Pericyte-derived bone morphogenetic protein 4 underlies white matter damage after chronic hypoperfusion

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Abstract

Subcortical small vessel disease (SVD) is characterized by white matter damage resulting from arteriolsclerosis and chronic hypoperfusion. Transforming growth factor beta 1 (TGFβ1) is dysregulated in the hereditary SVD, CARASIL (cerebral autosomal recessive arteriopathy with subcortical infarcts and leukoencephalopathy). However, very little is known about the role of the largest group in the TGFβ superfamily – the bone morphogenetic proteins (BMPs) – in SVD pathogenesis. The aim of this study was to characterize signaling abnormalities of BMPs in sporadic SVD. We examined immunostaining of TGFβ1 and BMPs (BMP2/BMP4/BMP6/BMP7/BMP9) in a total of 19 post-mortem human brain samples as follows: 7 SVD patients (4 males, 76–90 years old); 6 Alzheimer’s disease (AD) patients (2 males, 67–93 years old) and 6 age-matched disease controls (3 males, 68–78 years old). We subsequently investigated the effects of oxygen–glucose deprivation and BMP4 addition on cultured cells. Furthermore, adult mice were subjected to chronic cerebral hypoperfusion using bilateral common carotid artery stenosis, followed by continuous intracerebroventricular infusion of the BMP antagonist, noggin. In the SVD cases, BMP4 was highly expressed in white matter pericytes. Oxygen–glucose deprivation induced BMP4 expression in cultured pericytes in vitro. Recombinant BMP4 increased the number of cultured endothelial cells and pericytes and converted oligodendrocyte precursor cells into astrocytes. Chronic cerebral hypoperfusion in vivo also upregulated BMP4 with concomitant white matter astrogliogenesis and reduced oligodendrocyte lineage cells, both of which were suppressed by intracerebroventricular noggin infusion. Our findings suggest ischemic white matter damage evolves in parallel with BMP4 upregulation in pericytes. BMP4 promotes angiogenesis, but induces astrogliogenesis at the expense of oligodendrocyte precursor cell proliferation and maturation, thereby aggravating white matter damage. This may explain white matter vulnerability to chronic hypoperfusion. The regulation of BMP4 signaling is a potential therapeutic strategy for treating SVD.

INTRODUCTION

Vascular cognitive impairment develops as a consequence of various types of cerebrovascular alterations. Subcortical white matter changes caused by small vessel alterations are frequently observed in vascular cognitive impairment and are referred to as subcortical ‘small vessel disease (SVD)’ (39). Disturbances in cerebrospinal fluid production (40), cerebral edema (22), breakdown of the blood–brain barrier and increased permeability (21, 54), oxidative stress (4), and inflammation have been cited as important causes in the development of white matter changes (19). However, the exact mechanisms have yet to be fully elucidated. Recent studies have noted that attenuations of vasculature and white matter are also frequently observed in other neurodegenerative disorders, especially in Alzheimer’s disease (AD) (46, 59). However, vascular risk factors are related to a lesser degree to a pure type of AD (10, 38) or mixed type (AD with vascular pathology) dementia (23) compared with vascular cognitive impairment. The involvement of different
etiolologies has been suggested in white matter damage between SVD and AD (13, 20). Therefore, clarification of causative factors that underlie various forms of white matter changes should enable the development of novel strategies for tackling cognitive impairment.

There are several recognized forms of inherited SVD (9), including CARASIL (cerebral autosomal recessive hereditary cerebral artery disease and arteriosclerosis with subcortical infarcts and leukoencephalopathy). The gene responsible for CARASIL is HTRA1 (high-temperature requirement A 1), and the resultant upregulation of transforming growth factor beta (TGFB) family signaling is postulated to underlie the small vessel changes observed in CARASIL (18). The TGFB superfamily also includes the bone morphogenetic protein (BMP) family proteins.

BMPs are involved in oligovascular pathologies in the ischemic brain, in addition to their physiological roles in embryonic and bone tissues. In an ischemic intrauterine growth retardation model, oxidative stress upregulates BMP4 and mediates periventricular white matter injury with a paucity of mature oligodendrocytes and hypomyelination, while BMP deletion reverses these defects (36). In neonatal mouse brains, BMP4 expression increases as a consequence of global hypoxia-ischemia, while the BMP antagonist noggin protects the white matter against damage (12). Furthermore, in adult mouse brains with focal cerebral ischemia, overexpressed noggin reduces infarct volume and motor deficits (44). These experimental results highlight the potential of BMP antagonism in the treatment of ischemic demyelinating disorders.

