



RESEARCH ARTICLE

White matter degeneration in remote brain areas of stroke patients with motor impairment due to basal ganglia lesions

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Funding information

Nanjing Health Science and Technology Development Special Fund Project, Grant/Award Number: YKK18218; National Natural Science Foundation of China, Grant/Award Numbers: 6590000127, 81801680

Abstract

Diffusion tensor imaging (DTI) studies have revealed distinct white matter (WM) characteristics of the brain following diseases. Beyond the lesion-symptom maps, stroke is characterized by extensive structural and functional alterations of brain areas remote to local lesions. Here, we further investigated the structural changes over a global level by using DTI data of 10 ischemic stroke patients showing motor impairment due to basal ganglia lesions and 11 healthy controls. DTI data were processed to obtain fractional anisotropy (FA) maps, and multivariate pattern analysis was used to explore brain regions that play an important role in classification based on FA maps. The WM structural network was constructed by the deterministic fiber-tracking approach. In comparison with the controls, the stroke patients showed FA reductions in the perilesional basal ganglia, brainstem, and bilateral frontal lobes. Using network-based statistics, we found a significant reduction in the WM sub-network in stroke patients. We identified the patterns of WM degeneration affecting brain areas remote to the lesions, revealing the abnormal organization of the structural network in stroke patients, which may be helpful in understanding of the neural mechanisms underlying hemiplegia.

KEYWORDS

diffusion tensor imaging, ischemic stroke, magnetic resonance imaging, motor impairment, white matter structural network

1 | BACKGROUND

Focal brain lesions can affect the overall performance of brain networks. Over the past decades, numerous neuroimaging studies using magnetic resonance imaging (MRI) have investigated the structural and functional reorganizations after stroke (Lim & Kang, 2015). Diffusion tensor imaging (DTI) is commonly used to evaluate the structural integrity of the white matter (WM). Previous DTI studies in stroke patients have calculated several diffusion tensor (DT) indicators

(e.g., fractional anisotropy [FA], mean diffusivity, radial diffusivity) in various regions along the corticospinal tract (CST) of the lesioned and contralateral hemispheres (Cunningham et al., 2015; Visser et al., 2019), or computed the CST integrity (Byblow, Stinear, Barber, Petoe, & Ackerley, 2015; Feng et al., 2015). These imaging indicators were subsequently correlated with functional outcomes, further establishing their value as predictive markers. Certain correlations have been identified between changes in specific fiber bundles and functional outcomes, such as aphasia (Meier, Johnson, Pan, & Kiran, 2019), neglect

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(Umarova et al., 2017), and paralysis (Jang et al., 2014). One study measured FA and axial diffusivity (AD) across 48 different WM tract regions in the brains of five hemiparetic patients. They found that patients with lesions involving the corona radiata (CR) and middle cerebral artery showed widespread reductions in perilesional FA and described longitudinal changes in the perilesional and remote FA and AD in relation to kinematic parameters of elbow flexion in the subacute poststroke period (Oey et al., 2019). This globally reduced FA after stroke may reflect extensive Wallerian degeneration (WD) of the descending pathways. Another study investigated global alterations of the lesion-spared network architecture in the acute and chronic stroke phases in patients showing spatial neglect. In addition to anterograde and retrograde axonal degeneration, structural network alterations can represent remote remodeling of fibers not directly connected to the lesion, that is, transneuronal degeneration. Moreover, the results showed that longitudinal WM changes, also transneuronal changes, may follow the persisting deficit (Umarova et al., 2017).

Unlike the CST command of motor functions, the involvement of other fibers and brain areas in movement control is not well understood. Voluntary movements are mostly controlled by the CST. The primary motor cortex is the source of most corticospinal axons, but its activity is strongly influenced by the pallidum, striatum, cerebellum, and many other cortical regions, including the somatosensory area of the cortex. Different cortical regions are involved in specific motor functions, such as motor learning, motor planning, motor preparation, and coordination. A complete action includes motor initiation and termination, feed-forward and feedback control loops, and feedback processing. Central processing systems also regulate movement through integration of sensory information with the motor plan. When the CST is damaged by large forebrain lesions in humans, the loss of fine movement is accompanied by hypertonia and hyperreflexia. These deficits do not appear in monkeys with lesions restricted to the CST, so they are probably attributable to damage to adjacent forebrain structures, such as the striatum, in humans (Desrochers, Brunfeldt, Sidiropoulos, & Kagerer, 2019; Watson, Kirkcaldie, & Paxinos, 2010). Therefore, neural degeneration in some regions may cause specific functional manifestations. We performed a pattern-recognition classification of FA maps of the brain in an attempt to explore regions that are closely related to motor function.

Multivariate pattern analysis (MVPA) is an imaging data analysis method based on machine learning and pattern recognition. It involves the use of pattern-classification algorithms to extract spatial patterns from neuroimaging data for the analysis of individual characteristics (Lao et al., 2004). MVPA offers the advantage of considering interregional correlations and searching for abnormalities throughout the entire brain because it adopts an unbiased and whole-brain method without artificially setting the region of interest (ROI) (Pereira, Mitchell, & Botvinick, 2009). This method has been widely used in psychiatry-related studies to analyze mild changes observable in sub-clinical populations (Janssen, Mourão-Miranda, & Schnack, 2018; Li et al., 2014).

WM structural connectivity can be modeled as a network. A network-based statistics (NBS) method can control the family-wise

error rate during mass univariate testing of every connection of the network. NBS can be also used to investigate the interregional correlations of the brain on a global level (Fortanier et al., 2019; Zalesky, Fornito, & Bullmore, 2010). We utilized these methods to explore subtle relevant changes and abnormalities of structural networks of the brain for a deeper understanding of the intrinsic brain structural basis of residual motor dysfunctions in ischemic stroke patients.

