## RESEARCH ARTICLE

# Mapping Motor Cortical Network Excitability and Connectivity Changes in De Novo Parkinson's Disease

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ABSTRACT: Background: Transcranial magnetic stimulation-electroencephalography (TMS-EEG) has demonstrated decreased excitability in the primary motor cortex (M1) and increased excitability in the pre-supplementary motor area (pre-SMA) in moderate-advanced Parkinson's disease (PD).

**Objectives:** The aim was to investigate whether these abnormalities are evident from the early stages of the disease, their behavioral correlates, and relationship to cortico-subcortical connections.

Methods: Twenty-eight early, drug-naive (de novo) PD patients and 28 healthy controls (HCs) underwent TMS-EEG to record TMS-evoked potentials (TEPs) from the primary motor cortex (M1) and the pre-SMA, kinematic recording of finger-tapping movements, and a 3T-MRI (magnetic resonance imaging) scan to obtain diffusion tensor imaging (DTI) reconstruction of white matter (WM) tracts connecting M1 to the ventral lateral anterior thalamic nucleus and pre-SMA to the anterior putamen.

Results: We found reduced M1 TEP P30 amplitude in de novo PD patients compared to HCs and similar pre-SMA TEP N40 amplitude between groups. PD patients exhibited smaller amplitude and slower velocity in fingertapping movements and altered structural integrity in WM tracts of interest, although these changes did not correlate with TEPs.

Conclusions: M1 hypoexcitability is a characteristic of PD from early phases and may be a marker of the parkinsonian state. Pre-SMA hyperexcitability is not evident in early PD and possibly emerges at later stages of the disease. © 2024 The Author(s). Movement Disorders published by Wiley Periodicals LLC on behalf of International Parkinson and Movement Disorder Society.

Key Words: Parkinson's disease; transcranial magnetic stimulation-electroencephalography; diffusion tensor imaging; motor cortex; pre-supplementary; motor area

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Motor impairment in Parkinson's disease (PD) is thought to stem from nigrostriatal degeneration, leading to dysfunctional activity within the basal ganglia-thalamo-cortical network.<sup>[1,2](#page-8-0)</sup> These changes cause an abnormal corticostriatal transmission, resulting in a net increase in basal ganglia–mediated inhibition of the motor thalamus projections to the motor areas.<sup>[3](#page-8-0)</sup> This process is believed to alter the excitability of motor cortical areas, potentially leading to bradykinesia, one of the cardinal signs of  $PD<sup>4-9</sup>$  $PD<sup>4-9</sup>$  $PD<sup>4-9</sup>$  However, despite extensive research, the pathophysiological significance and the behavioral correlates of these motor cortical changes in PD remain elusive.

A novel approach to assess motor cortical activation in humans involves using transcranial magnetic stimulation

coupled with electroencephalographic recordings (TMS-EEG)[.10](#page-8-0) This technique directly probes the excitability and connectivity of cortical areas with high temporal resolution by recording the TMS-evoked potentials (TEPs). TEPs manifest as a sequence of positive and negative components occurring within  $\sim$ 300 ms after the TMS pulse. TEPs provide a comprehensive view of the interplay between local excitability and effective connectivity of the stimulated area within its functional network.<sup>[11,12](#page-8-0)</sup> In a previous TMS-EEG study, we observed a reduction in primary motor cortex (M1) excitability and an increase in pre-supplementary motor area (pre-SMA) excitability in PD patients with moderate to advanced disease stages. $13$  Notably, in these patients, changes in M1 excitability correlated with the severity of clinically evaluated bradykinesia, and both M1 and pre-SMA changes were normalized by dopaminergic therapy. Furthermore, TEPs from motor areas reflect propagation through reverberating cortico-basal ganglia-thalamo-cortical pathways, $14-16$ implying that TEP abnormalities might represent cortical changes secondary to underlying basal ganglia dysfunctions. Yet several issues remain unclear, and unraveling the mechanisms underlying these motor cortical changes and their clinical implications will enhance our understanding of PD's pathophysiology.

The first aim of this study was to determine if the M1 and pre-SMA excitability changes are also evident in the early stages of the disease, before the introduction of dopaminergic therapy, and if they correlate with bradykinesia. Second, the study aimed to investigate whether TEP changes in PD mirror abnormal structural integrity in pathophysiologically relevant white matter (WM) tracts. $17,18$ 

To accomplish the first aim, we compared TEPs from M1 and pre-SMA between early, drug-naive PD patients ("de novo") and a control group of healthy controls (HC). We also examined the correlation between TEP alterations and objective kinematic measures of bradykinesia during finger-tapping movements. For the second aim, we assessed the structural integrity of cortico-subcortical connections using diffusion tensor imaging (DTI) and investigated potential correlations between DTI changes and TEP abnormalities.

