## Journal of the American Heart Association

## **ORIGINAL RESEARCH**

# Imaging of Angiogenesis in White Matter Hyperintensities

**BACKGROUND:** White matter hyperintensities (WMHs) are areas of increased signal intensity on T2-weighted magnetic resonance imaging (MRI). WMH penumbra may be a potential target for early intervention in WMHs. We explored the relationship between angiogenesis and WMH penumbra in patients with WMHs.

METHODS AND RESULTS: Twenty-one patients with confluent WMHs of Fazekas grade  $\geq$ 2 were included. All the participants underwent <sup>68</sup>Ga-NOTA-PRGD2 positron emission tomography/magnetic resonance imaging. WMH penumbra was analyzed with masks created for the WMH and 7 normal-appearing white matter layers; each layer was dilated away from the WMH by 2 mm. Angiogenesis array and ELISA were used to detect the serum levels of angiogenic factors, inflammatory factors, HIF-1 alpha, and S100B. Fourteen patients with increased <sup>68</sup>Ga-NOTA-PRGD2 maximum standardized uptake (>0.17) were classified into group 2. Seven patients with maximum standardized uptake  $\leq$ 0.17 were classified as group 1. WMH volume and serum levels of integrin ανβ3, vascular endothelial growth factor receptor 22, and interleukin-1β tended to be higher in group 2 than in group 1. In group 2, <sup>68</sup>Ga-NOTA-PRGD2 uptake was significantly increased at the border between the WMH and normal-appearing white matter than in WMHs (P=0.004). The structure penumbra, defined by fractional anisotropy, was wider in group 2 (8 mm) than in group 1 (2 mm). The cerebral blood flow penumbra was 12 mm in both groups. Angiogenesis showed a correlation with reduced cerebral blood flow and microstructure integrity.

**CONCLUSIONS:** Our study provides evidence that angiogenesis occurs in the WMH penumbra. Further studies are warranted to verify the effect of angiogenesis on WMH growth.

Key Words: angiogenesis ■ cerebral blood flow ■ PET/MRI ■ white matter hyperintensities

hite matter hyperintensities (WMHs) are part of the spectrum of cerebral small-vessel disease markers, which are considered age-specific changes in magnetic resonance imaging (MRI) and extend over time. In recent years, an increasing number of studies have observed the impact of WMH progression on predicting a more rapid decline in global cognitive performance. Sabayan et al found an association between accelerated progression of WMHs and a high risk of all-cause death (hazard ratio, 1.22 [95% CI, 1.09–1.37] per mL/y increase in WMHs). Therefore, preventing or slowing the progression of

white matter injury has emerged as a major goal of current research.

The term white matter hyperintensity penumbra has been used to characterize the normal-appearing white matter (NAWM) tissue surrounding the WMHs and may represent subtle white matter injuries associated with the progression of WMHs. Previous diffusion tensor imaging (DTI) studies have shown that structural WMH penumbra, as measured by fractional anisotropy (FA), extends  $\approx$ 2 to 9 mm distal to WMH, whereas the cerebral blood flow (CBF) WMH penumbrae are  $\approx$ 11 to 14 mm distal to WMH. The WMH penumbra has

Correspondence to: Bin Peng, MD, Department of Neurology, Peking Union Medical College Hospital, Chinese Academy of Medical Science and Peking Union Medical College, No. 1, Shuaifuyuan, Dongdan, Dongcheng District, Beijing 100730, China. Email: <a href="mailto:pengbin3@hotmail.com">pengbin3@hotmail.com</a>

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#### **CLINICAL PERSPECTIVE**

#### What Is New?

- 68Ga-NOTA-PRGD2 positron emission tomography can be used to detect the expression of integrin ανβ3, a molecular biomarker for angiogenesis.
- Our study provides evidence that angiogenesis exists in white matter hyperintensity penumbra.

#### What Are the Clinical Implications?

- Angiogenesis is correlated with reduced cerebral blood flow and microstructural integrity.
- This study provides new insights into angiogenesis as a novel target for therapeutic intervention to arrest the progression of white matter hyperintensities.

#### **Nonstandard Abbreviations and Acronyms**

CBF cerebral blood flow
DTI diffusion tensor imaging
FA fractional anisotropy

**FOV** field of view **MD** mean diffusivity

MMP matrix metalloproteinase
 NAWM normal-appearing white matter
 SUV<sub>max</sub> maximum standardized uptake values
 SUV, standardized uptake value ratios

TE echo time
TR repetition time

**WMHs** white matter hyperintensities

an increased likelihood of conversion to WMHs.<sup>5-8</sup> Therefore, we hypothesized that the WMH penumbra is likely to become a potential target for early intervention in WMHs.

