td>1.95</td>
<td>3.22</td>
<td>3.83</td>
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<tr>
<td>APB-ADM</td>
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<td>APB-ADM</td>
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<td>13.31</td>
</tr>
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<td>5.82</td>
<td>8.24</td>
<td>13.59</td>
<td>16.15</td>
</tr>
<tr>
<td>APB-EDC</td>
<td>−0.30</td>
<td>6.13</td>
<td>8.66</td>
<td>14.29</td>
<td>16.98</td>
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ICC – intraclass correlation coefficient (a value between 0 and 1).
SDC – smallest detectable change.
SEMeas – standard error of measurement.
SEMeas and SDC values for the cases when ICC values are low should be considered with caution.
CI – confidence interval.
X-axis corresponds to the lateral to medial direction, y-axis corresponds to the posterior to anterior direction.
4.1 Somatotopy gradient

The existence of a robust within-limb somatotopy in the primary motor cortex (M1), obtained with TMS, is still debated, although “homunculus-like” TMS motor cortical maps were described from the earliest years of TMS mapping (Gentner & Classen, 2006; Metman et al., 1993). A commonly reported output for different muscles' TMS MCRs interaction is the distances/shifts between the COGs (Dubbioso et al., 2017; Schabrun et al., 2009; Tyc, Boyadjian, Allam, & Brasil-Neto, 2012) or hotspots (Bashir et al., 2013; Lüdemann-Podubecká & Nowak, 2016) of these different muscles' MCRs. To the best of our knowledge, the clearest recent demonstration of the somatotopy gradient of the hand TMS MCRs has been made using “along-sulcus” TMS mapping, when one investigates MEPs produced by stimulation only along the central sulcus, taking into account its shape (Dubbioso et al., 2017; Raffin et al., 2015; Raffin &

---

**FIGURE 9** Correlation matrices for inter-muscle topography parameters in the individual space (for different muscle cortical representation (MCR) pairs: 1—APB-ADM, 2—ADM-EDC, 3—APB-EDC) across days. Diagonal elements in the matrix indicate within-muscle MCR comparison, while off-diagonal elements indicate between-muscle MCR comparison. COG, center of gravity; EMD, earth mover’s distance; FWER, family-wise error rate
Schieber, 2019). We aimed to confirm and extend such a somatotopy gradient among the investigated hand muscle MCR COGs using a large grid approach. We observed that APB MCR COGs are more lateral compared to ADM and EDC MCR COGs. Interestingly, in two of our participants all three muscle MCRs were shifted rostrally, so it was impossible to trace MEPs along the central sulcus in a substantial part of the “hand knob” (see example in Figure 5). In this case, we believe that such placement of points might correspond to a “dorsal premotor” sub-type of TMS MCRs, as opposed to the “M1” sub-type (Dubbioso, Sørensen, Thielscher, & Siebner, 2019). This supports further the notion that in order to obtain a comprehensive motor mapping in a variety of subjects one should use an extended grid to capture individual profiles of MCRs. Considering M1 division to the “new” caudal M1 (having more monosynaptic connections with the spinal motoneurons) and the “old” rostral M1 (Lemon, 2019; Rathelot & Strick, 2006) we investigated whether the “along-sulcus” MCRs (where presumably mostly “new” M1) would be more segregated compared to the “whole” MCRs. On the contrary, we found that the mediolateral somatotopy gradient is greater for the “whole” TMS mapping approach compared to the “along-sulcus” one. So, at least for our data, it appears that additional features relating to whole M1 TMS mapping do not mask the within-limb TMS somatotopy compared to the along-sulcus-only approach and might in fact provide additional topographic aspects.