2 | METHODS

2.1 | Participants

Ten right-handed stroke patients (mean age, 56.7 ± 10.5 years) from the Southeast University-affiliated Zhongda Hospital were recruited for this study from March 2019 to December 2019. The inclusion criteria for patients were as follows: (a) age ≥ 20 and ≤ 80 years; (b) first onset of ischemic stroke with the involvement of the basal ganglia; (c) pure motor deficits; and (d) stable condition after treatment of acute stroke without recurrence. The exclusion factors were as follows: (a) a history of neurological or psychiatric disorders prior or subsequent to symptomatic stroke; (b) brain abnormalities unrelated to the infarct lesions; and (c) MRI contraindications. All of the affected extremities were evaluated for motor function. The motor outcome of the affected limbs was evaluated by the Fugl-Meyer assessment (FMA), including the upper and lower extremities (Feng et al., 2015). The Brunnstrom stage was also recorded. Recovery of the affected extremities was scored on a 6-point scale (1 = severe; 6 = normal) (Naghdi, Ansari, Mansouri, & Hasson, 2010). The clinical characteristics of the stroke patients are summarized in Table 2. Eleven demographically matched healthy control participants (mean age, 61.5 ± 7.8 years) were also recruited.

The study was approved by the local Ethics Committee of the Southeast University-affiliated Zhongda Hospital. All participants provided written informed consent to participate in accordance with the Declaration of Helsinki.

2.2 | Image acquisition

Diffusion-tensor images were acquired using a 3.0-Tesla Philips (Ingenia) Medical System equipped with a Synergy-L Sensitivity Encoding (SENSE) head coil and a single echo planar imaging sequence, and 33 diffusion-weighted images ($b = 1,000$ s/mm²) and a reference T2-weighted image with no diffusion weighting ($b = 0$ s/mm²) were obtained with the following acquisition parameters: voxel size = $2 \times 2 \times 2$ mm³, gap = 0 mm; echo time (TE) = 107 ms; repetition time (TR) = 5,835 ms; field of view (FOV) = 256×256 mm²; flip angle (FA) = 90°; matrix = 128×128 ; and slices = 75.

High-resolution T1-weighted axial images covering the whole brain were obtained by a 3D-magnetization prepared rapid gradient-echo (MP-RAGE) sequence with the following parameters: TR = 9.6 ms; TE = 3.7 ms; FA = 9°; matrix = 256×256 ; FOV = 256×256 mm²; voxel size = $1 \times 1 \times 1$ mm³; gap = 0 mm; and number of slices = 140.

Additionally, sagittal fluid attenuated inversion recovery (FLAIR) images were obtained with the following parameters: TE = 110 ms; TR = 7,000 ms; inversion time = 2,200 ms; FA = 90°; matrix size = 480 × 480; FOV = 250 × 250 mm²; slice thickness = 5 mm; number of slices = 20.

2.3 | Lesion mapping

Lesion-side normalization was performed to place lesions on the left side of the brain. For patients with lesions in the right hemisphere, the images were flipped from the right to the left along the midsagittal line to simplify the comparison with other patients with stroke. In accordance with previous investigations, lesion masks of each patient were manually segmented on individual structural MRI images (T1-weighted MP-RAGE and FLAIR images) using MRIcron software (<http://www.mricron.com>). After spatial normalization of all individual lesion masks, a lesion overlap image for all patients was constructed (Figure 1) (Chen & Schlaug, 2013; Grefkes et al., 2008; Zhang et al., 2016).

2.4 | DTI data preprocessing and network definition

DTI data analysis was performed by a pipeline toolbox for analyzing brain diffusion images (PANDA, <http://www.nitrc.org/projects/panda>) (Cui, Zhong, Xu, He, & Gong, 2013). The main procedure includes the following steps: (a) correction for head motion and eddy current effects using FMRIB's Diffusion Toolbox; (b) calculation of the DT metric, that is, FA, for each voxel by using the DTIFIT tool; and (c) normalization by registration of all the individual FA images to the FMRIB58_FA template by calling the FNIRT tool (Guo et al., 2019). The FA maps were obtained, and the FA values of 50 WM labels were computed according to the WM atlas "r1CBM_DTI_81_WMPM_FMRIB58.nii.gz." This atlas is created by hand-segmenting a standard-space average of diffusion

MRI tensor maps from 81 normal participants according to histology criteria and consists of 50 core regions (Mori et al., 2008).

Whole-brain tractography for each participant was performed in the native diffusion space based on fiber assignment by the continuous tracking algorithm (Mori, Crain, Chacko, & van Zijl, 1999). All voxels with FA values ≥ 0.2 were used as seed points; the FA and curvature thresholds of path tracing were set to 0.2 and 45°, respectively.

FA-weighted networks were constructed using 116 nodes defined according to the automated anatomical labeling (AAL) atlas (Tzourio-Mazoyer et al., 2002). The entire cerebral cortex was automatically partitioned into 116 anatomical ROIs (45 ROIs for each cerebral hemisphere, 26 ROIs for the cerebellum) using the AAL algorithm. The weight of the edges in the network was defined as the mean FA value of the connected fibers between each pair of nodes. In order to reduce false-positive connections, two nodes were considered structurally connected only when at least three fibers were reconstructed between them (Shu et al., 2011).

2.5 | Multivariate pattern analysis

Pattern classification analysis could be used to examine the differences in the FA values between groups. A specific MVPA approach known as support vector machine (SVM) was implemented using the Pattern Recognition for Neuroimaging toolbox (PRoNT) software (<http://www.mnl.cs.ucl.ac.uk/pronto/>) (Schrouff et al., 2013). Individual FA maps were treated as points located in a high-dimensional space. A linear decision boundary in this high-dimensional space was defined by a hyperplane that separated the individual brain images according to a class label (i.e., patients vs. controls) (Figure 2a). A more detailed description of the SVM can be found in previous reports (Li et al., 2014). The receiver operating characteristic curve (ROC), sensitivity, and specificity of the FA classifications and the weight of each brain region in the classification analysis were obtained. The brain regions with voxels showing values $\geq 30\%$ of the maximum weight vector value of the discrimination map were considered to be the key areas.