Nevertheless, the exact role of BMP family members remains poorly understood in the context of chronic hypoperfusion in the adult human brain. Therefore, the present study explored the underlying etiology of white matter abnormalities, specifically focusing on BMP expression through the use of postmortem human brains, cultured cells exposed to oxygen–glucose deprivation (OGD), and chronic hypoperfusion mouse brains.

**MATERIALS AND METHODS**

**Experimental design of postmortem brain material**

Study samples were selected from 371 autopsied brains retained at Kyoto University Hospital from 1983 to 2009. This collection was approved by the institutional research committee, Kyoto University Graduate School and Faculty of Medicine, Ethics Committee. Neuropathological diagnoses were made according to thorough histopathological examination of extensively sampled brain sections, as previously described (1, 35). Cases with mixed SVD and AD pathological diagnoses were excluded to select for brain samples with relatively single processes. We analyzed a total of 19 postmortem brain samples as follows: 7 SVD patients (4 males, 76–90 years old, brain weight 1101±89.9 g); 6 AD patients (2 males, 67–93 years old, brain weight 1056±67.7 g) and 6 age-matched disease controls (3 males, 68–78 years old, brain weight 1190±113.7 g). Table 1 provides demographics and pathological features of the 19 subjects. The clinical diagnosis of dementia met the criteria of the Diagnostic and Statistical Manual of Mental Disorders IV (2). The AD neuropathological diagnoses were made according to the presence of frequent senile plaques in the neocortex (32), and no less than stage V of Braak stage of neurofibrillary tangles (6, 7). The diagnosis of SVD, or subcortical ischemic vascular dementia was clinically made (5) and was retrospectively found to meet the pathological criteria outlined by Kalaria et al (24). Two observers (M.T.U. and M.I.) individually assessed senile plaque and

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AD = Alzheimer’s disease; ALS = amyotrophic lateral sclerosis; BW = brain weight; FMI = frontal myelin index; NFT = neurofibrillary tangle; SP = senile plaque; SVD = small vessel disease.
neurofibrillary tangle stages, and if required, a joint assessment was scrutinized under a two-headed microscope.

**Tinctorial staining and immunohistochemistry for postmortem human brain samples**

Six micrometer-thick paraffin-embedded tissue sections were cut from the frontal lobes at the level of the olfactory bulbs and stained for Luxol fast blue. Some sections were also incubated with specific primary antibodies (listed in Supporting Information Table S1) overnight at 4°C. Sections were pretreated at 90°C for 20 min in Tris-ethylenediamine tetraacetic acid (pH 9.0) for actin, alpha 2, smooth muscle, aorta (ACTA2, also known as αSMA) and platelet-derived growth factor receptor alpha (PDGFRA), and some were pretreated at 121°C for 15 min in 0.01 M citrate buffer (pH 6.0) using an autoclave for BMP2. This was followed by appropriate biotinylated secondary antibodies (1:200, Vector Laboratories, Burlingame, CA, USA) and avidin-biotin-peroxidase complex (1:200, Vector Laboratories, Burlingame, CA, USA) and visualized with 0.1 M phosphate-buffered saline (PBS, pH 7.4) and polymer Detection System (Histofine Simple Stain MAX PO MULTI, Nichirei Biosciences, Tokyo, Japan) application. Between steps, the sections were washed three times for 5 min with 0.1 M phosphate-buffered saline (PBS, pH 7.4) and visualized using 0.01% diaminobenzidine tetrahydrochloride and 0.003% H2O2 in 50 mM Tris-HCl (pH 7.5).

For triple-immunofluorescence, deparaffinized sections were incubated with a mixture of primary antibodies overnight at 4°C, followed by Alexa Fluor 488, 594 and 647 conjugated secondary antibodies (1:200, Invitrogen, Carlsbad, CA, USA) for 2 h at room temperature. The images of interest were captured with a microscope (BX2700, Keyence, Osaka, Japan) or a confocal laser-scanning microscope (FV1000, Olympus, Tokyo, Japan).