We also checked the COG somatotopy gradient for the MNI version of the MCRs and showed that APB MCR is again the most lateral one. Comparing MNI MCRs across subjects, we found that the ADM MCR area tends to be smaller than the APB and EDC MCR areas, and for EDC, this difference is significant. While in the “classical homunculus” the forearm representation is not only more medial but also smaller than the hand one, such a direct “homunculus” interpretation might be speculative (Nazarova & Blagovechtchenski, 2015; Schieber, 2001) and is not completely accurate even for the very same data from the original Penfield & Boldrey article (Penfield & Boldrey, 1937), as shown in the recent re-analysis (Catani, 2017). For TMS mapping, to our knowledge, no clear difference has been reported for the sizes of the proximal upper limb muscle MCRs versus intrinsic hand muscle MCRs (Devanne et al., 2006). Considering that MNI normalization, being routine for many brain mapping techniques such as fMRI or magnetoencephalography is still not a common procedure for TMS mapping (see examples of TMS data MNI normalization (Grab et al., 2018; Kraus & Gharabaghi, 2016; Niskanen et al., 2010; Weiss et al., 2013), we believe that this confirmation of the within-hand TMS somatotopy gradient and MCR size differences in MNI space may be useful for future studies using multiple muscle TMS mapping across subjects. There is a possibility that the difference between EDC and ADM TMS MCRs can be partially attributed to the parameters of TMS based on 110% of APB RMT, since it was reported that the degree to which the MCR area depends on stimulation intensity may vary among muscles (Thordstein, Saar, Pegenius, & Elam, 2013). However, we believe that this cannot be a major cause of the MCR area difference because no substantial difference among hand distal muscle RMTs either in healthy subjects (Schabrun & Ridding, 2007; Ziemann, Ilić, Alle, & Meintzschel, 2004), or stroke patients (in the contralesional hemisphere) (Nazarova et al., 2021) was found in previous studies. Importantly, the difference between EDC and ADM MCRs is not prone to cross-talk between muscles, because they are located far from each other in the upper limb (Selvanayagam, Riek, & Carroll, 2012).

4.2 | Reliability of single-muscle MCRs

A good understanding of TMS MCR reliability ranges is crucial for the TMS mapping application in longitudinal studies for the assessment of motor cortex plasticity. Here, we investigated both relative reliability, allowing stratifying subjects/patients based on their MCRs, and absolute reliability, reflecting the minimal change of TMS MCR parameters, which may be traced with the current nTMS mapping approach.

4.2.1 | Relative reliability

In the early TMS mapping studies without navigation, the relative reliability of the MCR parameters was shown to be primarily high (Malcolm et al., 2006; Plowman-Prine et al., 2008; Wolf et al., 2004). These observations were further confirmed using non-individual brain nTMS (Cavaleri et al., 2018; Jonker et al., 2019; Sankarasubramanian et al., 2015; van de Ruit et al., 2015), as well as individual brain nTMS (Forster et al., 2014; Kraus & Gharabaghi, 2016; Solmann et al., 2013; Zdunczyk et al., 2013). The problem is that only COG location reliability was always reported in TMS mapping papers, while the inclusion of other parameters was inconsistent. Using individual brain nTMS mapping it was shown that MCR parameters such as areas and volumes may be less reliable compared to COGs and mean MEP per MCR (Kraus & Gharabaghi, 2016). Previously, the calculations of MCR parameters were based on variable custom-made scripts, so we aimed at accessing the reliability of MCR parameters obtained using the TMSmap software (Novikov et al., 2018), which we suggested as a possible unifying approach for the fast and easy quantitative assessment of TMS motor mapping results. We observed high ICC values for all the investigated standard metrics of TMS MCRs (areas, volumes, mean MEPs, hotspot, and COG locations), with the highest ICC values for COGs and hotspots (ICC > 0.9). However, for the normalized MNI data, the ICC for hotspots and COGs was substantially lower because of the lower data variance among the brains after normalization, but still generally remained good.