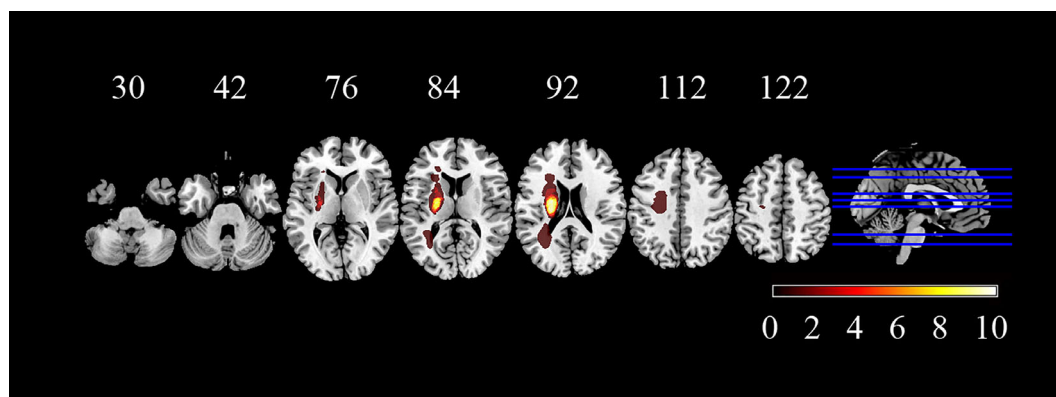


FIGURE 1 Lesion incidence map of patients with stroke. Stroke lesions were projected to the left hemisphere for each patient and overlaid onto a T1 template in MNI standard space. The color bar indicates the number of patients with stroke lesions in the corresponding voxel. The numbers above the brain images are Z values marking the MNI coordinates of the transverse sections

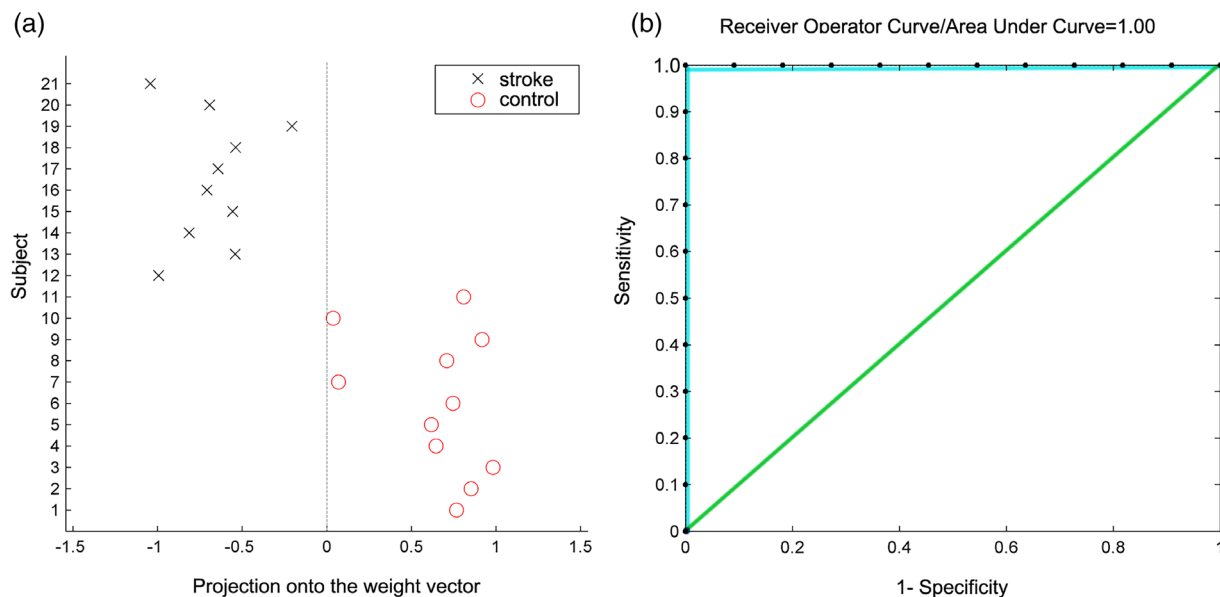


FIGURE 2 The results of multivariate pattern analysis (MVPA) classification. The classification plot (a) and receiver operating characteristic (ROC) curve (b) for the comparison between stroke patients and controls using fractional anisotropy (FA) maps derived from diffusion tensor imaging (DTI) data, which yielded an accuracy of 100% (100% sensitivity, 100% specificity), with statistical significance at $p < .005$

2.6 | Statistical analysis

Two-sample t tests were employed to test the group differences in demographic data and imaging measures, while the differences in FA values of 50 WM labels were compared between the two groups by using the t test. Correlations between FA values in the core regions and clinical scores were computed using Pearson's correlation test. These statistical tests were performed using the Statistical Package for Social Science (SPSS) 22.0 software. The two-sample t test was adopted for the comparisons of inter-nodal connections, followed by NBS using GRETNA (v2.0.0) (Wang et al., 2015) to analyze the FA networks between groups. In addition, the backbone extraction for the structural connectivity matrix (the subnetwork) of each group was calculated to show the group probability matrix. Visualization of the results was performed by the BrainNet Viewer (Xia, Wang, & He, 2013). The comparisons of the two sets of FA values in the key areas derived from MVPA were performed in SPM12 with false discovery rate (FDR) correction ($P_{FDR} < .05$, cluster size >5) (Liang, Cai, Zhou, Huang, & Zheng, 2020).

3 | RESULTS

3.1 | Demographic data and patient characteristics

The demographic data of the patients and the controls are presented in Table_1_SupplInfo. No significant differences in age or sex were observed between the two groups. The mean interval from stroke onset to DTI scans was 10.3 ± 9.0 weeks (Table 1). Stroke lesions were projected to the left hemisphere for each patient and overlaid onto a T1 template in MNI standard space (Figure 1).

WM labels with significantly different FA values between control and stroke groups are shown in Table 3. The WM labels that correlated with movement scores were evaluated with r values. Except for the FA of the fornix, which increased, all other FA values reduced in the stroke group. The most significant WM labels that showed correlation with FMA ($p < .001$) were the left posterior limb of internal capsule (PLIC, $r = .795$), superior CR ($r = .720$), and superior fronto-occipital fasciculus (SFOF, $r = .744$).

3.2 | Overall classifier performance

Figure 2a shows the results of the MVPA classification between the 10 stroke patients and the 11 controls based on FA values. The overall accuracy was 100%, and was significant at $p < .005$ ($p = .001$). Both the sensitivity and the specificity were 100% (with ROC shown in Figure 2b). The overall classification accuracy of the algorithm measures its ability to correctly sort the two groups.