**Assessment of myelin density or immunohistochemical staining**

Myelin density was assessed using immunohistochemistry and myelin index as previously described (20, 58). Briefly, to determine the myelin index, the white matter was automatically outlined using the wand tool of Image J. If the border between the cortex and white matter was obscured, the white matter was manually outlined using the semiautomatic trace tool instead of the wand tool. The detected range of gray levels within the white matter, corresponding to the staining intensity, from 0 to 127 (0, white; 255, black) was divided into four quartiles (the first quartile 0–29, the second 30–62, the third 63–94 and the fourth 95–127). The percent area for each quartile was calculated. The median gray level of each quartile (14.5, 46.0, 78.5 and 111.0), an estimated staining intensity, was then multiplied by percent area in each quartile and summed up, providing the total myelin index (Figure 1A).

For immunohistochemical staining, images from 15 to 20 randomly selected ROIs were captured. The images were then converted into 8-bit grayscale, binarized by threshold using the Triangle Method, and the area above threshold measured using Fiji and Image J. For quantification of BMP4 immunoreactivity in PDGFRB- and ACTA2-positive cells, images of PDGFRB or ACTA2 immunostaining were firstly converted into binarized images using the Triangle Method and the area above threshold was selected. The selected area was applied to the same section with BMP4 immunostaining, and the mean intensity and percent area of BMP4 in the selected area measured.

The above analyses were performed blinded to the diagnosis by labeling sections with arbitrary numbers.

**Cell culture**

Mouse brain vascular pericytes (1200, ScienCell, Carlsbad, CA, USA) and human brain microvascular endothelial cells (ACBRI, HBDMC, Cell Systems, Kirkland, WA, USA) were cultured with pericyte medium (1201, ScienCell) or endothelial cell growth medium (CC-3156 and CC-4147, Lonza, Basel, Switzerland) containing 10% fetal bovine serum and 1% penicillin/streptomycin. Endothelial cells were placed on pre-coated dishes with 3.2% bovine collagen type IV alpha 1 chain (COL4A1) (A1064401, Invitrogen) in PBS. Oligodendrocyte precursor cells (OPCs) were isolated from cerebral cortices from 1- to 2-day-old Sprague Dawley rats as previously described (8, 30).

**Oxygen–glucose deprivation**

Upon confluency, pericyte media were replaced with glucose- and serum-free pericyte media. The following day, culture plates were placed into a hypoxic culture kit (BIONIX II, Sugiyama-Gen, Tokyo, Japan) with 5% O2 and 3% CO2, and incubated at 37°C for 1, 2 and 3 days.

**RT-PCR**

After preparing the cell lysate, mRNA was rapidly purified using the RNeasy Plus Mini Kit (74136, Qiagen, Hilden, Germany) and subjected to reverse transcriptase-polymerase chain reaction (RT-PCR) analysis using appropriate primers (listed in Supporting Information Table S2).

**BMP4 and noggin treatment**

The media were replaced with new medium with or without recombinant human BMP4 (1 or 10 ng/ml, 314-BP, R&D systems) and/or recombinant human noggin (100 ng/ml for proliferation assay and 500 ng/ml for differentiation assay; 120–10°C, PeproTech, IL, USA). The cells treated with both BMP4 and noggin were pretreated with only noggin (100 ng/ml for proliferation assay and 500 ng/ml for differentiation assay) for 2 h.

**WST-8 assay**

Two days after BMP4 and/or noggin administration, 1% of 2-(2-methoxy-4-nitrophenyl)-3-(4-nitrophenyl)-5-(2,4-disulfophenyl)-2H-tetrazolium, monosodium salt solution (WST-8, 07553–44, Nacalai Tesque, Kyoto, Japan) was added to each medium. Then, the plates were incubated for 1 h at 37°C in the incubator. Absorbance at 450 nm and 630 nm were measured using a multi-label plate reader (2030 ARVO X, PerkinElmer, Waltham, MA, USA). The results of absorbance at 450 nm were corrected to absorbance at 630 nm.
The cells were fixed with 4% paraformaldehyde/PBS for 15 min at room temperature, treated with 3% bovine serum albumin for 1 h, incubated with primary antibodies (listed in Supporting Information Table S3) overnight at 4°C and appropriate secondary antibodies for 2 h at room temperature. Between steps, the cells were washed three times for 5 min with PBS.