WMH remains a clinical challenge due to its poorly understood pathogenesis. WMHs are thought to have an ischemic pathogenesis caused by chronic hypoperfusion due to localization in the watershed areas formed by the terminal arteries. Photogenesis are due to CBF in the WMH penumbra has been associated with the development of WMHs. Potential interventions to increase white matter blood flow for the prevention of further white matter damage and their clinical consequences have been discussed. Collateral revascularization in the penumbra areas has been correlated with recovery and longer survival times in patients with ischemic stroke. Page 12,13 Angiogenesis is a form of collateral circulation and a crucial restorative mechanism under

ischemic conditions. Recently, several studies have demonstrated a relationship between angiogenic factors and WMHs.<sup>14,15</sup>

Whether angiogenesis exists and plays a role in the WMH penumbra is unclear. There are still challenges in the study of angiogenesis in vivo. However, because positron emission tomography (PET) imaging is widely used in cerebrovascular studies, previous PET studies have shown that arginine–glycine–aspartic acid uptake is significantly correlated with integrin  $\alpha\nu\beta3$  expression and microvessel density. The integrin  $\alpha\nu\beta3$ -targeted arginine–glycine–aspartic acid sequence-based probe showed high specificity for angiogenesis imaging.  $^{16-18}$   $^{68}$ Ga-NOTA-PRGD2 PET has been used to evaluate angiogenic activity in cerebral ischemic diseases and cancer.  $^{19,20}$ 

In this study, we hypothesized that angiogenesis in the WMH penumbra may be correlated with reduced CBF and white matter integrity. We tested this hypothesis by detecting integrin  $\alpha\nu\beta3$  expression using simultaneous <sup>68</sup>Ga-NOTA-PRGD2 PET/MRI.

#### **METHODS**

#### **Data Availability**

The data that support the findings of this study are available from the corresponding author upon reasonable request.

#### Study Design and Participants

Twenty-one participants with cerebral small-vessel disease from Peking Union Medical College Hospital between November 2016 and February 2018 were included. The inclusion criterion was confluent WMHs of Fazekas grade ≥2 on MRI. The patients with any cause of WMHs other than small-vessel disease, tumors, systemic inflammatory disease, brain trauma, and contraindications to PET/MRI were excluded. This study was approved by the institutional review board of Peking Union Medical College Hospital. Written informed consent was obtained from all included patients or their representatives.

#### PET/MRI Protocol

All images were acquired using an integrated time-of-flight (TOF) PET/MRI scanner (SIGNA PET/MRI; GE Healthcare, Waukesha, WI). <sup>68</sup>Ga-NOTA-PRGD2 was synthesized as described previously.<sup>21</sup> A dynamic PET scan was performed for 60 minutes after an intravenous injection of <sup>68</sup>Ga-NOTA-PRGD2, at a dose of 1.85 MBq (0.05 mCi) per kilogram of body weight. The interval between injection and scanning was about 30 to 60 minutes. PET and 3T MRI data were simultaneously acquired. The magnetic resonance

sequences were performed as below: T2-weighted imaging with fast recovery fast spin echo (repetition time [TR]=4691 ms, echo time [TE]=89 ms, field of view [FOV]=22×22 cm, slice thickness=3 mm, slice spacing=0 mm, and matrix=320×320); T1-weighted imaging with 3-dimensional fast spoiled gradient echo (TR=7.4 ms, TE=3.2 ms, preparation time=4.00 ms, number of excitations=1.00, FOV=25.6 cm×23.2 cm, slice thickness=1 mm, slice spacing=0.6 mm, and matrix=256×232); 3-dimensional pseudocontinuous arterial spin labeling (TR=4790 ms, TE=10.2 ms, post label delay (PLD) =)=2025 ms, number of excitations=5, FOV=24 cm×24 cm, slice thickness=5 mm, slice spacing=0 mm, and matrix=320 × 224); fluid attenuated inversion recovery (TE= =130 ms, TR=12000 ms, time of inversion =) =2711 ms, number of excitations= =2.00, FOV= =21 cm×18.9 cm, slice thickness=3 mm, slice spacing=0 mm, and matrix=256 × 192); and DTI sequence (TR=10999 ms, TE=72.9 ms, slice thickness=2.5 mm, FOV=24 cm, and 15 diffusion-weighted directions with b=1000 s/mm<sup>2</sup>).

#### **Imaging Analysis**

To calculate the uptake of <sup>68</sup>Ga-NOTA-PRGD2 more precisely, PET images were reconstructed using the time-of-flight method and postprocessed using the GE Advantage Workstation version 4.6 (GE Healthcare). In this study, images were reconstructed using the ordered subset expectation maximization reconstruction method (32 subsets, 8 iterations, filter cutoff frequency of 1.5 mm) combined with high definition (Sharp IR). The full width at half maximum of the Gaussian time-of-flight kernel with a width of 400 ps was used in the reconstruction. Magnetic resonance attenuation correction was performed on the basis of the double-point Dixon sequence.<sup>22</sup>

PET/MRI fusion images were visually evaluated for focal  $^{68}\text{Ga-NOTA-PRGD2}$  uptake by 2 experts. The maximum standardized uptake values (SUV\_max) were obtained for  $^{68}\text{Ga-NOTA-PRGD2}$  PET images. The participants were divided into 2 groups according to  $^{68}\text{Ga-NOTA-PRGD2}$  SUV\_max in the WMH, NAWM, or border between the WMH and NAWM.