4.2.2 | Absolute reliability

Absolute reliability (which may be reported by SEM, SDC, or limits of agreement) is crucial for the method's use on an individual level in clinical practice or sport (Atkinson & Nevill, 1998; Beaulieu, Flamand, et al., 2017). The absolute reliability of TMS motor mapping results has been described in the literature less frequently than the relative...
one (Jonker et al., 2019; Ngomo et al., 2012; Potter-Baker et al., 2016; van de Ruit et al., 2015). To the best of our knowledge, previous TMS motor mapping studies reporting classic absolute reliability metrics, were performed using non-individual MRI navigation, except one study (Ngomo et al., 2012). Yet the importance of considering individual anatomy in TMS motor mapping has been emphasized (Bashir et al., 2013; Raffin et al., 2015). When absolute reliability was reported, it was recommended to be interpreted “with caution on an individual level” (Jonker et al., 2019). In our study, we provide a comprehensive range of SDC values for different MCR parameters for three hand muscles. We can also interpret SDC as being generally sizeable: a particularly thought provoking SDC was found for the hotspot coordinates (around 1 cm), it also corresponds with a recent finding that hotspot may be considered as an area rather than a point taking into account MEPs variability and EF-spread (Reijonen et al., 2020). Hotspot is a crucial parameter for TMS motor mapping studies (Bashir et al., 2013; Rossini et al., 2015; Sollmann et al., 2013), where shifts of just several millimeters between hotspots of different MCRs are reported (Bashir et al., 2013). Hotspot is also highly relevant for a wide range of TMS applications with repetitive sessions of stimulation, including therapeutic TMS. Thus, we suggest that for longitudinal TMS design (either for diagnostic or therapeutic purposes) it is important to recheck the hotspot location with a roughly 1 cm radius, at least every several days.

4.2.3 | Muscle-specificity

We were interested in the possibility of reliably tracing musclespecific features of different muscle MCRs. Thus, we introduced a term “muscle-specificity” of the TMS MCR parameter, which we defined as the possibility to predict a given muscle among others (but only when using the data across subjects), using a certain MCR parameter. We observed that among standard parameters of single-muscle MCRs only mean MEP tended to be muscle-specific. Additionally, we investigated the reliability of a more novel parameter of the EP, which we previously implemented in the TMSmap software (Novikov et al., 2018), similar to the EP probed with the along-sulcus mapping approach (Raffin et al., 2015; Raffin & Siebner, 2019). We showed that the difference between the EPs, reflected in the EMD values (Novikov et al., 2018; Rubner et al., 1998) of the same muscle between days being smaller than for the different muscles MCR EPs between days. We believe that the idea of the MCR EP is important because it relates to the key phenomenon of motor cortex organization—convergence (Schieber, 2001), manifested in the well-known fact that different neurons in M1 may be involved in a given muscle activity depending on the goal/motor task or other conditions (Capaday, Ethier, Van Vreeswijk, & Darling, 2013). Moreover, using fMRI it has been shown that a single-muscle MCR is not homogeneous in its brain connectivity patterns (Smith et al., 2017). A similar approach of discrete peak calculation in an MCR was proposed recently (Elgueta-Cancino, Marinovic, Jull, & Hodges, 2019; Massé-Alarie et al., 2017), but it has been already shown to be less reliable than the standard TMS MCR metrics (Cavalieri et al., 2018). We also consider EPs comparison using EMD to be more direct than discrete peak calculation because while using EMD one does not need to define an arbitrary threshold level of the peak, and it is possible to account for the complex two-dimensional shape of the MCR. However, further validation of the MCR EP comparison using EMD in an interventional longitudinal TMS mapping studies is necessary.