3.3 | Discrimination map

In the whole-brain voxel weight maps, the weight vector value indicates the relative importance of the voxel in the decision function, that is, the discrimination between patients and controls (Figure 3). Note that all voxels in the WM mask contribute to the decision function since the analysis is multivariate. The spatial distribution of the weight vector provided information about the contribution of different areas to classification.

The brain regions that contributed the most to the discrimination between stroke patients and controls were identified by setting the

ID	Age (years)	Side	Localization of infarct	BRS	FMA	Scan time (week)
1	49	R	BG	3, 3, 5	58	22
2	67	R	BG	2, 1, 4	37	2
3	57	L	BG, PV	2, 1, 4	35	4
4	51	L	BG, PV	3, 2, 4	41	10
5	60	R	BG, CR	4, 4, 5	86	20
6	57	L	BG, CR	2, 1, 5	38	3
7	74	L	BG, CR	5, 4, 5	82	14
8	60	L	BG	2, 1, 3	19	2
9	35	L	BG	2, 1, 3	25	3
10	57	L	BG	5, 5, 5	89	24

TABLE 1 Patient characteristics

Note: Side: the hemisphere of lesions on brain; separate functional evaluation of proximal and distal portions of the upper and entire lower extremities; (full score = 100); scan time: interval of DTI acquisition from stroke onset.

Abbreviations: BG, basal ganglia; BRS, Brunnstrom stage; CR, corona radiata; DTI, diffusion tensor imaging; FMA, Fugl-Meyer assessment; IC, internal capsule; PV, periventricular.

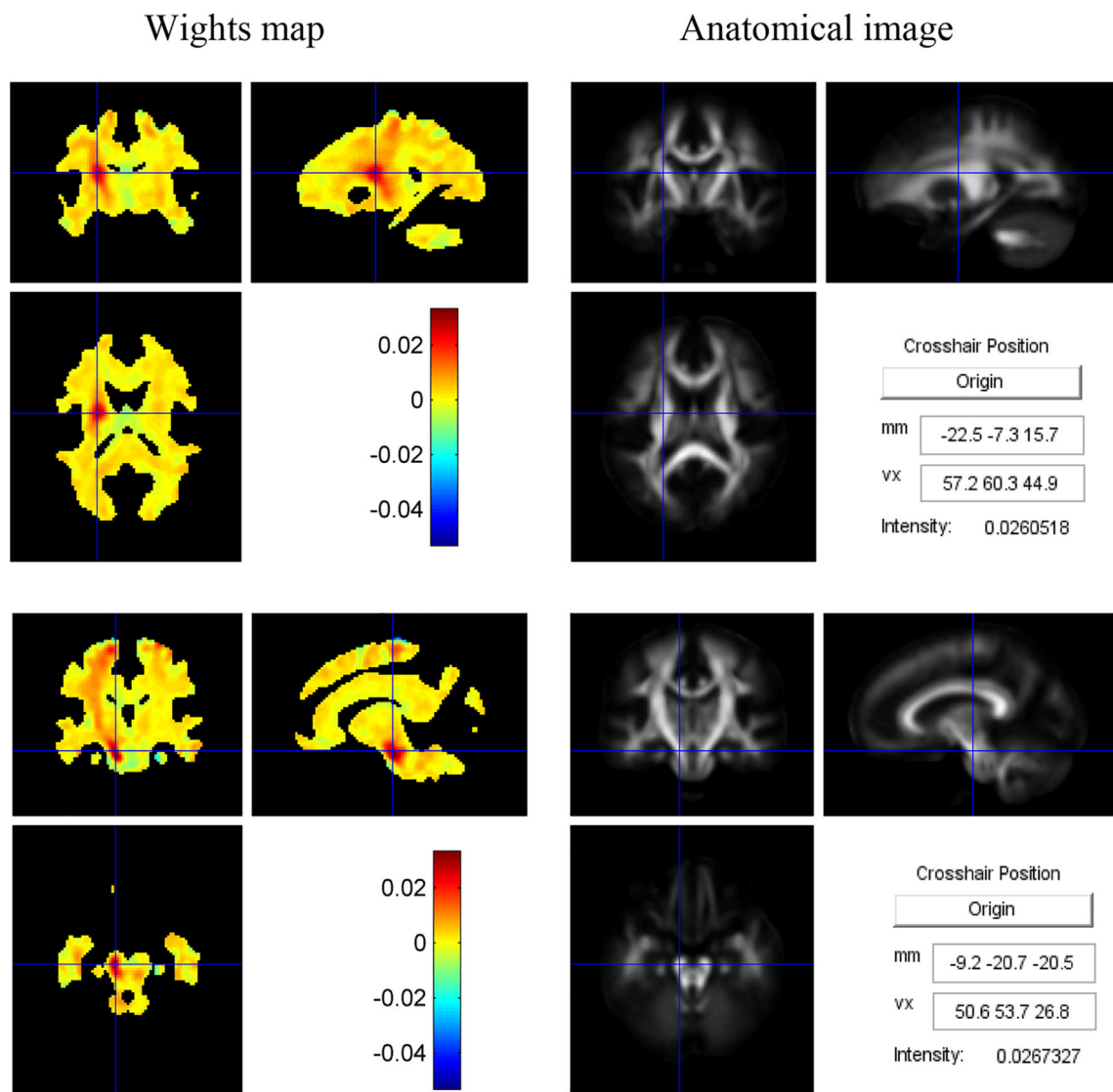


FIGURE 3 Whole-brain voxel weight map. It shows the white matter regions contributing to discrimination between groups based on fractional anisotropy (FA) values. The color bar indicates the weight vector value of the voxel, which is also indicated in the intensity field of the anatomical image (white fiber atlas “JHU-ICBM-FA-2 mm”) panel

threshold to $\geq 30\%$ of the maximum weight vector scores, consistent with previous studies using MVPA for disease classification (Ecker et al., 2010; Li et al., 2014). The most frequently identified classifying features of the FA maps included the perilesional basal ganglia and brainstem, with a few features appearing in the bilateral frontal lobes (Figures 3 and 4a), which were considered as the key areas.