# Tissue Segmentation and Spatial Normalization

CBF maps of arterial spin labeling were generated on a GE AW 4.5 workstation using a software 3-dimensional ASL Functool kit. The model used for calculating CBF (mL/min per 100 g) was based on a previous study.<sup>23</sup>

DTI images were postprocessed using the FMRIB Software Library (FSL version 5.0; http://www.fmrib.ox.ac.uk/fsl). We calculated DTI parameters including FA and mean diffusivity (MD). To further investigate the correlation between <sup>68</sup>Ga-NOTA-PRGD2 uptake and

changes in CBF (mL/min per 100 g tissue) and white matter structural integrity, all PET-MRI images were preprocessed using SPM12 (statistical parametric mapping; https://www.fil.ion.ucl.ac.uk/spm/software/spm12/). Each image was checked to ensure common orientation.

Fluid attenuated inversion recovery and 3-dimensional T1-weighted images were aligned using affine coregistration for each individual. Three-dimensional T1 images were used to segment brain tissue into cortical gray matter, white matter, and cerebrospinal fluid. WMH lesions were obtained from the 3-dimensional T1- and T2-weighted fluid attenuated inversion recovery images using the Lesion Segmentation Tool in SPM12. Manual refinement was performed by an expert to ensure an accurate representation of WMH using ITK-SNAP (www.itksnap.org). NAWM masks were created for each individual to determine the WMH penumbra for each measure. Each layer was dilated away from the WMH by 2mm for a total of 7 NAWM layers. We then applied the WMH and NAWM L1 to L7 masks to the PET, CBF, FA, and MD maps, which were previously linearly aligned to their T1-weighted images (Figure 1). The mean CBF, FA, MD, and <sup>68</sup>Ga-NOTA-PRGD2 uptake for the WMH and each NAWM layer were measured; the surrounding brain tissue regions were excluded through the brain mask. Standardized uptake value ratios (SUV<sub>r</sub>) were calculated for each NAWM layer, divided by the respective WMH.

#### **Serological Test**

Serum samples were collected from all participants and stored at  $-80\,^{\circ}$ C. The angiogenesis array (RayBiotech, Norcross, GA) was used to detect the serum concentrations of angiogenic factors, including E-selectin, matrix metalloproteinase (MMP)-2, MMP-9, vascular endothelial growth factor (VEGF) receptor 2, VEGF receptor 1, and VEGF-A. An ELISA was used to detect serum concentrations of integrin  $\alpha v \beta 3$  (LifeSpan BioSciences), S100B (LifeSpan BioSciences), and hypoxia-inducible factor 1-alpha (HIF-1 alpha) (RayBiotech).

#### **Statistical Analysis**

Baseline characteristics with normal distribution are reported as mean±SD. Nonnormally distributed variables are presented as medians (interquartile range). Independent-sample t tests or the Mann–Whitney U test were used for continuous variables, and chisquare tests were used for categorical variables to compare the differences between the 2 groups. We compared  $^{68}$ Ga-NOTA-PRGD2 PET with SUV<sub>r</sub>, CBF, FA, and MD in WMH and NAWM L1 to L7. The values from NAWM L7 were used as references. Structural penumbra was defined as the NAWM region with a reduced FA or MD. CBF penumbra was defined as

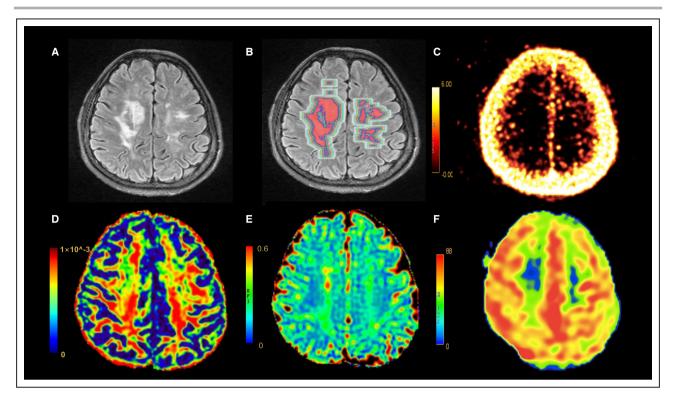


Figure 1. 68Ga-NOTA-PRGD2 PET/MRI.

**A**, FLAIR sequence showed WMHs at centrum semiovale. **B**, WMH and NAWM masks for each individual were created. Each NAWM layer was dilated away from the WMHs by 2 mm for a total of 7 NAWM layers. **C**, <sup>68</sup>Ga-NOTA-PRGD2 PET. **D**, The FA map. **E**, The MD map. **F**, The CBF map. CBF indicates cerebral blood flow; FA, fractional anisotropy; FLAIR, fluid attenuated inversion recovery; MD, mean diffusivity; MRI, magnetic resonance imaging; NAWM, normal-appearing white matter; PET, positron emission tomography; and WMHs, white matter hyperintensities.

the NAWM region with a reduced CBF.<sup>8</sup> The relationships among CBF, FA, MD, and SUV<sub>r</sub> were assessed using Spearman's correlation analysis. The serum concentrations of angiogenic factors were compared between the 2 groups. Bonferroni correction was used to perform a correction for multiple comparisons. Standardized mean difference (SMD) and 95% CI were calculated. Analyses were performed using the Statistical Package for the Social Sciences version 25.0 software (SPSS; IBM, Armonk, NY). Statistical significance was set at *P*<0.05.