### Patients and Methods

### **Participants**

We consecutively enrolled 28 de novo PD patients (18 men, median age: 57) (range: 43–81) as well as 28 age- and gender-matched HCs (19 men, median age: 60, range: 42–77) (Table [1A](#page-2-0)). Patients were enrolled in the Movement Disorders Outpatient Unit at the Department of Human Neurosciences, Sapienza University of Rome, Italy. The study protocol was approved by the

institutional review board and conducted in accordance with the latest revision of the Declaration of Helsinki. All patients gave their written informed consent before participating in the study.

Inclusion criteria for patients included a PD diagnosis confirmed by a movement disorder expert neurologist based on international criteria<sup>[19](#page-8-0)</sup> and had to be drug naive for antiparkinsonian medications, with an Hoehn & Yahr (H&Y) stage <II, and with clinical onset  $\leq$ 2 years.<sup>[20](#page-8-0)</sup> Exclusion criteria included a diagnosis of a neurological or psychiatric condition other than PD and cognitive decline defined as a Montreal Cognitive Assessment (MoCA) score <26.

#### Experimental Sessions

Participants underwent a clinical evaluation, a TMS-EEG, and kinematic recording on a single day. Participants also underwent a magnetic resonance imaging (MRI) session on a separate day, between 7 and 14 days from the other experimental day. We focused on the most affected upper limb (clinical and kinematic assessments) and contralateral hemisphere (TMS-EEG and kinematic recordings) in PD patients and matched it one to one with the corresponding side (either left or right) of HCs. In particular, for each PD patient who had a greater bradykinesia on the right upper limb, the limb (and left hemisphere) was assessed in a HC of similar age and the same gender, and vice versa when patients had the left side more affected. This matching method an ensured equal distribution of the left and right sides investigated between groups.

#### Clinical Assessment

Clinical assessment included the determination of disease duration, H&Y scale; Movement Disorder Societysponsored Unified Parkinson's Disease Rating Scale (MDS-UPDRS), Part III; MoCA; and Nonmotor Symptoms Scale. We defined the most affected upper limb in PD patients as the limb with the highest summed scores of MDS-UPDRS, Part III, subitems 3.4, 3.5, and 3.6.

#### TMS-EEG Study

Participants were comfortably seated on a chair designed for TMS (EMS, Italy), with forearms pronated and resting on armrests; they were then instructed to stay relaxed and keep their eyes open, fixed on a point (a black cross) displayed on a PC screen at 70 cm. A Magstim SuperRapid stimulator (Magstim Company, UK) connected to a 70-mm figure-of-eight coil delivered single-pulse TMS. Using a neuronavigation system (SofTaxic, EMS, Italy) with an optical tracking system (Polaris Vicra, Northern Digital Inc., Canada), we sampled 23 points from each participant's scalp. Using nonlinear fitting, we adapted the reconstructed brain to the Montreal Neurological Institute (MNI)

<span id="page-2-0"></span>TABLE 1 Demographic and clinical data of Parkinson's disease patients and healthy controls

Groups			Subjects Gender Age (y)	Duration (y)	$H\&Y$	MoCA	<b>NMSS</b>	<b>MDS-UPDRS,</b> Part III	Most bradykinetic arm
A. All participants: TMS-EEG study									
De novo <b>PD</b>	28	18M 10F	$57(43-81)$	$1(1-2)$			$1(1-2)$ 27 $(26-30)$ 12.5 $(0-150)$	$19(7-38)$	$4.5(0-7)$
HCs	28	19M 9F	$60(42-77)$						
B. Subgroup: TMS-EEG, kinematic, and DTI studies									
De novo PD.	20		$12M$ 8F 57 (43-71)	$1(1-2)$	$1(1-2)$	$27(26-30)$	$11(0-150)$	$17.5(7-38)$	$4.5(0-7)$
HCs	20		13M 7F 60 (44-73)						

Most bradykinetic arm = summed scores of MDS-UPDRS, Part III, subitems 3.4, 3.5, and 3.6 of more affected arm; values are expressed as median (minimum–maximum). Abbreviations: H&Y, Hoehn & Yahr; MoCA, Montreal Cognitive Assessment; NMSS, Nonmotor Symptoms Scale; MDS-UPDRS, Movement Disorder Society-Sponsored Unified Parkinson's Disease Rating Scale; TMS-EEG, transcranial magnetic stimulation-electroencephalography; PD, Parkinson's Disease; M, male; F, female; HC, healthy control; DTI, diffusion tensor imaging.

space to monitor coil positioning over the hot spot for all stimulation sites.