**Immunocytochemistry**

The cells were fixed with 4% paraformaldehyde/PBS for 15 min at room temperature, treated with 3% bovine serum albumin for 1 h, incubated with primary antibodies (listed in Supporting Information Table S3) overnight at 4°C and appropriate secondary antibodies for 2 h at room temperature. Between steps, the cells were washed three times for 5 min with PBS.

**Tube formation assay**

After thawing overnight at 4°C, the matrigel (354230, BD Biosciences, Franklin Lakes, NJ, USA) was applied on each well of a slide (81506, Ibidi, Madison, WI, USA) and polymerized for 1 h at 37°C. Then, 50 μl of suspension at a density of 4 × 10⁵ cells/ml endothelial cells were added to the matrigel surface. After seeding the cells, the slides were incubated for 24 h at 37°C. The number of tubes was quantified under a microscope.

**OPC proliferation assay and differentiation assay**

For the proliferation assay, OPCs were cultured with OPC proliferation media (Neurobasal medium containing 2 mM glutamine, 1% penicillin/streptomycin, 10 ng/ml platelet-derived growth factor-AA, 10 ng/ml basic fibroblast growth factors and 2% B27 supplement). At 30% confluency, the cells were treated with BMP4 and/or noggin and incubated for 2 days until harvest and analysis by RT-PCR.

For the differentiation assay, when the cells were 70% confluent, the OPC proliferation media were replaced with OPC differentiation media (Dulbecco's-modified Eagle Medium containing 1% penicillin/streptomycin, 10 ng/ml ciliary neurotrophic factor, 10 ng/ml triiodothyronine and 2% B27 supplement). Simultaneously, BMP4 and/or noggin were administered. Four days after treatment, the cells were analyzed by immunocytochemistry or RT-PCR.

Changes in OPC morphologies were captured using a time-lapse camera (BZ-X700, Keyence).

**Animals and surgical procedure**

The protocol for this study was approved by the Institutional Animal Care and Use Committee, Institute of Laboratory Animals Graduate School of Medicine, Kyoto University.
Adult C57BL/6J male mice (10–12 weeks old) were subjected to bilateral common carotid artery stenosis (BCAS) using microcoils as previously described \( (n = 6) \) (29, 48). Control mice underwent sham surgery \( (n = 6) \). Four weeks after BCAS, mice were euthanized and their brains analyzed by immunohistochemistry and western blot (Figure 6A).

In a separate experiment, mice received continuous intracerebroventricular infusion \( (cICV) \) 2 days prior to BCAS using brain infusion kit 3 \((0008851, \text{Alzet, Cupertino, CA, USA})\) and microosmotic pump \((1004, \text{Alzet})\) under stereotaxis. The pumps were filled with recombinant human noggin \((500 \text{ ng/day or } 1000 \text{ ng/day})\) in artificial cerebrospinal fluid \((\text{ACSF, 3525, Funakoshi, Tokyo, Japan})\), or ACSF only, as previously described \((42, 43)\). Control mice underwent sham surgery \( (n = 6 \text{ for sham control}; n = 6 \text{ for BCAS and noggin } 500 \text{ ng/day}; n = 6 \text{ for BCAS and noggin } 1000 \text{ ng/day}) \) (Figure 6H).

**Immunohistochemistry for mouse brain**

Mice were anesthetized and intracardially perfused with cooled PBS. The brains were quickly frozen using powdered dry ice. Coronal sections \((20-\mu \text{m thick})\) were cut on a cryostat at \(-20^\circ \text{C}\) and collected onto glass slides. As previously described \((47)\), sections were fixed with \(4\% \text{ paraformaldehyde/PBS for 15 min and incubated with antibodies (listed in Supporting Information Table S4).} \)

Quantitative analyses of glial fibrillary acidic protein (Gfap)-positive astrocytes, oligodendrocyte transcription factor (Olig2)-positive oligodendrocyte lineage cells, and myelin basic protein (Mbp)-positive mature oligodendrocytes were performed in the bilateral paramedian corpus callosum using Image J.