#### **RESULTS**

Twenty-one patients were included in this study. Demographic characteristics of the participants are presented in Table 1. The mean age of the study population was 55.3 (SD=14.2) years. The patients were classified into 2 groups on the basis of a median SUV<sub>max</sub> threshold of 0.17. Patients with SUV<sub>max</sub>  $\leq$  0.17 were classified as group 1. Patients with increased  $^{68}$ Ga-NOTA-PRGD2 uptake (SUV<sub>max</sub> > 0.17) were classified into group 2. The patients in group 2 had a significantly higher SUV<sub>max</sub> than those in group 1 (0.45±0.22 versus 0.06±0.08, P<0.001) (Figure 2).

### Spatial Distribution of <sup>68</sup>Ga-NOTA-PRGD2 Uptake

We analyzed <sup>68</sup>Ga-NOTA-PRGD2 uptake in WMH, NAWM, and the border between WMH and NAWM. We found that <sup>68</sup>Ga-NOTA-PRGD2 uptake was more obvious on the border between the WMH and NAWM (Figure 3). In group 2, <sup>68</sup>Ga-NOTA-PRGD2 SUV<sub>max</sub> was significantly higher than that of WMH (*P*=0.004). In group 1, the SUV<sub>max</sub> was not significantly different among the 3 regions (Figure 4).

Figure 5A shows a comparison of  $^{68}$ Ga-NOTA-PRGD2 uptake between the WMH and NAWM layers (L1–L7) in groups 1 and 2.  $^{68}$ Ga-NOTA-PRGD2 SUV, was significantly higher in NAWM L1 to L7 than in WMH (P<0.001) in group 2. However, there was no significant increase in  $^{68}$ Ga-NOTA-PRGD2 SUV, between WMH and NAWM L1 to L7 in group 1 (P=0.251).

#### **Influencing Factors of Angiogenesis**

Patients in group 2 had a larger WMH volume than those in group 1 (P=0.002) (Table 1). The serum levels of integrin  $\alpha\nu\beta3$ , VEGF receptor 2, and interleukin-1 $\beta$  were significantly higher in group 2 than in group 1 (P=0.038, P=0.03, and P=0.012, respectively). There were no significant differences in the serum levels of S100B and

Table 1. Summary of Participant Characteristics

	Total (n=21)	Group 1 (n=7)	Group 2 (n=14)
Age, y, mean±SD (range)	55.33±14.20 (25-79)	58.29±20.10 (25–79)	53.86±10.81 (31-66)
Sex, female, n (%)	15 (71.4%)	7 (100%)	8 (57.1%)
Hyperlipidemia, n (%)	8 (38.1%)	4 (57.1%)	4 (28.6%)
Hypertension, n (%)	16 (76.2%)	5 (71.4%)	10 (71.4%)
Diabetes, n (%)	3 (14.3%)	2 (28.6%)	1 (7.1%)
eGFR [mL/(min×1.73 m <sup>2</sup>	95.43±29.14	84.25±22.40	101.46±31.33
MMSE score, mean (SD)	23.33±6.70	22.14±7.02	25.71±5.74
BPF, mean (SD)	0.70±0.07	0.73±0.09	0.69±0.06
WMH volume, mL, mean (SD)	41.80±30.88	17.56±15.30	53.92±29.73
Number of lacunes, median (IQR)	3 (1.5–8)	4 (2-5)	3 (1–10)
Number of CMBs, median (IQR)	10 (2.5–21)	6 (5–15)	12 (0.75–24)
Severe CSO-PVS, n (%)	15 (71.4%)	5 (71.4%)	10 (71.4%)
Severe BG-PVS, n (%)	8 (38.1%)	3 (42.9%)	5 (35.7%)
WMH SUV <sub>max</sub> , mean (SD)	0.14±0.14	0.01±0.01	0.21±0.12
WMH-NAWM SUV <sub>max</sub> , mean (SD)	0.29±0.24	0.03±0.05	0.41±0.19
NAWM SUV <sub>max</sub> , mean (SD)	0.24±0.24	0.04±0.08	0.34±0.24

BG indicates basal ganglia; BPF, brain parenchymal fraction; CMBs, cerebral small-vessel diseases; CSO, centrum semiovale; eGFR, estimated glomerular filtration rate; MMSE, Mini-Mental State Examination; NAWM, normal-appearing white matter; PVS, perivascular spaces; SUV<sub>max\*</sub> maximum standardized uptake value; and WMH, white matter hyperintensity.

HIF-1 $\alpha$  between the 2 groups (P=0.689 and P=0.062, respectively) (Table 2). After applying multiple comparisons to the number of tests, no significant associations were found between the 2 groups (P<0.05/13=0.004).

Serum levels of integrin  $\alpha v\beta 3$  were associated with WMH volume (r=0.545, P=0.011) and MMP-2 (r=0.689, P<0.001). In group 2 (n=14), serum levels

of integrin  $\alpha$ v $\beta$ 3 were associated with WMH volume (r=0.648, P=0.012) and MMP-2 (r=0.731, P=0.003) (Figure 6A and 6B). In group 1 (n=7), SUV<sub>max</sub> was associated with E-selectin (rho=0.875, P=0.010), interleukin-6 (rho=0.867, P=0.012), and S100B (rho=-0.835, P=0.019). In group 2, SUV<sub>max</sub> was associated with WMH volume (r=-0.683, P=0.007) (Figure 6C).