M1 was stimulated over the hot spot, evoking the most consistent motor-evoked potential (MEP) in the contralateral first dorsal interosseous (FDI) muscle with the posterior–anterior current direction. Resting motor threshold (RMT) was defined separately for each M1 as the minimum intensity required to elicit MEPs of  $\geq$ 50-µV peak-to-peak amplitude in at least 5 of 10 consecutive trials. For pre-SMA stimulation, the coil was placed with the handle pointing backward, parallel to the interhemispheric fissure, and centered over the MNI coordinates  $x = 0$ ,  $y = 10$ ,  $z = 68<sup>13</sup>$  $z = 68<sup>13</sup>$  $z = 68<sup>13</sup>$  The coil orientation was chosen following recent evidence showing that this orientation ensures substantial estimated electric field coverage of the pre- $SM^{13}$  $SM^{13}$  $SM^{13}$  and larger TEPs com-pared to other orientations.<sup>[21](#page-8-0)</sup> In two separate recording sessions, 100 TMS pulses were delivered at 110% RMT intensity during continuous EEG over either M1 or the pre-SMA randomly. For M1 stimulation, electromyography (EMG) was also recorded from the contralateral FDI through pairs of Ag/AgCl surface electrodes arranged in a belly-tendon montage. EMG signal was bandpass filtered between 10 and 1000 Hz, amplified  $(\times 1000)$  (D360, Digitimer, UK), and digitized at 5 kHz (CED 1401, Cambridge Electronic Design, UK).

EEG was recorded from 32 passive electrodes on a TMS-compatible cap (BrainCap, EASYCAP, Germany) mounted on an elastic cap, according to the international 10-20 system, namely Fp1, Fp2, Afz, F7, F3, Fz, F4, F8, FC5, FC1, FCz, FC2, FC6, T7, C3, Cz, C4, T8, TP9, CP5, CP1, CP2, CP6, TP10, P7, P3, Pz, P4, P8, O1, O2, and Iz. All electrodes were grounded to Fpz and online referenced to POz. Impedance for each channel was kept below 5 kΩ. The EEG signal was bandpass filtered at DC-2.5 kHz and digitized at 10 kHz using a TMS-compatible system (NeurOne, Bittium Corporation, Finland). To reduce auditory contamination, participants used ear protectors (signalto-noise ratio  $= 30$ ) over earphones that consistently emitted a sound to conceal the TMS click. To reduce the TMS click bone conduction and the feeling of coil vibration on the scalp, a 0.5-cm foam padding was positioned beneath the coil.

TMS-EEG signal preprocessing was performed as reported in our previous study,  $13,22$  using the EEGLAB<sup>[23](#page-9-0)</sup> and TMS-EEG Signal Analyser<sup>22</sup> open-source MATLAB (version 2022b) toolboxes. Signal preprocessing steps are presented in the Supplementary materials (Data S1). The cleaned TMS-EEG epochs were converted into reference-free, current source density (CSD) estimates using "CSD,"<sup>[24](#page-9-0)</sup> an open-source toolbox for Fieldtrip.<sup>25</sup> Final TEPs were obtained by averaging TMS-EEG epochs (averaged number of epochs  $94 \pm 3$ ) in each block. As our aim was to characterize previously identified cortical anomalies, we focused specifically on regions of interest (ROI) and times of interest that demonstrated differences between PD patients and HCs, as well as between OFF-and ON-dopaminergic states in our previous research.<sup>[13](#page-8-0)</sup> Based on this, the P30 amplitude after M1 stimulation was measured at its peak value between 23 and 33 ms in the TEP averaged over C3 and FC1 electrodes. Similarly, the N40 amplitude post pre-SMA stimulation was measured between 34 and 44 ms in the TEP averaged over Fz and Afz electrodes.

#### Kinematic Evaluation

Kinematic recordings were conducted using an optoelectronic system (SMART motion system, BTS Engineering), with three infrared cameras. This system tracked the motion of five reflective markers attached

to the participants' hands, enabling the measurement of three-dimensional hand movements.<sup>[26,27](#page-9-0)</sup> The study involved participants engaged in a finger-tapping exercise, and the participants were asked to tap their index finger repetitively against their thumb and to perform the movement as widely and as quickly as possible. Reflective markers were attached to the tip of the index finger and the tip of the thumb, and additional markers were placed on the hand, at the head and base of the second metacarpal bone, and on the base of the fifth metacarpal bone. A preliminary trial was conducted before the official recording. The data collection consisted of three 15-second finger-tapping sessions interspersed with 60-second rest periods to minimize fatigue.