By comparing the FA values of the two groups in the discrimination map by SPM12 ($P_{FDR} < .05$, cluster size > 5), we found that the stroke group showed reduced FA in the key areas, as shown in Figure 4a and reported in Table 2. According to the WM atlas, these clusters were mainly located in the CST pathway. Moreover, the FA values of these areas showed a positive correlation with that of the ipsilesional CST ($r = .888$, $p < .001$) (Figure 4b).

3.4 | Decreased connection of the component network in stroke patients

FA-weighted networks were constructed from the nodes (brain areas) defined according to the AAL atlas. The weight of the edges in the network was defined as the mean FA value of the connected fibers between each pair of nodes. The NBS approach was used on the structural networks constructed by deterministic tractography. We identified several significantly decreased connections of a component network (subnetwork) in stroke patients ($p < .001$, $p = .00099$) (Figure 5a and Table_2_SupplInfo).

Figure 5a presents the subnetwork that showed deterioration in the brain structure of the stroke patients. Figure 5b,c demonstrates

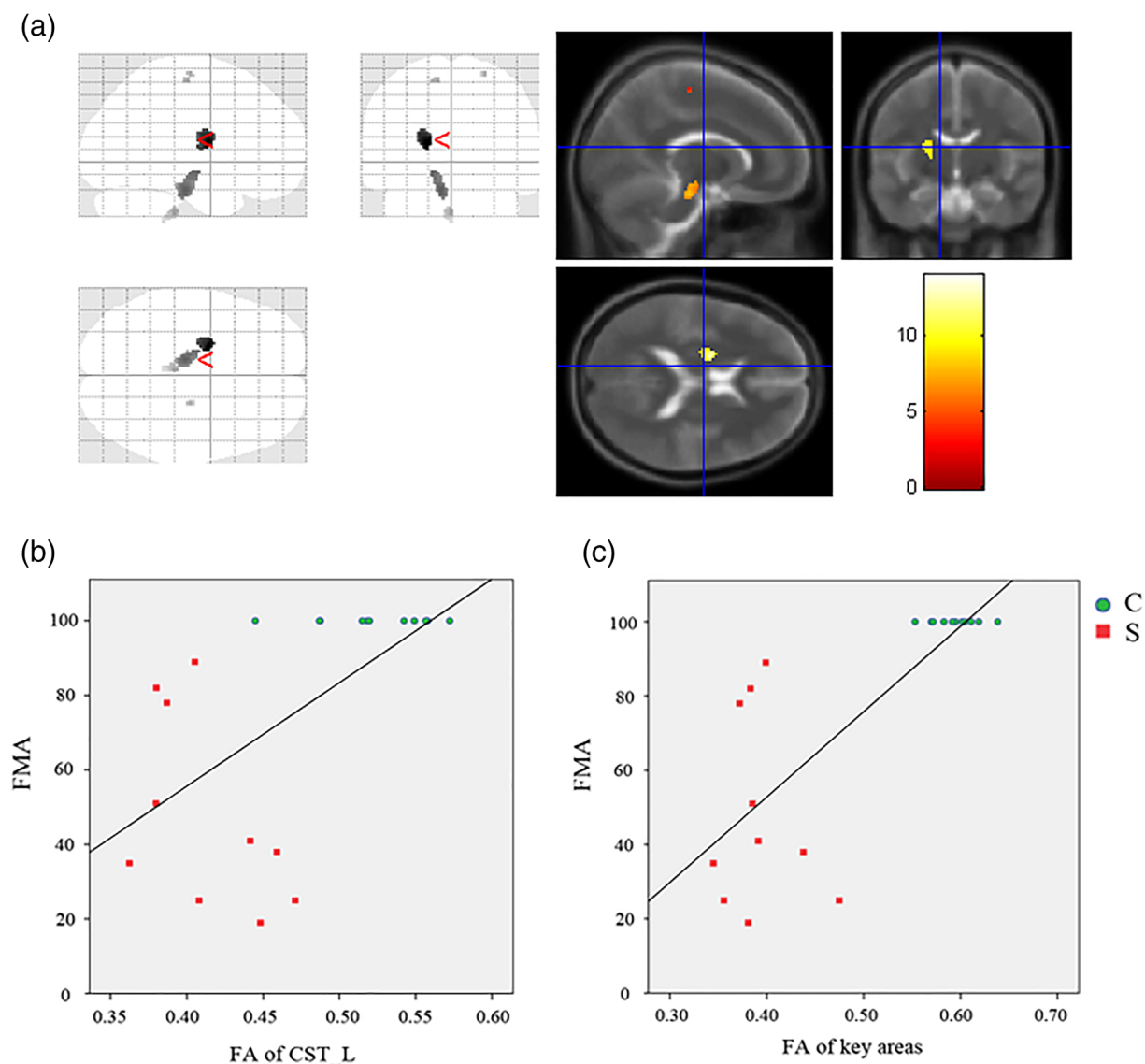


FIGURE 4 (a) The brain areas showing decreased fractional anisotropy (FA) in the patients compared to the controls, including the left brainstem, left basal ganglion and a fraction of voxels in the bilateral frontal lobe. These regions were identified by setting the threshold to $\geq 30\%$ of the maximum weight vector scores on the basis of the whole-brain voxel weight map. The color bar indicates the T value in two-sample t tests. (b,c) The FA values of the ipsilesional corticospinal tract (CST) and the key areas both showed positive correlations with Fugl-Meyer assessment (FMA) ($r = .588$, $p \leq .005$; $r = .784$, $p < .001$). C: control group; S: stroke group

TABLE 2 The brain areas showing decreased FA in patients in comparison with controls

ID	Voxel size	Peak MNI coordinate (x, y, z)	Peak intensity	Brain regions	White matter regions (voxel size)
1	36	-4, -36, -46	0.028	Pons; Medulla.L	CST.L 20
2	114	-10, -20, -22	0.032	Midbrain; Pons.L	CP.L CST.L 57 53
3	112	-22, -6, 16	0.027	Extra-nuclear; Lentiform Nucleus.L	PLIC.L SCR.L SFOF.L ALIC.L 43 30 15 12
4	9	-14, -20, 60	0.019	MFG.L	Not in the atlas
5	7	24, -18, 66	0.023	MFG.R	Not in the atlas

Note: ID: the index of the cluster; voxel size: number of voxels in the cluster; peak MNI coordinate: the location of the voxel with the maximum weight vector scores (also peak intensity) in each cluster. *Cluster Locator* in PANDA software was used to locate the cluster image according to *JHU ICBM-DTI-81 White-Matter Labels*. White matter atlas (voxel size): the atlas regions this cluster involves and the quantity of voxels in this cluster overlapped with each atlas region.