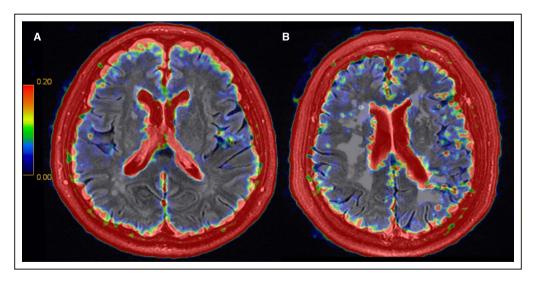


Figure 2. WMHs and 68Ga-NOTA-PRGD2 uptake.

**A**, Example of a patient in group 1. The integrated PET/MRI of <sup>68</sup>Ga-NOTA-PRGD2 PET and FLAIR imaging show no significant increase in <sup>68</sup>Ga-NOTA-PRGD2 uptakes in WMH and NAWM. **B**, Example of a patient in group 2. The integrated PET/MRI of <sup>68</sup>Ga-NOTA-PRGD2 PET and FLAIR imaging show significantly increased <sup>68</sup>Ga-NOTA-PRGD2 uptakes in WMH and NAWM. FLAIR indicates fluid attenuated inversion recovery; MRI, magnetic resonance imaging; NAWM, normal-appearing white matter; PET, positron emission tomography; and WMHs, white matter hyperintensities.

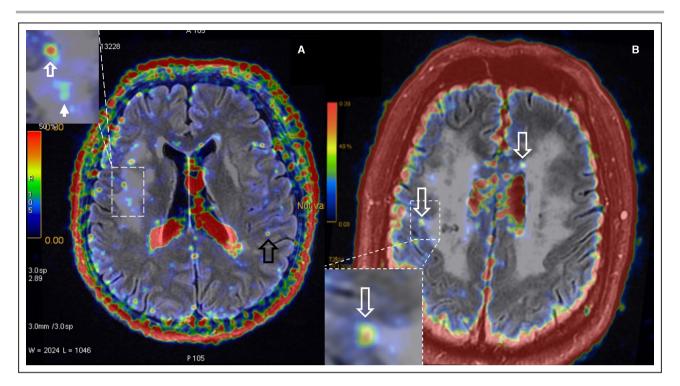


Figure 3. Distribution of <sup>68</sup>Ga-NOTA-PRGD2 uptake.

**A**, Example of a patient in group 2, the integrated PET/MRI of <sup>68</sup>Ga-NOTA-PRGD2 PET and FLAIR imaging show the pattern of <sup>68</sup>Ga-NOTA-PRGD2 uptakes in WMHs (solid white arrow), NAWM (faint black arrow), and the border of WMH and NAWM (faint white arrow). **B**, Example of a patient in group 2, the integrated PET/MRI of <sup>68</sup>Ga-NOTA-PRGD2 PET and FLAIR imaging show the pattern of <sup>68</sup>Ga-NOTA-PRGD2 uptakes in the border of WMH and NAWM (faint white arrow). FLAIR indicates fluid attenuated inversion recovery; MRI, magnetic resonance imaging; NAWM, normal-appearing white matter; PET, positron emission tomography; and WMHs, white matter hyperintensities.

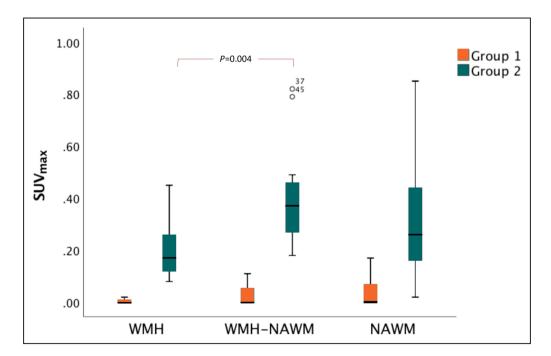


Figure 4. Spatial distribution of <sup>68</sup>Ga-NOTA-PRGD2 uptake.

<sup>68</sup>Ga-NOTA-PRGD2 uptakes were significantly increased at the border of WMH and NAWM than in WMH in group 2. NAWM indicates normal-appearing white matter; SUV<sub>max</sub>, maximum standardized uptake values; and WMH, white matter hyperintensities.

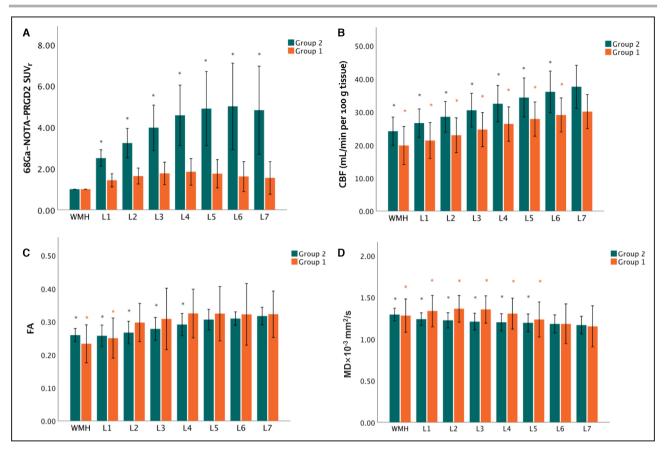


Figure 5. WMH penumbra.