A dedicated software tool (SMART Analyzer, BTS Engineering) was used to quantify the movement mean amplitude, mean velocity, and decrement in amplitude and velocity during the repetitive finger-tapping task namely the sequence effect—as described in previous studies.<sup>[26,27](#page-9-0)</sup>

#### MRI Study

During the MRI acquisition, participants were instructed to relax and rest to avoid movement artifacts. All participants underwent a standardized MRI protocol on a 3-T scanner (Siemens Magnetom Verio) and a 12-channel head coil designed for parallel imaging (generalized autocalibrating partially parallel acquisitions). The following MRI sequences were obtained: (a) high-resolution 3D, T1-weighted magnetization-prepared rapid acquisition with gradient echo: repetition time  $(TR) = 1900$  ms, echo time (TE) = 2.93 ms, flip angle =  $9^{\circ}$ , field of view (FOV)  $= 260$  mm, matrix  $= 256 \times 256$ , 176 contiguous 1-mmthick sagittal slices; (b) dual turbo spin-echo proton density and T2-weighted images:  $TR = 3320$  ms, TE = 10/103 ms, FOV = 220 mm, matrix =  $384 \times 384$ , 25 4-mm-thick axial slices, 30% gap; (c) DTI single-shot, echo-planar, spin-echo sequence with 10 interspersed volumes of  $b = 0$  (b0) and 64 gradient directions,  $TR = 4600$  ms,  $TE = 78$  ms, multiband acceleration factor  $= 2$ , monopolar diffusion scheme,  $FOV = 192$  mm, matrix =  $96 \times 96$ , b = 1000 s/mm<sup>2</sup>, 72 contiguous axial 2-mm-thick slices.

An expert radiologist (P.P.) examined all MRIs to exclude the presence of concomitant brain lesions and focal T2 WM hyperintensities.

Structural preprocessing was performed using FMRIB's Software Library (FSL), version 6.0.5.1 ([https://fsl.fmrib.](https://fsl.fmrib.ox.ac.uk/fsl) [ox.ac.uk/fsl\)](https://fsl.fmrib.ox.ac.uk/fsl). Diffusion data were visually inspected for artifacts and preprocessed using different tools from FDT (FMRIB Diffusion Toolbox, part of FSL). Images were corrected for eddy current distortion and head motion using a 12-parameter affine registration to each subject's first no-diffusion weighted volume, and the gradient directions were rotated accordingly. $28$  Nonbrain tissue

was removed from the eddy-corrected images using the Brain Extraction  $Tool<sub>2</sub><sup>29</sup>$  creating a binary mask of the brain. Then, maps of fractional anisotropy (FA), mean diffusivity (MD), axial diffusivity (AD), and radial diffusivity (RD) were estimated at the individual level using the DTIFIT tool by fitting a tensor model to the eddy-corrected and brain-masked diffusion data. Registration between diffusion, structural, and standard space images was performed within FDT. Transformation matrices and their inverses were created to transform images between spaces.

To reconstruct WM tracts between cortical and subcortical structures, ROIs were defined. We decided a priori to restrict our DTI analysis to the corticosubcortical tracts most likely to contribute to TEP changes at pre-SMA and M1 levels in PD. We investigated structural connectivity between the pre-SMA and the anterior putamen because pre-SMA hyperactivation in PD is associated with increased connectivity to this region, $30$  to which pre-SMA predominantly projects.<sup>[31](#page-9-0)</sup> We also investigated structural connectivity between the ventral lateral anterior thalamic nucleus (Vla)—the main output station of the basal ganglia<sup>[32](#page-9-0)</sup>—and M1, as M1 hypoactivation in PD likely follows a decreased thalamocortical output, as a consequence of dopaminergic denervation.<sup>3</sup> We also selected ROIs to reconstruct a control tract connecting the primary visual cortex (V1) and the pulvinar thalamic nucleus. This nonmotor control tract was included to verify the specificity of our possible correlation findings between motor tract DTI changes and TEPs from motor areas, with the hypothesis that TEP changes would not correlate with this tract.

Cortical ROIs were created using 12-mm-radius spheres centered on reference MNI coordinates. The pre-SMA ( $x = 0$ ,  $y = 10$ ,  $z = 68$ ) sphere was centered on TMS stimulation sites and then divided on the sagittal plane  $x = 0$  in the right and left regions. For each ROI, we selected the most affected side in PD patients (ie, contralateral to the most bradykinetic upper limb) and matched it one to one with the corresponding side of HCs. For the M1 ROI, we identified the hand's area based on the method described by Chris Rorden $33$  (left:  $x = -34$ ,  $y = -22$ ,  $z = 52$ ; right:  $x = 34$ ,  $y = -22$ ,  $z = 52$ ). The center of V1 was selected based on previous works (left:  $x = -8$ ,  $y = -76$ ,  $z = 10$ ; right:  $x = 8$ ,  $y = -76$ ,  $z = 10$ ).<sup>[34](#page-9-0)</sup> Left- and right-putamen ROIs were selected from the Harvard-Oxford Subcortical Structural Atlas ([https://fsl.fmrib.ox.ac.uk/fsl/fslwiki/](https://fsl.fmrib.ox.ac.uk/fsl/fslwiki/atralses) [atralses\)](https://fsl.fmrib.ox.ac.uk/fsl/fslwiki/atralses). The anterior portion of the putamen was obtained by dividing the putamen along the coronal plane that intersects the anterior commissure.<sup>[35](#page-9-0)</sup> Thalamic nuclei (Vla and pulvinar) were obtained for each subject (in structural space) using a multi-atlas segmentation technique (THOMAS [Thalamus Optimized Multi Atlas Segmentation]).<sup>[36](#page-9-0)</sup>