Abbreviations: ALIC, anterior limb of the internal capsule; CP, cerebral peduncle; FA, fractional anisotropy; L, left; MFG, medial frontal gyrus; PLIC, posterior limb of the internal capsule; R, right; SCR, superior corona radiata; SFOF, superior fronto-occipital fasciculus (could be a part of the anterior internal capsule).

the group probability matrix for the subnetwork of each group. The table in supplementary information reported the regions as nodes in the subnetwork. The subnetwork contained parts of the cerebral cortical regions (the frontal, parietal, and occipital lobes), the subcortical areas (the basal ganglion), and the cerebellum. Together with the results of MVPA, these findings showed obvious changes involving the frontal parietal lobe and the basal ganglia, and those in the brainstem region may overlap with some of the fibers connected to the cerebellum.

4 | DISCUSSION

Beyond the well-known concept of lesion-symptom mapping, some lesions in a single location in the brain could disrupt brain functions routed to widespread neural networks (Burke Quinlan et al., 2015; Lim et al., 2014). Our analysis confirmed that local destruction of basal ganglia could affect remote areas in the brain.

4.1 | WM degeneration in the CST pathway of stroke patients

The degree of anisotropy depends on the level of organization, the integrity of the WM tract, and the degree of freedom for water diffusion caused by the oriented axonal membranes and myelin sheaths (Virta, Barnett, & Pierpaoli, 1999). Reduced anisotropy along the CST far from the original lesions has been interpreted as WD (Thomalla et al., 2004). DTI can quantify the FA values to evaluate pathology changes in the WM, such as WD. Using MVPA, we reported that apart from the basal ganglia region where the infarcts are localized, brain areas with significantly decreased FA values were also located in the brainstem of the lesioned hemisphere, and a few areas were present in the bilateral frontal lobes, which might be indicative of the degenerative lesions caused by WD.

The acute and chronic phases of stroke probably differ in terms of WM changes since neural changes can include anterograde and retrograde degeneration, or refactoring. However, regardless of the type of alteration, it should generate specific structural changes and affect the corresponding functions. Therefore, we performed pattern-recognition classification using the whole-brain FA map and explored the key brain regions important for distinguishing stroke patients from controls. MVPA analysis explored the key brain regions important for distinguishing stroke patients from controls. Most of the regions were located in the WM based on the atlas, but some voxels were still not in the WM areas. They may have been present at the junction of gray and WM. Therefore, we can investigate the areas that were embedded in a network by NBS. In comparison with the whole brain network, we referred to the decreased FA-weighted component network as a subnetwork. Its contents are displayed in Figure 5, and include the frontal lobe, limbic lobe, occipital lobe, parietal lobe, basal ganglia, temporal lobe, and cerebellum. We thought that this subnetwork may be specific to basal ganglia stroke and could be generalizable for patients with hemiplegia.

Our study verified the degenerative changes in the WM of stroke patients. The infarcted lesions were mainly located in the basal ganglia region, but the FA reduction in some remote areas had reached the point where they could be differentiated from the controls. Thus, the damaged structural anatomy of the subcortical areas may induce deterioration of key WM areas in the brain.

4.2 | Decreased WM connections were widely distributed across the brain regions of stroke patients

Many patients showed motor dysfunctions after the occurrence of cerebral infarction. These dysfunctions were usually related to injury of the CST. The CST originates from multiple motor and somatosensory cortices, including the premotor cortex, supplementary motor cortex (SMA),