**A**, The <sup>68</sup>Ga-NOTA-PRGD2 SUV<sub>r</sub> was significantly increased in NAWM layers (L1–L7) than in WMHs in group 2. **B**, The mean CBF of WMHs and NAWM L1 to L6 was significantly lower than that of the NAWM L7 in groups 1 and 2. **C**, The mean FA of WMHs and NAWM L1 was significantly lower than that of NAWM L7 in group 1. The mean FA of WMHs and NAWM L4 was significantly lower than that of NAWM L7 in group 2. **D**, The MD of WMHs and NAWM L1 to L5 in groups 1 and 2. CBF indicates cerebral blood flow; FA, fractional anisotropy; MD, mean diffusivity; NAWM indicates normal-appearing white matter; SUV<sub>r</sub>, standardized uptake value ratios; and WMHs, white matter hyperintensities.

#### Angiogenesis and CBF WMH Penumbra

For patients without significantly increased  $^{68}$ Ga-NOTA-PRGD2 uptake (group 1), the mean CBF of WMH and NAWM L1 to L6 was significantly lower than that of NAWM L7 (P<0.001). For patients with significantly increased  $^{68}$ Ga-NOTA-PRGD2 uptake (group 2), the mean CBF of WMH and NAWM L1 to L6 was also significantly lower than that of NAWM L7 (P<0.001) (Figure 5B). We further investigated the correlation between the mean CBF and  $^{68}$ Ga-NOTA-PRGD2 uptake. The results showed a negative correlation between SUV, and CBF in group 2 (r=-0.238, P=0.018). However, there was no significant correlation between SUV, and CBF across different levels.

# Angiogenesis and Structural WMH Penumbra

For patients without significantly increased  $^{68}$ Ga-NOTA-PRGD2 uptake (group 1), the mean FA of WMH and

NAWM L1 was significantly lower than that of NAWM L7 (P=0.005 and P=0.017, respectively) (Figure 5C). The mean MD of WMH and NAWM L1 to L5 was higher than that of NAWM L7 (P=0.004) (Figure 5D).

For patients with significantly increased <sup>68</sup>Ga-NOTA-PRGD2 uptake (group 2), a comparison between WMH and NAWM demonstrated that the mean FA of WMH and NAWM L1 to L4 was significantly lower than that of NAWM L7 (*P*<0.001) (Figure 3C). The mean MD of WMH and NAWM L1 to L5 was higher than that of NAWM L7 (*P*<0.001) (Figure 5D).

We investigated the correlation between white matter integrity and  $^{68}\text{Ga-NOTA-PRGD2}$  uptake. The results showed significant correlation between SUV<sub>r</sub> and FA in group 2 (L2 [r=-0.552, P=0.041]; L3 [r=-0.684, P=0.007]; L4 [r=-0.741, P=0.002]; L5 [r=-0.789, P<0.001]; L6 (r=-0.798, P<0.001); L7 [r=-0.714, P=0.004]) (Figure 7). There was no significant correlation between SUV<sub>r</sub> and MD in WMH and NAWM L1 to L7.

Table 2. Angiogenic and Inflammatory Factors

Factors (ng/mL)	Group 1	Group 2	P value	SMD (95% CI)
hs-CRP, mg/L	1.16±1.083	1.84±2.96	0.596	-0.68 (-3.18 to 1.81)
Interleukin-1β	0.01±0.01	0.04±0.07	0.012	-0.04 (-0.09 to 0.02)
Interleukin-6	0.02±0.02	0.09±0.25	0.743	-0.08 (-0.280 to 0.13)
Interleukin-8	0.01±0.02	0.04±0.06	0.636	-0.08 (-0.28 to 0.13)
E-selectin	3.00±4.14	4.06±3.40	0.36	-0.02 (-0.06 to 0.01)
VEGF-A	0.03±0.02	0.03±0.02	0.971	0.00 (-0.02 to 0.02)
Integrin ανβ3	0.32±0.26	0.58±0.30	0.038	-0.26 (-0.54 to 0.02)
MMP-2	0.81±0.45	1.61±1.11	0.085	-0.78 (-1.71 to 0.15)
MMP-9	7.85±1.71	8.07±3.24	0.873	-0.22 (-2.99 to 2.54)
VEGF receptor 2	8.10±1.64	1.09±3.77	0.030	-0.27 (-0.53 to 0.02)
VEGF receptor 1	0.47±0.32	0.61±0.25	0.269	-0.14 (-0.41 to 0.13)
HIF-1α	0.02±0.02	0.05±0.04	0.062	-0.03 (-0.061 to 0.00)
S100B	3.91±5.53	2.90±2.20	0.689	1.01 (-2.48 to 4.50)

 $HIF-1\alpha$  indicates hypoxia-inducible factor-1 alpha; hs-CRP, high-sensitivity C-reactive protein; MMP, matrix metalloproteinase; SMD, standardized mean difference; and VEGF, vascular endothelial growth factor.

#### DISCUSSION

In this study, we found that angiogenesis was present in patients with WMH. A higher WMH burden is a predictive factor for angiogenesis in cerebral small-vessel disease. More specifically, we found that integrin  $\alpha\nu\beta3$  expression was significantly increased on the border between the WMH and NAWM. These findings suggest that angiogenesis may occur in the WMH penumbra in patients with a high WMH burden.