Finally, cortical and subcortical regions were transformed from standard or structural space into diffusion space using previously generated registrations and visually checked for accuracy.

Probabilistic tractography was performed within each participant's diffusion space using Bedpost $X^{37}$  $X^{37}$  $X^{37}$ with default parameters. We generated streamlined probability distribution maps between each predefined subcortical and cortical ROI. In each reconstructed map, we specified the subcortical region as the seed, the cortical region as the target, and the contralateral hemisphere as the exclusion mask. We also specified the cortical target region as a termination mask to identify the only and exact connections between the given seed and the target. $38$  Pathway probability maps were normalized for seed size by dividing the probability maps by the overall number of successfully generated streamlines, and spurious connections were removed by thresholding the resulting maps by  $5\%$ .<sup>[38](#page-9-0)</sup> Thresholded probability maps were then binarized and overlaid on FA, MD, AD, and RD individual maps, from which average values were extracted and used for further statistical analyses.<sup>[39](#page-9-0)</sup>

#### Statistical Analysis

An unpaired  $t$  test was conducted to compare the ages between groups. A  $\chi^2$  test was performed to compare gender distribution between groups.

We used Mann-Whitney U tests to assess the differences in TEP amplitude (M1 P30, pre-SMA N40) between de novo PD and HCs.

Mann-Whitney U tests were used to compare kinematic measures (mean amplitude, mean velocity, amplitude sequence effect, velocity sequence effect) and DTI parameters (FA, MD, AD, and RD) in each tract between the de novo PD and HCs.

Spearman's correlation coefficient was used to investigate possible correlations in the PD group between TEP amplitude (M1 P30, pre-SMA N40) and MDS-UPDRS, Part III, score, most affected upper-limb bradykinesia subscores, and with kinematic and DTI features found significantly different between the group of PD patients and HCs.

Results are reported as significant when  $P < 0.05$ . False discovery rate (FDR) correction for multiple comparisons has been applied to the kinematic, DTI, and correlation tests.

We conducted a power analysis using G\*Power for group comparisons in TEPs utilizing the Mann-Whitney U test, with estimated effect sizes of 0.7 for M1 and 0.8 for pre-SMA derived from our previous work.<sup>[13](#page-8-0)</sup> Setting α at 0.05 and aiming for a power of 0.8, the analysis determined that the required sample sizes were 28 for M1 and 21 for pre-SMA.

### **Results**

#### Clinical and Demographic Findings

All 28 de novo PD patients and all 28 HCs completed the TMS-EEG study (Table [1A](#page-2-0)). Analysis of age and gender distribution revealed no significant differences between groups (age:  $t(54) = 0.392$ ,  $P = 0.697$ ; gender:  $\chi^2 = 0.080$ ,  $P = 0.778$ ). In addition, 20 de novo PD patients and 20 HCs completed both the kinematic recordings and MRI studies (Table [1B](#page-2-0)). Similarly, we found no significant difference in age and gender distribution between the subgroups that completed the kinematic and MRI studies (age:  $t(38) = 0.563$ ,  $P = 0.576$ ; gender:  $\chi^2 = 0.107$ ,  $P = 0.774$ ). All participants were right handed and based on the most affected side in PD patients, the right upper limb (and thus the left hemisphere) was studied in 15 PD patients and 15 HCs, and the left upper limb was assessed in 13 patients and 13 HCs.

### Transcranial magnetic stimulationelectroencephalography

When stimulated over M1, de novo PD patients exhibited significantly smaller median P30 amplitude compared to HCs  $(2.17 \text{ vs. } 6.53 \text{ µV}, U = 234,$  $P = 0.01$ ). Removing an outlier in the HC group with values exceeding three standard deviations above the mean (53.24), the comparison remained significant (2.17 vs. 6.33  $\mu$ V, U = 234, P = 0.015) (Fig. [1\)](#page-5-0). In contrast, we found no significant difference in median N40 from pre-SMA stimulation between de novo PD patients and HCs  $(-3.22 \text{ vs. } -1.61 \text{ }\mu\text{V}, \text{ } U = 352,$  $P = 0.512$ ). Removing an outlier in the HC group with values exceeding three standard deviations above the mean  $(25.56)$  did not affect the result  $(-3.22)$ vs.  $-1.65 \mu V$ , U = 352, P = 0.66[2\)](#page-5-0) (Fig. 2).