**Western blot**

Western blot analysis was performed as previously described \((28)\). Briefly, the dorsal half of the brain at 0–0.5 mm anterior to bregma was homogenized in radio-immunoprecipitation assay buffer, followed by sonication for 5 min with a 30-s interval and centrifugation at 14 000 rpm for 5 min at 4°C. The supernatant was boiled in sample buffer \((1\% \text{ w/v) sodium dodecyl sulfate (SDS), 12.5\% (w/v) glycerol, 0.005\% (w/v) bromophenol blue and 2.5\% (v/v) 2-mercaptoethanol in 25 mM Tris-HCl, pH 6.8.} \)

Samples were separated onto 10\% (w/v) gels for SDS-PAGE or 4–12\% (w/v) gels for NuPAGE \((\text{NP032330X, Invitrogen})\), followed by transfer to polyvinylidene difluoride membranes \((\text{Millipore, Billerica, MA, USA})\). The membranes were then incubated overnight at 4\°C with primary antibodies (listed in Supporting Information Table S5) followed by the appropriate horseradish peroxidase-conjugated secondary antibodies for 1 h at room temperature. Immunoreactive bands were detected with chemiluminescence assay kits \((02230, \text{Nacalai Tesque}), \text{ECL Western Blotting Substrate (32106, Thermo Fisher Scientific, Waltham, MA, USA)}\), and Amersham Imager 600 \((\text{GE Healthcare, Buckinghamshire, UK})\).

**Statistical analysis**

Statistical analysis was performed using the R statistical package \((\text{www.r-project.org})\). Statistical significance was evaluated using two-tailed paired Student’s \( t \)-test, the Wilcoxon rank-sum test or one-way ANOVA followed by the Bonferroni post-hoc test. Pearson correlation analysis was performed to observe a possible correlation. For all analyses, the level of statistical significance was set at \( **P < 0.01 \) or \( *P < 0.05 \).

**RESULTS**

**Myelin loss in SVD and AD compared with controls**

Luxol fast blue staining \((\text{Figure 1B})\) and MBP immunohistochemical staining \((\text{Figure 1C})\) represented severe myelin attenuation in the SVD group. The myelin index was significantly different between the SVD group and the other two groups as follows: controls \((80.9) > \text{AD (71.5) > SVD (56.3)} \) (Figure 1D), indicating that myelin loss was greatest in the SVD group. Age did not correlate with myelin index \( (r = -0.329, P = 0.170, \text{data not shown}) \). The percent area of MBP immunoreactivity also decreased in the SVD group \((\text{Figure 1E})\).

**Capillary bed density**

Capillary bed density was determined by immunohistochemistry using antibody specific to COL4A1, a major constituent of the basement membrane \((\text{Figure 1F})\). Quantitative analysis showed that the percentage of COL4A1 density was not different in the white matter or cortex between the three groups \((\text{white matter, } P = 0.095; \text{cortex, } P = 0.095) \) (Figure 1G).

**Pericyte density in the postmortem human brain**

Pericyte density was assessed by two different markers: ACTA2 and platelet-derived growth factor receptor beta \((\text{PDGFRB})\). The percentage of ACTA2 and PDGFRB expression, respectively, were divided by the percentage of COL4A1 stained area to correct for vascular density.

The ACTA2 immunoreactivity was not different in the white matter and cortex between the three groups \((\text{white matter, } P = 0.123; \text{cortex, } P = 0.220) \) (Figure 2A,B). Conversely, the percentage of PDGFRB expression was significantly increased in white matter of the SVD group compared with the other two groups \((\text{Figure 2C,D})\). There was a significant inverse correlation between the percentage of PDGFRB/COL4A1 expression in the white matter and myelin index in all 19 cases combined \( (r = -0.568, *P = 0.013) \) (Figure 2E).

**BMP4 expression in PDGFRB-positive pericytes of damaged white matter in the postmortem human brain**

BMP4 was strongly expressed in abluminal surface of capillaries and tunica media of arterioles, and mildly expressed in macrophages, and activated astrocytes in the white matter of SVD brains. Triple immunofluorescence studies revealed abundant BMP4 expression in PDGFRB- and ACTA2-positive pericytes in the white matter of SVD cases. However, in control or AD brains, BMP4 expression was weak \((\text{Figure 3A})\).

Similar to the percentage of PDGFRB/COL4A1 expression, the percentage of BMP4/COL4A1 expression was significantly increased in the white matter of SVD cases but was not different in the cortex between the three groups \((\text{Figure 3B})\). BMP4 expression...
negatively correlated with the myelin index ($r = -0.549$, *$P = 0.015$) (Figure 3C) in all 19 cases combined. Fluorescence intensity and percentage of BMP4-positive area in PDGFRB- and ACTA2-positive cells were increased in SVD cases (Figure 3D–G), suggesting BMP4 expression was increased in pericytes. BMP4 expression was also observed in activated astrocytes and macrophages in SVD, but not control or AD, cases (data not shown).