Previous investigations have investigated the WMH penumbra in the NAWM tissue surrounding WMHs, which may represent a subtle vascular and neuronal injury.  $^{5-8}$  In the present study, we found that the structural penumbras, as defined by FA and MD, were  $\approx\!2$  to 10 mm surrounding the WMH; CBF penumbras, as defined by CBF, were  $\approx\!12\,\text{mm}$  from WMH. These results are consistent with those of previous studies, which reported that the CBF penumbra was  $\approx\!11$  to 14 mm surrounding the WMH, whereas the DTI-FA penumbra covered  $\approx\!2$  to 9 mm from the WMH.  $^{7.8}$  We observed that the structural penumbra was narrower in the group without significant angiogenesis and with lower WMH volume. Additionally, our results confirmed that the CBF penumbra was more extensive than the structural penumbra.

However, whether hypoxia promotes angiogenesis in the central nervous system remains unknown. Angiogenesis is a complex process that involves numerous factors. We found that serum levels of integrin  $\alpha\nu\beta3$ , VEGF receptor 2, and interleukin-1 $\beta$  were higher in the group with significantly increased  $^{68}\text{Ga-NOTA-PRGD2}$  uptake, indicating the regulatory effects of the angiogenic factors in this study. Previous studies have shown that cerebrospinal fluid biomarkers of angiogenesis, such as VEGF, VEGF receptor 1, VEGF receptor 2, and placental growth factor are associated

with more pronounced white matter lesions in patients with Parkinson disease. Hypoperfusion and hypoxia associated with orthostatic hypotension have been proposed as potential mechanisms.  $^{15}$  It is likely that the state of hypoperfusion and chronic ischemia that leads to the manifestation of WMH on brain MRI also acts as a trigger for the upregulation of angiogenic factors. Higher levels of intercellular adhesion molecule 1, E-selectin, neopterin, and vascular cell adhesion molecule 1 were observed in patients with greater WMH volumes and lacunar infarcts. Previous studies have also emphasized that interleukin-1 $\beta$  could have an impact on white matter lesions and angiogenesis.  $^{24,25}$ 

In this study, we found that both reduced CBF and microstructural integrity are correlated with angiogenesis. We hypothesized that compromised CBF precedes microstructural integrity changes and induces angiogenesis. Khan et al<sup>26</sup> reported that chronic remote ischemic conditioning can induce angiogenesis and cerebral vascular remodeling. Previous studies using animal models of chronic cerebral hypoperfusion have shown that bilateral common carotid artery stenosis—induced ischemic white matter damage is significantly improved after transplantation of bone marrow mononuclear cells through endothelial nitric oxide synthase activation, upregulation of the VEGF–VEGF receptor 2 signaling pathway, and subsequent angiogenesis.<sup>27,28</sup>

The WMH penumbra has been associated with expansion of the WMH. However, the causal relationship between the reduction in CBF and microstructural integrity with WMH progression remains controversial. The deep white matter is particularly susceptible to injury from hypoperfusion because this area is supplied exclusively by perforating arterioles arising from the leptomeningeal border zone and must travel a long distance to reach the deep white matter.<sup>9,29–31</sup>

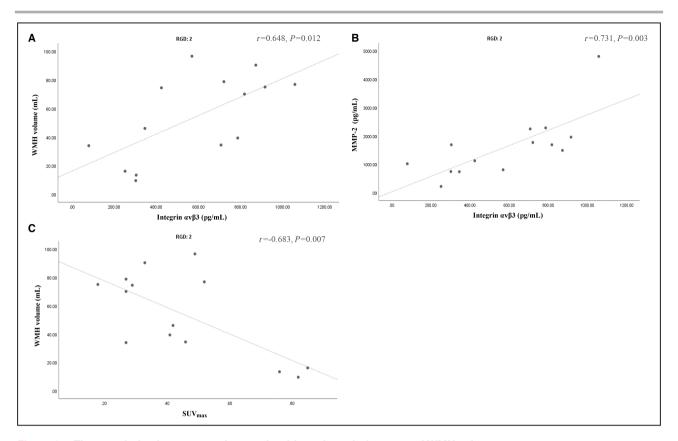


Figure 6. The correlation between angiogenesis with angiogenic factors and WMH volume.

A, The correlation between serum integrin  $\alpha v \beta 3$  and WMH volume in group 2. B, The correlation between serum integrin  $\alpha v \beta 3$  and MMP-2 in group 2. C, The correlation between SUV<sub>max</sub> and WMH volume in group 2. MMP indicates matrix metalloproteinase; SUV<sub>max</sub>, maximum standardized uptake value; and WMH, white matter hyperintensity.

Staffaroni et al<sup>32</sup> found that longitudinal low baseline CBF was associated with an increased WMH burden. Promjunyakul et al<sup>33</sup> found that baseline reduced CBF and microstructural characteristics of the WMH penumbra region were predictors of WMH growth, especially the inner 5 mm of the penumbra.