4.2.4 | Reliability of the MCR overlaps

While we confirmed the somatotopy gradient between APB MCR versus ADM and EDC MCRs, using COG comparison, the relative reliability of the shifts between different muscle MCR COGs and hotspots is very poor and these metrics are not muscle-specific. We evaluated MCR overlap reliability as another parameter of the within-limb somatotopy. Extensive overlaps among different muscle MCRs were described from the early TMS studies (Devanne et al., 2006; Melgari, Pasqualetti, Pauri, & Rossini, 2008; E. M. Wassermann et al., 1993). Such overlaps reflect yet another key principle of M1 organization—its divergence, meaning that the output of a single M1 neuron may reach multiple spinal motoneurons, resulting in the activation of different limb segments (Schieber, 2001). It was demonstrated both non-invasively by TMS (Gerachshenko, Rymer, & Stinear, 2008) and invasively by microstimulation (M. Graziano, 2006), that the stimulation of the same cortical point may evoke activity in different muscles depending on a number of factors such as limb position (M. Graziano, 2006), anticipated movement (Gerachshenko et al., 2008; Uehara, Morishita, Kubota, & Funase, 2013), and so forth. Moreover, it was recently reported that in pathological cases, such as a tetraplegia, the “hand knob” in the precentral gyrus may be tuned not only to other upper limb segments but to the entire body (Willett et al., 2020). There is also some proof that the extent of TMS MCR overlaps may change in pathological conditions like dystonia or chronic pain (Schabrun et al., 2009; Schabrun et al., 2015). We observed that the overlaps between different muscle MCRs, and, to a lesser extent, EMDs between the different muscle MCRs, were more reliable and tended to be muscle-specific. To our knowledge, this is the first demonstration of the reliability of the inter-muscle interactions for multiple muscle TMS motor mapping using individual MRI data.

Another hypothesis about MCR overlaps is that they may be a neural substrate underlying muscle synergies (Beaulieu, Flanmand, et al., 2017; Capaday et al., 2013; Piltkänen et al., 2017; Schabrun et al., 2009; Schabrun et al., 2015). For normal synergies, there are many clues about cortical involvement and one of the most striking example was obtained using a prolonged electrical stimulation of the motor cortex, which yields synergistic motor patterns both in animals (Brown & Teskey, 2014; Graziano et al., 2002, Graziano et al., 2012; Graziano, Taylor, & Moore, 2002) and humans (Desmurget et al., 2014). As for the after-stroke pathological synergies, it has been discussed that their primary source may lay in the medulla...
(Karbasforoushan, Cohen-Adad, & Dewald, 2019; McPherson et al., 2018; Zaaim, Dean, & Baker, 2018), but there are also data that the MCR overlap might at least partly represent a cortical manifestation for this phenomenon as well (Giszter, 2015; Huffmaster, Van Acker, Luchies, & Cheney, 2018; Klockov, Khizhnikova, Nazarova, & Chernikova, 2017). Importantly, being a prominent phenomenon in humans, pathological synergies do not have a good animal model (Klockov et al., 2017; Krakauer & Carmichael, 2017). So, it is pretty unavoidable to primarily use non-invasive techniques like TMS to study them. Therefore, finding the high reliability of MCR overlaps and their muscle-specificity may be encouraging for new studies where such overlaps may be used as a traceable parameter reflecting training/rehabilitation dedicated to synergy manipulation.

5 | METHODOLOGICAL CONSIDERATIONS AND FUTURE STUDIES

An important strength of this study is the simultaneous investigation of multiple muscle MCRs. To our knowledge, this is the first work where the absolute reliability of multiple muscle MCRs was probed, using individual brain navigated TMS mapping for the whole motor cortex, in addition to the along-sulcus TMS mapping (Dubbioso et al., 2017; Raffin et al., 2015). Another feature of this study is its robust mapping design: we used up to 350 points per subject, because the literature about the necessary number of points for TMS motor mapping is contradictory: while some parameters of MCR were reported to be reliably traced already with 60 stimuli (van de Ruit et al., 2015), there have also been indications that accuracy of MCR parameters continued to increase without saturation up to a much higher number of stimuli (Nazarova et al., 2019; Sinitsyn et al., 2019). Another methodological strength of our study is the homogenous group of male participants with no special motor skills. Yet it may also be viewed as a limitation, meaning that in a more diverse multi-gender population the reliability of motor parameters may be less pronounced, thus a separate reliability study in females, taking, for instance, into account the menstrual cycle, may be needed. Another issue is that we probed only hand muscles and thus the reliability for distal versus proximal upper limb muscle MCRs is left for further investigation. Also, we did not employ any motor task to assess motor abilities, thus, the question of the functional relevance of TMS within-limb somatotopy and its reliability should be investigated in the future. It would also be interesting to add an assessment of the evoked movements in parallel with EMG recording, using accelerometry as in (Zartl, Kapfer, & Muellbacher, 2014) or motion capture.