Other TGFβ superfamily members (TGFB1, BMP2, BMP6, BMP7 and BMP9)

Macrophages and activated astrocytes within the vicinity of microinfarcts were immunoreactive to all six TGF superfamily members. BMP2 was highly expressed in the endothelial cells of arterioles, venules and capillaries, and pericytes of capillaries in all three groups. BMP7 and TGFB1 were slightly expressed in the tunica media of arterioles and capillaries. BMP6 and BMP9 were not expressed in blood vessels (Supporting Information Figure S1).

Oligodendrocyte precursor cells and astrocytes in the postmortem human brain

PDGFRA-positive OPCs were found in the subventricular zone (SVZ), and cellular processes of some OPCs were found around arterioles, venules and capillaries (Figure 3H, Supporting Information Figure S2). The number of brain interstitial OPCs was measured in the SVZ of all cases and found to be increased relative to white matter attenuation, as previously described (33). However, when the SVD cases were divided into two groups by the cut-off point of 0.31 for BMP4/COL4A1, the higher BMP4/COL4A1 group (SVD 2, $n = 4$) tended to have a smaller number of OPCs than the lower BMP4/COL4A1 group (SVD 1, $n = 3$) (Figure 3I). The GFAP-positive astrocytes were increased in the white matter of SVD cases (Figure 3J).

BMP4 expression in cultured pericytes under oxygen–glucose deprivation

To confirm whether BMP4 is increased in pericytes under chronic hypoperfusion, we analyzed $Bmp4$ mRNA expression in cultured pericytes under continuous OGD. The mouse brain microvascular pericyte cell line expressed both $Pdgfrb$ and $Acta2$ (Figure 4A). $Bmp4$ mRNA expression exhibited a time-dependent increase under continuous OGD for 3 days (Figure 4B). $Pdgfrb$, but not $Acta2$, mRNA expression increased under OGD (Figure 4C,D). Expression of each mRNA sample was normalized to hypoxanthine-guanine phosphoribosyltransferase expression ($Hprt$) expression, which did not fluctuate under continuous OGD.

Effects of BMP4 on pericyte proliferation and $Pdgfrb$/$Acta2$ mRNA expression

The WST-8 assay showed that high-dose BMP4 induced pericyte proliferation (Figure 4E). BMP4 also increased $Pdgfrb$ (Figure 4F) and $Acta2$ (Figure 4G) mRNA expression levels.

Effects of BMP4 on endothelial cell proliferation and tube formation

Endothelial cell proliferation in response to BMP4 treatment was assayed by immunocytochemistry (Figure 4H) and WST-8 assay.
Endothelial cell proliferation and tube formation (Figure 4J,K) were facilitated in a dose-dependent manner by BMP4, but were blocked by the BMP4 antagonist, noggin.

**Effects of BMP4 on oligodendrocyte precursor cells**

In the OPC proliferation assay, treatment of BMP4 converted OPCs into astrocyte-like cells, with thick and long processes (Figure 5A and Supporting Information Movie S1). Immunocytochemistry showed that BMP4 treatment decreased Pdgfra (OPC marker) immunoreactivities and increased Gfap (astrocyte maker) (Figure 5B). RT-PCR also showed that BMP4 decreased Pdgfra mRNA levels, and significantly increased Gfap mRNA levels in a dose-dependent manner; noggin blocked the conversion (Figure 5C,D). Treatment with noggin alone increased Pdgfra mRNA levels in OPCs due to inhibition of endogenous BMP4 (Figure 5C).