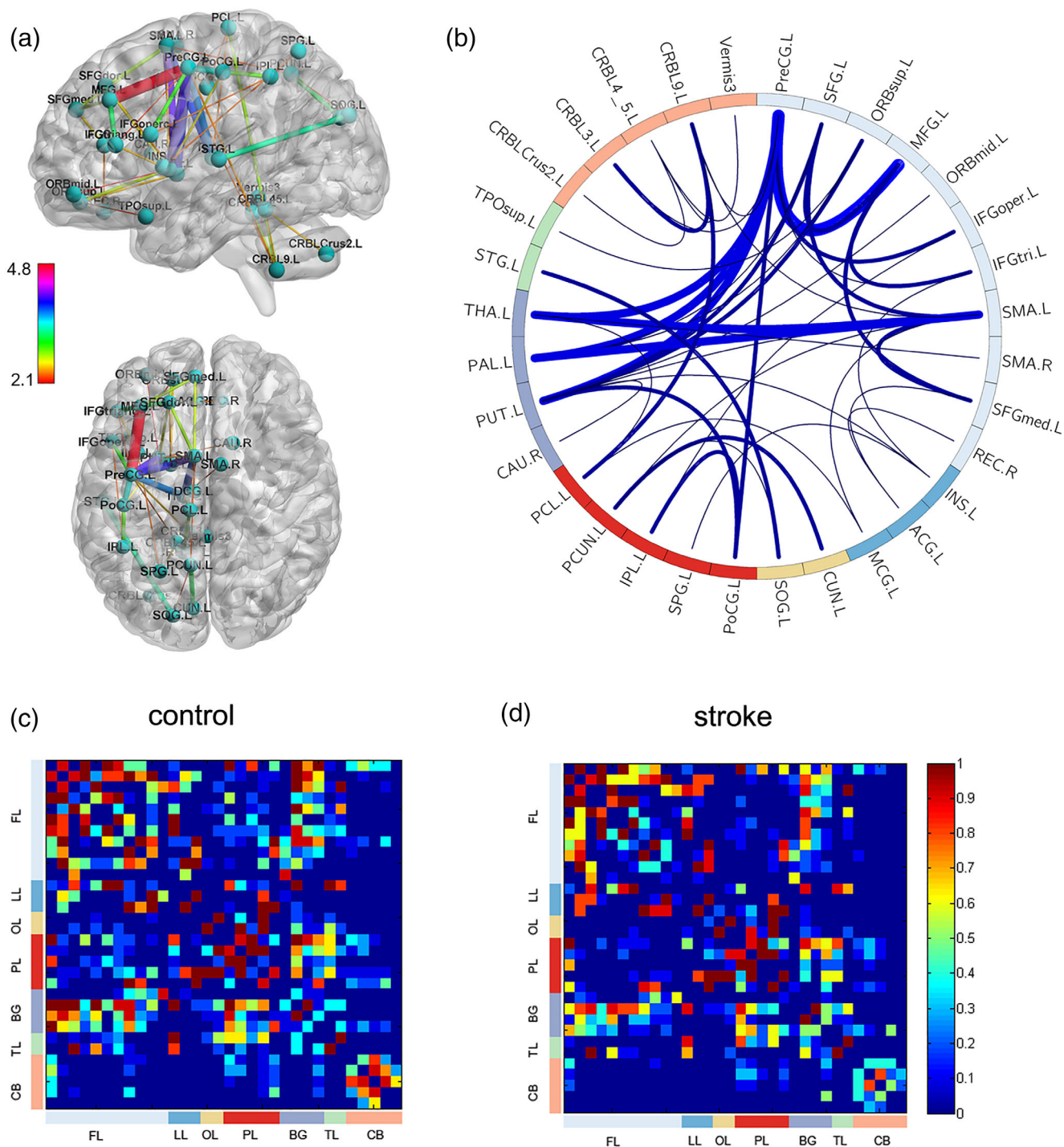


FIGURE 5 The subnetwork identified by the network-based statistics (NBS) analysis. (a,b) The subnetwork demonstrated reduced connectivity in stroke patients in comparison with controls. The connections in the subnetwork shown in a brain model (a) and a circle (b). The color bar represents the *t* value derived from two-sample *t* test for each connection and the thickness of the edges represents how significantly the two groups are different (thickness of lines in b: $t < 2.5$, thin; $2.5 \leq t < 3.5$, moderate; $t \geq 3.5$, thick). (c,d) The group averaged fractional anisotropy (FA)-weighted structural connectivity network for the control and stroke groups. The color bar indicates the connection probability in the groups. The abbreviations in (b) are illustrated in the Supporting Information. BG, basal ganglia; CB, cerebellum; FL, frontal lobe; LL, limbic lobe; OL, occipital lobe; PL, parietal lobe; TL, temporal lobe

primary motor cortex, as well as primary and secondary somatosensory cortices. The CST is crucial for proper execution of a volitional movement (Chenot et al., 2019; Lemon, 2008). Apart from the CST and the

motor areas of the cortex, the proper execution of movements involving balance and coordination also requires the extrapyramidal tract and other brain regions such as basal ganglia and the cerebellum (Moreno-López,

Olivares-Moreno, Cordero-Erausquin, & Rojas-Piloni, 2016). The NBS analysis showed that the structural subnetwork connection of the stroke group was weaker than that of the control group, indicating that the nerve fibers involved in the subnetwork were affected. Destruction of the integrity and order of the brain structure may be reflected in the WM. Not only was the motor cortex directly related to motor commands, but other regions that regulated movement were also involved in the subnetwork.

In addition, since the MVPA analysis is multivariate, all voxels in the WM mask contribute to the decision function in the processing stage. The CST passes through the frontal lobe and midbrain, and MVPA verified its pathway degeneration in stroke patients with hemiplegia. Other brain areas influenced were explored by NBS (e.g., the occipital lobe, parietal lobe, and cerebellum). The results of MVPA more obviously showed the damage to the pathway from the basal ganglia to the brain stem (Figure 4). The results of NBS showed that the affected brain tissues were distributed across a wide range of brain regions, and the damage to the pathway from the basal ganglia to the cortex was more obvious (Figure 5). Both methods showed reduced FA in the frontal lobe and basal ganglia. The connections between the frontal lobe and the cerebellum pass through the cerebral peduncle (CP) in the brainstem. Therefore, the weakening of these connections in subnetworks derived from NBS was partially consistent with the decreased FA in the brainstem based on MVPA. The affected contralesional areas were verified with the difference of the corpus callosum and fornix between groups. In short, the decreased FA appears to spread out from the original infarct area as time elapses after the acute stage of the stroke onset. The subnetwork specific to basal ganglia stroke implicated the involvement of an internal model in patients with hemiplegia. Next, we briefly discuss the brain regions involved in the subnetwork.

4.3 | The bilateral frontal lobe

Similar to the results of previous studies showing infarct-related focal thinning of the motor area in the remote cortex via degeneration of inter-hemispheric connection fibers of the corpus callosum (Duering et al., 2015; Hayward et al., 2017), we found changes in the connection between the frontal hemispheres, as well as reduced FA values in a small area located in the contralesional frontal cortex. In studies calculating the mean kurtosis values of manually drawn ROIs from diffusion kurtosis imaging, secondary degeneration has been reported to occur in the ipsilesional precentral gyrus (PreCG) at the 6-month follow-up after subcortical stroke involving the CST (Wei, Shang, Zhou, Zhou, & Li, 2019).

In Figure 5, the connection between the left PreCG and middle frontal gyrus (MFG) in the subnetwork derived from NBS was reduced in the stroke group, which may well be linked to reduced FA in the frontal lobe in Figure 4. This cluster was not present in the WM atlas, but was present in the MFG (Table 2). The CST originates from motor cortices, including the PreCG, premotor cortex, and SMA in MFG. In Table 3, FA values of the superior CR, anterior CR, superior

TABLE 3 WM labels with different FA values between groups

ID	WM label	t test	Correlation test	
		p	p	r
Increased FA in the stroke group				
1	Fornix	.010	.024	-.490
Decreased FA in the stroke group				
2	Splenium of the corpus callosum	.035	.088	
3	CST.L	.000	.005	.588
4	Superior CP.R	.001	.032	.468
5	Superior CP.L	.026	.214	
6	CP.L	.000	.006	.579
7	ALIC.R	.014	.048	.436
8	ALIC.L	.003	.118	
9	PLIC.R	.040	.065	
10	PLIC.L	.000	.000	.795
11	RIC.R	.018	.086	
12	RIC.L	.001	.003	.610
13	Anterior CR.L	.039	.309	
14	Superior CR.R	.007	.064	
15	Superior CR.L	.000	.000	.720
16	Posterior CR.R	.005	.062	
17	Posterior CR.L	.003	.038	.456
18	Sagittal stratum.R	.001	.006	.582
19	Sagittal stratum.L	.001	.033	.466
20	External capsule.L	.001	.037	.457
21	SLF.R	.004	.111	
22	SLF.L	.007	.110	
23	SFOF.L	.000	.000	.744
24	Uncinate fasciculus.R	.008	.179	

Note: WM labels with significantly different FA values between the control and stroke groups using the t test ($p < .05$). The correlation between FA values and FMA scores was determined using Pearson correlation test. The correlation coefficient r is shown when $p < .05$. The analyses included the control group which showed the full FMA score (FMA = 100).