Whether angiogenesis in the WMH penumbra has an impact on WMH growth remains unknown. Ischemiainduced angiogenesis may be a possible mechanism to compensate for hypoperfusion in WMH and may be beneficial for remyelination and white matter tract rewiring. Recent studies have demonstrated that angiogenesis plays a beneficial role in accelerating white matter remodeling. Oligodendrocytes may secrete angiogenic factors, such as MMP-9, which accelerate angiogenesis and promote white matter remodeling after white matter injury.<sup>25</sup> In contrast, VEGF, MMPs, and other angiogenesis factors could lead to blood-brain barrier disruption. 14,34-36 Blood-brain barrier disruption is considered to play a role in the pathophysiology of WMH. However, we did not find significant differences in S100B levels in this study. Future studies using neuroimaging to precisely measure the blood-brain barrier will be more instructive.

This study is the first to provide in vivo evidence of an association between angiogenesis and WMH

penumbra. We used a novel noninvasive method,  $^{68}$ Ga-NOTA-PRGD2 PET/MRI, to detect angiogenesis in vivo. Nuclear medicine and molecular imaging are complementary imaging methods for a better understanding of the mechanisms underlying central nervous system diseases. Integrated PET/MRI enables the analysis of both anatomic structures and metabolic processes in cerebrovascular diseases. Furthermore, integrin  $\alpha v \beta 3$  has been a popular treatment modality for pro- or antiangiogenesis approaches. This insight will provide new therapeutic directions for cerebral small-vessel disease treatment.

This study has several limitations. First, our cross-sectional analysis did not prove causality. Further prospective studies are needed to verify the impact of angiogenesis on WMH growth. Second, our results were limited by the small sample size. The multiple comparisons may lead to a high rate of false negatives or false positives in such small sample size, and the possibility of residual confounding cannot be excluded. The results showed some floor effects and nonlinear associations between angiogenesis and structural WMH penumbra. Further study with a large sample size should be performed to prove the linear correlations. Third, in group 2 of patients with high WMH burden, the WMH penumbra and NAWM were closer to the cortex, which

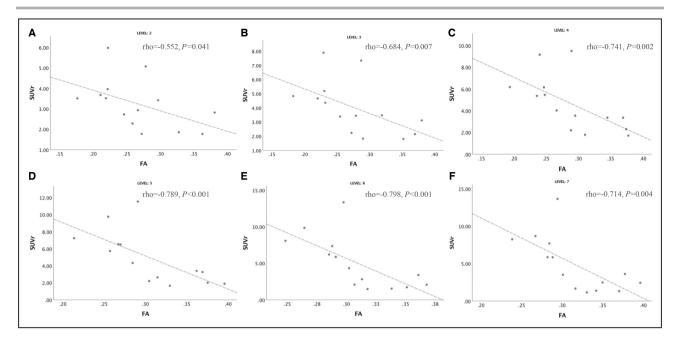


Figure 7. The correlation between <sup>68</sup>Ga-NOTA-PRGD2-SUV<sub>r</sub> and FA.
The <sup>68</sup>Ga-NOTA-PRGD2 SUV<sub>r</sub> was correlated with FA in NAWM layers (L2–L7) in group 2. FA indicates fractional anisotropy; NAWM, normal-appearing white matter; and SUV<sub>r</sub>, standardized uptake value ratios.

may have induced relatively greater CBF values. Future studies with larger sample sizes should be conducted to analyze the differences in CBF penumbra in different locations, including deep WMH and periventricular WMH. Fourth, magnetic resonance attenuation correction was performed in this study, which disregards the potential influence of cortical bone. Previous studies have shown that magnetic resonance attenuation correction led to underestimation of PET values in comparison to computed tomography-based attenuation correction, which may explain the relatively low standardized uptake values in our study. However, the difference is typically <10%.37,38 Besides, there are several advantages. Combined PET/MRI examinations may facilitate improvements in the additional use of magnetic resonance for retrospective motion correction or magnetic resonance-based partial volume correction of PET.<sup>22</sup> Finally, the integrity of the blood-brain barrier may affect <sup>68</sup>Ga-NOTA-PRGD2 uptakes. The consistency of <sup>68</sup>Ga-NOTA-PRGD2 uptakes and angiogenesis still needs to be confirmed by pathology study.

#### CONCLUSIONS

This study further defined the WMH penumbra pattern. Our results demonstrate that angiogenesis was present in the WMH penumbra. Additionally, our findings indicate that angiogenesis is correlated with reduced CBF and microstructural integrity. This study suggested the potential therapeutic value of angiogenesis in the prevention and treatment of WMH. Further

studies are warranted to verify the important role of angiogenesis in the development of WMHs and integrin  $\alpha\nu\beta$ 3-targeted molecular imaging as a noninvasive monitoring technique.

#### ARTICLE INFORMATION

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#### **Affiliations**

Department of Neurology, Beijing Tiantan Hospital, Capital Medical University, Beijing, China (L.D.); China National Clinical Research Center for Neurological Diseases, Beijing, China (L.D.); Department of Radiology, Peking Union Medical College Hospital, Chinese Academy of Medical Science and Peking Union Medical College, Beijing, China (B.H., T.S., F.F.); Department of Nuclear Medicine (J.Z., Z.Z.) and Department of Neurology (B.P.), Peking Union Medical College Hospital, Chinese Academy of Medical Science and Peking Union Medical College, Beijing, China and Department of Neurology, State Key Laboratory of Complex Severe and Rare Diseases, Beijing, China (B.P.).

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#### **Disclosures**

None.

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