#### Kinematic Analysis

Compared to HCs, de novo PD patients exhibited significantly lower median scores of mean amplitude (79.7 vs. 39.9 degrees<sup>o</sup>, U = 33, FDR-adjusted  $P < 0.001$ ) and mean velocity (714.6 vs. 373.3 degrees/s,  $U = 30$ , FDRadjusted  $P < 0.001$ , and significantly higher amplitude sequence effect  $(-0.10 \text{ vs. } -0.69 \text{ degrees/number of})$ movements,  $U = 66$ , FDR-adjusted  $P < 0.001$  and velocity sequence effect  $(0.48 \text{ vs. } -5.24 \text{ (degrees/s)/num--}$ ber of movements,  $U = 70$ , FDR-adjusted  $P < 0.001$ ) (Fig. S1).

#### Diffusion Tensor Imaging

Examples of reconstructed WM tracts are visualized in Supplementary Figure S2. PD patients exhibited significantly lower FA values with respect to HCs in all the reconstructed WM tracts (Vla–M1, pre-SMA– anterior putamen, pulvinar–V1) and significantly higher

<span id="page-5-0"></span>

FIG. 1. Grand average TMS-evoked potentials (TEPs) from primary motor cortex (M1) stimulation. (A, B) Butterfly plots; (C) current source density (CSD) within the region of interest (ROI). Common average reference (top) and CSD (bottom) topoplots in (D) healthy controls (HC) and (E) de novo PD patients. Yellow bars: time of interest, white stars: ROI. (F) P30 difference (\*P < 0.05). [Color figure can be viewed at [wileyonlinelibrary.com\]](http://wileyonlinelibrary.com)



FIG. 2. Grand average TMS-evoked potentials (TEPs) from pre-supplementary motor area (pre-SMA) stimulation. (A, B) Butterfly plots; (C) current source density (CSD) within the region of interest (ROI). Common average reference (top) and CSD (bottom) topoplots in (D) healthy controls (HC) and (E) de novo PD patients. Yellow bars: time of interest, white stars: ROI. (F) N40 difference. [Color figure can be viewed at [wileyonlinelibrary.com](http://wileyonlinelibrary.com)]

<b>Tracts WM property</b>	$HCs$ (mean $\pm$ SD)	PD (mean $\pm$ SD)	W	$P-value^{\star}$
Pre-SMA-anterior putamen				
AD	$0.0011199 \pm 0.0000518$	$0.0010918 \pm 0.0000463$	144.5	<b>NS</b>
FA	$0.4745723 \pm 0.0329053$	$0.4399771 \pm 0.0305911$	87	< 0.05
MD	$0.0007236 \pm 0.0000366$	$0.0007248 + 0.0000358$	220	<b>NS</b>
RD	$0.0005253 \pm 0.0000379$	$0.0005413 \pm 0.0000368$	255.5	NS.
$Vla-M1$				
AD	$0.0010890 \pm 0.0000769$	$0.0010501 \pm 0.0000564$	123	NS.
FA	$0.4996382 \pm 0.0486346$	$0.4396030 \pm 0.0339368$	53	< 0.01
MD	$0.0006863 \pm 0.0000406$	$0.0006951 \pm 0.0000326$	234.5	<b>NS</b>
RD	$0.0004849 \pm 0.0000434$	$0.0005179 \pm 0.0000338$	298.5	< 0.05
Pulvinar-V1				
AD	$0.0014460 \pm 0.0001926$	$0.0013094 + 0.0000860$	102.5	<b>NS</b>
<b>FA</b>	$0.5394423 \pm 0.0541856$	$0.4791342 \pm 0.0681371$	49	< 0.001
MD	$0.0008828 \pm 0.0001689$	$0.0008363 \pm 0.0000876$	206.5	<b>NS</b>
RD	$0.0006014 \pm 0.0001626$	$0.0005996 \pm 0.0000956$	276.5	<b>NS</b>

TABLE 2 Differences between Parkinson's disease patients and healthy controls for white matter measures for each tract

Differences were assessed using Mann-Whitney U test; NS:  $P > 0.05$ .

Abbreviations: WM, white matter; HC, healthy control; SD, standard deviation; AD, axial diffusivity; FA, fractional anisotropy; MD, mean diffusivity; RD, radial diffusivity; Vla, ventral lateral anterior nucleus; M1, primary motor cortex; V1, primary visual cortex.