Abbreviations: ALIC, anterior limb of the internal capsule; CP, cerebral peduncle; CR, corona radiata; FA, fractional anisotropy; FMA, Fugl-Meyer assessment; L, left; PLIC, posterior limb of the internal capsule; R, right. Sagittal stratum (includes the inferior longitudinal fasciculus and inferior fronto-occipital fasciculus); RIC, retrolenticular part of the internal capsule; SFOF, superior fronto-occipital fasciculus (could be a part of the anterior internal capsule); SLF, superior longitudinal fasciculus; WM, white matter.

longitudinal fasciculus, and parts of the internal capsule were shown to be decreased in the stroke group. We speculated that the FA changes in the CST pathway and that these changes may be linked with the related motor supplementary areas in frontal lobes.

4.4 | The basal ganglia region

The patients recruited in this study all had infarcts localized to the basal ganglia. The comparison results for the WM regions in Table 3

showed that the PLIC, anterior limb of the internal capsule, retrolenticular part of internal capsule, and external capsule were affected after stroke. These WM tracts were located between the thalamus and basal ganglia. The advantage of the structural subnetwork derived from NBS was that it showed connections between smaller brain areas, such as the edges between the thalamus, putamen, pallidum, PreCG, SMA, and PoCG (Figure 5b). The CST section passing through the internal capsule may receive regulatory information from the nucleus of basal ganglia. Due to the cortico-basal ganglia-thalamocortical “motor” loop, any impact on the circuit constituents can lead to a shift in the balance between neural interactions in the direct and indirect pathways and subsequently lead to variations in the brain functions (Alexander, Crutcher, & DeLong, 1991; Silkis, 2001). In the early stage of rehabilitation, stroke patients with hemiplegia often have synergistic movements, which are thought to be related to this loop.

4.5 | The parietal and occipital lobes

Fibers connecting the frontal, parietal, and occipital lobes were affected after stroke (Figure 5), and the SFOF showed obvious degeneration in the stroke group (Table 3). The parietal lobe participates in sensory and motor integration, while the occipital lobe is related to visual effects. Stroke patients with dysfunction of normal voluntary movements may develop corresponding abnormal sensory modulations that gradually affect brain structure (Buaron, Reznik, Gilron, & Mukamel, 2020).

4.6 | The cerebellum

The cerebellum is often related to balance adjustment, and patients with hemiplegia usually experience problems with stability and coordination after stroke. The AAL atlas did not contain regions in the brainstem. Therefore, connections in the network could not precisely demonstrate fibers to the brainstem, including the red nucleus and substantia nigra. In our study, the CP was changed after stroke, and the reduced FA of the brainstem was probably related to changes in cortico-ponto-cerebellar tract. Previous DTI studies also showed decreased FA in the midbrain of stroke patients by manually plotting ROIs (Wei et al., 2019). A recent study indicated that the cerebellum plays a role in residual motor output by facilitating cortical excitability in chronic stroke (Guder et al., 2020). Taken together, these findings suggest that the inability to perform normal movements might gradually lead to abnormal balance, which is reflected in decreased cortico-cerebellar connectivity in these patients.

5 | LIMITATIONS

This study had multiple limitations. First, the sample size was small. Second, the interval of DTI acquisition from stroke onset ranged from 2 to 24 weeks. Recovery of FA in the penumbra regions occurs most

rapidly during the first 2 weeks following stroke, with continued slow increases in FA occurring for many weeks thereafter (Ding et al., 2008; Mandeville, Ayata, Zheng, & Mandeville, 2017). The phases of rehabilitation may influence WM organization, but we only discussed stroke patients with motor impairments in a cross-sectional manner. Future studies should aim to further explore these findings by expanding the sample size and dynamically observing changes from the acute to chronic phases. Additionally, emphasis should be placed on the classification and refinement of clinical behaviors of stroke patients, with identification of the specific brain feature changes that correspond to the functional outcomes.

6 | CONCLUSIONS

Our study recruited stroke patients with motor dysfunction and used MVPA and NBS methods based on DTI data to detect reduced FA values and abnormal WM connections at a global level in the brain of these patients. We found multiple WM structural abnormalities in the affected brain areas of stroke patients showing motor impairment. Our study may provide the basis for further exploration of the neural mechanisms involved in residual motor deficits in stroke patients. In future studies, we will further compare the neural changes in well-recovered patients to provide a basis for the development of adaptive rehabilitation training strategies.

ACKNOWLEDGMENTS

The authors thank all the patients and volunteers for participating in this study. This work was funded by the National Natural Science Foundation of China (grant Nos. 6590000127 and 81801680) and Nanjing Health Science and Technology Development Special Fund Project (grant No. YKK18218).

CONFLICT OF INTEREST

The authors declare no conflict of interests.

DATA AVAILABILITY STATEMENT

The data in the current study are available on reasonable request to the corresponding author.

PATIENT CONSENT STATEMENT

All participants gave written informed consent to participate in accordance with the Declaration of Helsinki.

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How to cite this article: Cao, X., Wang, Z., Chen, X., Liu, Y., Wang, W., Abdoulaye, I. A., Ju, S., Yang, X., Wang, Y., & Guo, Y. (2021). White matter degeneration in remote brain areas of stroke patients with motor impairment due to basal ganglia lesions. *Human Brain Mapping, 42*(14), 4750–4761. <https://doi.org/10.1002/hbm.25583>