A technical limitation of the study is that when tracing several muscle MCRs, the characteristics of the MCR parameters distribution (such as normality or homoscedasticity) may vary across muscles, so, for instance, the question of the logarithmic transformation for the parameters has to be carefully considered. Also, as we have mentioned above, we used the same intensity for all the muscles MCRs creation based on the RMT of APB muscle, which may be considered as a technical limitation. However, we have chosen this approach because: (a) the difference among hand distal muscles’ thresholds is not significant based on the literature (Nazarova et al., 2021; Schabrun & Ridding, 2007; Ziemann et al., 2004), (b) multiple hotspots and RMTs hunting is time consuming, and (c) an approach when the intensity is based on just one of the investigated muscles RMT is used in clinical nTMS mapping (e.g., Sollmann et al., 2017).

One general limitation applicable not only to our work but to all TMS studies, is the spatial resolution of TMS. We assume that the simultaneous activation of several muscles by TMS at one point is a result of several key factors: (a) the non-focality of TMS itself: indeed, while a peak of the EF is confined to just several millimeters, a spread of the induced EF leads to the activation of neighboring cortical areas with lower intensities (Wassermann et al., 2008); (b) the primarily indirect TMS effect on the pyramidal neurons (Seo, Schaworonkow, Jun, & Triesch, 2016; Spampinato, 2020); (c) the co-location of the cortical neuronal populations, innervating different spinal motoneurons pools (Capaday et al., 2013; Schieber, 2001); and (d) the cross-talk between the muscles at the peripheral level when using surface EMG (Selvanayagam et al., 2012). TMS focality depends on the coil configuration and on the intensity, shape and direction of the pulse (Koponen, Nieminen, Mutanen, Stenroos, & Ilmoniemi, 2017; Rossi et al., 2015; Sommer et al., 2006, 2018; Tugin et al., 2020). We utilized the figure-of-eight coil used for nTMS presurgical mapping (Krieg, 2017) and a biphasic pulse shape, allowing to use less intensity (Raffin et al., 2015). We used a biphasic pulse shape because it is the most common in TMS mapping studies with patients (Lüdemann-Podubecká & Nowak, 2016; Takahashi, Vajkoczy, & Pichlt, 2013). However, considering that a biphasic pulse may activate two distinct neuronal pools (Sommer et al., 2018); it may be informative to investigate how the reliability of the within-limb TMS somatotopy differs when using monophasic current configurations. Yet another aspect to consider is a more advanced induced EF modeling using several compartments for the head models (Stenroos & Koponen, 2019), which may be especially warranted when investigating structurally affected brains, such as brain with a cortical stroke lesion or tumor (Minjoli et al., 2017).

6 | CONCLUSIONS

Until now, the use of nTMS mapping for longitudinal purposes in fundamental and clinical studies remains challenging, especially when multiple MCRs are probed. In this work, we confirmed the existence of the somatotopy gradient for the hand MCR COGs for the whole motor cortex nTMS mapping in addition for the along-sulcus TMS mapping approach. Adding to previous reliability studies, we confirmed a high relative reliability of the standard MCR parameters for three hand muscles and present a range of SDC values for the MCR parameters. We provide novel evidence that inter-muscle MCR interaction can be reliably traced using MCR area overlaps, while shifts between the COGs and hotspots of different muscle MCRs are not suitable for this purpose. Our work has also a practical perspective—a
high reliability of MCR overlaps allows us to suggest them as a possible cortical biomarker for tracking the neuronal changes associated with the training/rehabilitation aimed at the modifications of muscle synergies.

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DATA AVAILABILITY STATEMENT
Data supporting the findings of this study are available from the corresponding author upon reasonable request.

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