\*FDR-corrected for multiple comparison.

RD in the Vla–M1 tract. No significant between-group differences were found in MD and AD values in the investigated tracts (Table 2).

#### TEP Correlations

We found no significant correlations between TEP P30 elicited from M1 or TEP N40 elicited from pre-SMA and clinical, kinematic, and DTI measures (Table [3\)](#page-7-0).

### **Discussion**

This study investigated M1 and pre-SMA excitability in early, de novo PD patients using TMS-EEG and their clinical, behavioral, and structural connectivity correlates. After TMS of the M1 contralateral to the most affected side, de novo PD patients exhibited lower TEP P30 amplitude than HCs. When pre-SMA was stimulated, we found similar TEP N40 amplitude in de novo PD patients and HCs. Kinematic analysis revealed smaller amplitude and slower velocity of finger-tapping movements in de novo PD patients compared to HCs. Structural MRI connectivity analysis uncovered notable differences between de novo PD patients and HCs in the tracts of interest connecting M1 to ventrolateral anterior (Vla) thalamic nucleus, the pre-SMA to the anterior putamen, and in the control tract from the pulvinar to

V1. There were no significant correlations between the M1 P30 amplitude and N40 amplitude and any clinical, kinematic, or DTI measures. Overall, the findings of the present study suggest that altered M1 excitability in PD is already present when motor signs first manifest, whereas pre-SMA abnormal hyperactivation is not. Motor cortical excitability alterations do not correlate with bradykinesia or structural changes in pathologically relevant cortical–subcortical tracts.

The first result of the present study is that de novo PD patients have a reduced TMS-evoked P30 amplitude from M1. The observed reduction in the P30 amplitude from M1 in de novo PD patients suggests that a dysfunction of cortical motor area excitability is present in the early stages of the disease. This extends previous findings from our group showing a decreased M1 P30 in moderate-advanced PD patients, which was modulated by dopaminergic treatment.<sup>[13](#page-8-0)</sup> The kinematic analysis showed that de novo PD patients had smaller and slower finger-tapping movements than HCs and a more signifi-cant sequence effect.<sup>[8](#page-8-0)</sup> We also found a lack of correlation between M1 TEP changes and clinical and objectively measured bradykinesia. The absence of correlation is likely not due to mild motor impairment in the patients we studied, as our findings, alongside previous studies, suggest that distinct features of bradykinesia, such as the sequence effect, are apparent or possibly more pronounced, even in the early stages of the disease.<sup>27</sup>

<span id="page-7-0"></span>



3.4, 3.5, and 3.6 of the more affected am. SE, sequence effect; pre-SMA, pre-supplementary motor area; AmPut, anterior putamen; FA, fractional anisotropic; Vla, ventral lateral anterior nucleus. M1, primary motor correx; R primary ΣÍ. nucleus. lateral anterior ventral Уlа. anisotropic: fractional area; AntPut, anterior putamen; FA, motor pre-SMA, pre-supplementary effect; sequence arm. SE. 3.4, 3.5, and 3.6 of the more affected<br>diffusivity.

The lack of correlation between M1 excitability, as assessed by TEPs, and the sequence effect expands upon previous findings,<sup>26</sup> which also reported no correlation between the sequence effect and corticospinal excitability, as measured by MEPs. Consequently, it appears that neither local M1 excitability nor corticospinal excitability plays a direct role in the pathophysiology of the sequence effect. The decreased TMS-induced activation of M1 may re flect a correlate of the impaired transition of this cortical area between akinetic and prokinetic states observed in  $PD<sup>40</sup>$  Studies using TMS coupled with deep brain stimulation have proposed that the neurophysiological mechanism for the impaired M1 activation results from an abnormal recruitment of the subthalamic nucleus via the hyperdirect pathway, $41$  notably that the latency of the P30 is compatible with a cortico-basal ganglia-thalamo-cortical loop recruited through TMS activation of the hyperdirect pathway (see further discussion on this point later).<sup>42,43</sup> However, it should be considered that TEPs were delivered in a resting condition, whereas bradykinesia mechanisms may be better exhibited during the preparation and execution of a movement.<sup>4,8</sup> This possibility could account for the observed lack of correlation between TEP measures and bradykinesia.

We observed no signi ficant differences in the pre-SMA N40 amplitude in de novo PD patients compared to HCs. This finding expands previous observations of larger N40 amplitude in patients in a moderate-to-advanced stage, $^{13}$ suggesting that pre-SMA hyperexcitability may develop at later disease stages. However, given the cross-sectional nature of data in the two studies, we can only make speculations about the relation between pre-SMA excitability changes and PD progression. Furthermore, the previous observation of pre-SMA hyperexcitability might have been in fluenced by the small sample sizes used, leaving open the possibility of a false positive. Future research, with larger sample sizes and longitudinal design, may confirm whether pre-SMA hyperexcitability is associated with advanced PD stages and whether it plays a pathophysio-logical<sup>30</sup> or compensatory role in motor impairment.<sup>[44,45](#page-9-0)</sup>

The second finding of the present study concerns whether TMS-EEG abnormalities in PD re flect structural connectivity changes in WM tracts, as measured by DTI. TEPs from motor cortical areas re flect cortical activation and propagation via reverberating corticobasal ganglia-thalamo-cortical pathways.  $14-16$  TMS likely activates monosynaptic projection to basal ganglia structures $46-48$  and reciprocal connections back to cortical areas through the thalamus, via polysynaptic pathways.[44,45,49](#page-9-0) Studies using deep brain stimulation coupled with EEG have shown that the stimulation of the subthalamic nucleus and internal globus pallidus generates EEG potentials, with onset latencies  $\sim$ 15 and 11 ms, respectively.  $44,45$  This suggests that activating cortico-subcortical-cortical loops by TMS could <span id="page-8-0"></span>contribute to TEPs as early as 20 ms. In our analysis, we concentrated on thalamic M1 connections, as the thalamus is the primary output of this loop and is believed to underlie M1 hypoactivation due to basal ganglia dysfunction.<sup>2,3</sup> For the facilitation of TEPs from pre-SMA in PD, our focus was on the connections between pre-SMA and anterior putamen, which is a major output of the pre-SMA and has been found to be functionally co-hyperactivated in PD. $30,31$  We found that PD patients exhibited reduced FA values, indicating microstructural alterations, in the tracts connecting the Vla thalamic nucleus to M1 and the pre-SMA to the anterior putamen, and in the control tract from the pulvinar to V1. Furthermore, PD patients exhibited increased RD in the Vla–M1 tract, suggesting differential patterns of WM pathology. These results are in line with previous studies showing widespread WM damage in early PD patients, involving both motor and nonmotor tracts,  $50,51$  indicating the presence of axonal abnormalities even in the early stage of the disease. The current study found that M1 and pre-SMA TEP alterations did not correlate with DTI abnormalities in PD patients. The lack of correlation suggests that changes in cortical excitability, as measured using TEPs, cannot strictly be attributed to alterations in structural connectivity along the WM tracts we investigated. TEP abnormalities in PD, therefore, may reflect intracortical changes or alternatively alterations in functional connectivity, within the cortico-subcortical circuits. However, the results of the present study do not exclude the possibility that TEP alterations might reflect structural connectivity damages in other pathways not examined in this study.

We acknowledge that our study has some limitations. We studied TEPs at rest and not during a finger-tapping task, which could have overlooked potential TEP modifications and correlations with bradykinesia. Despite the risk of misleading diagnosis in enrolling de novo PD patients, the diagnosis of PD was confirmed in subsequent clinical follow-up of these patients.

In conclusion, our study sheds light on mechanisms and clinical correlates of cortical excitability changes in PD. We found that M1 hypoexcitability is already present at the onset of the disease, whereas pre-SMA hyperexcitability is not. We propose that changes in M1 reflect a parkinsonian cortical state rather than changes correlating specifically with bradykinesia and speculate that pre-SMA alterations could develop at later disease stages. Future longitudinal studies will be useful to assess changes in M1 and pre-SMA excitability in PD during disease progression and chronic treatment.

### Data Availability Statement

Data are available from the corresponding author upon reasonable request.

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## Supporting Data

Additional Supporting Information may be found in the online version of this article at the publisher's web-site.

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### Author Roles

(1) Research project: A. Conception, B. Organization, C. Execution; (2) Statistical Analysis: A. Design, B. Execution, C. Review and Critique; (3) Manuscript Preparation: A. Writing of the First Draft, B. Review and Critique.

G.L.: 1A, 1B, 1C, 2A, 2B, 2C, 3A, 3B M.I.D.B.: 1A, 1B, 1C, 2A, 2B, 2C, 3A, 3B C.P.: 1C, 2A, 2B, 2C, 3B M.M.: 1C, 2B, 3B A.O.: 2B, 2C, 3B M.C.: 1C, 2B, 3B F.A.: 1C, 2B, 3B G.V.: 1A, 2C, 3B G.F.: 1A, 1B, 2C, 3B A.C.: 1A, 2A, 2C, 3B P.P.: 1A, 1B, 2A, 2C, 3B A.B.: 1A, 1B, 2A, 2C, 3B D.B.: 1A, 1B, 2A, 2C, 3B

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