The effects of BMP4 on OPCs were also observed in the OPC differentiation assay. In the differentiation media, primary OPCs differentiated into mature oligodendrocytes, whereas BMP4 treatment induced astrocyte-like cells (Supporting Information Figure S3A and Movie S2). BMP4 treatment decreased Mbp
**BMP4 derived from pericytes in SVD**

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**Figure 3.** Pericyte BMP4 expression in damaged white matter in the post-mortem human brain samples. A. Representative triple-immunofluorescent images for BMP4, PDGFRB, and ACTA2, and their merged images in the WM of all three groups. Differences are significant between any two groups, except between control and AD in the WM: control vs. AD, \( P = 0.894 \); control vs. SVD, \( *P = 0.003 \); AD vs. SVD, \( *P = 0.012 \). Differences are not significant in the cortex (\( P = 0.659 \)). B. An inverse correlation between the percentage of BMP4 expression and the myelin index in all 19 cases combined (\( r = -0.549, *P = 0.015 \)). D. The percent area of BMP4 expression in PDGFRB-positive cells in the WM of all three groups. Differences are significant between any two groups, except between control and AD: control vs. AD, \( P = 0.593 \); control vs. SVD, \( *P = 0.001 \); AD vs. SVD, \( *P = 0.011 \). E. The mean intensity of BMP4 in PDGFRB-positive cells in the WM of all three groups. Differences are significant between any two groups: control vs. AD, \( *P < 0.001 \); control vs. SVD, \( *P < 0.001 \); AD vs. SVD, \( *P = 0.017 \). F. The percent area of BMP4 expression in ACTA2-positive cells in the WM of all three groups. Differences are significant between any two groups, except between control and AD: control vs. AD, \( *P < 0.001 \); control vs. SVD, \( *P < 0.001 \); AD vs. SVD, \( *P = 0.017 \). G. The mean intensity of BMP4 in ACTA2-positive cells in the WM of all three groups. Differences are significant between any two groups: control vs. AD, \( *P < 0.001 \); control vs. SVD, \( *P < 0.001 \); AD vs. SVD, \( *P = 0.018 \). H. Representative images of PDGFRB expression in the interstitium of the SVZ in the cases of control, AD, and SVD of post-mortem human brain samples. According to expression levels of BMP4 (BMP4-to-COL4A1 ratio), the SVD cases are further classified into subgroup: a lower BMP4 expressing group (SVD 1) and a higher one (SVD 2) with the cut-off ratio of 0.31. Insets show enlarged images of OPCs. Bars indicate 50 \( \mu m \) and 10 \( \mu m \) (insets). **I.** The number of OPCs in all 4 groups. SVD 2 group shows smaller number of OPCs compared with SVD 1 group. **J.** Representative images of GFAP expression in the WM of post-mortem human brain samples. The WM of SVD shows severe astrogliosis with high GFAP expression. Bar indicates 50 \( \mu m \). Vertical bars represent mean \( \pm SEM \). Abbreviations are as follows: AD, Alzheimer’s disease; Cont, control; SVD, small vessel disease; SVZ, subventricular zone; WM, white matter.

(oligodendrocyte marker) and increased Gfap in a dose-dependent manner; noggin also blocked the conversion (Supporting Information Figure S3B–D).

**Bmp4 expression and effect of noggin in the mouse brain under chronic hypoperfusion**

Adult mice subjected to BCAS and noggin cICV (or ACSF only) treatment were analyzed by immunohistochemistry and western blot (Figure 6A,H). The inhibitory effect of noggin on BMP signaling was confirmed by a reduction of phosphorylated Smad immunoreactivity (data not shown).

Bmp4 was expressed in the Acta2/Pdgfrb double-positive pericytes in capillaries of 28-day-BCAS mouse brains (Figure 6B). Western blot showed significantly increased expression of Bmp4 precursor (Figure 6C,D and Supporting Information Figure S4A) and Pdgfrb (Figure 6E,F and Supporting Information Figure S4B), but not Acta2 (Figure 6E,G and Supporting Information Figure S4C) after BCAS, compared with sham controls. The Bmp4 precursor (41) was detected by both C-terminal (ab39973, Abcam, Cambridge, UK) and N-terminal (MAB1049, Millipore, and sc-393329, SantaCruz, Dallas, TX, USA) domain antibodies (data not shown). The expressions of western blot sample were normalized to tubulin gamma 1 (Tubg1) expression, which did not fluctuate under hypoperfusion.

The percent area of Gfap expression increased after BCAS, which was suppressed by noggin cICV (Figure 6J). The percent area of Olig2 expression was decreased by BCAS, which was ameliorated by noggin cICV at a dosage of 1000 ng/day (Figure 6I,K). Western blot also showed significantly increased Gfap expressions in the BCAS mice, which was suppressed by noggin cICV (both 500 ng/day and 1000 ng/day) (Figure 6L,M and Supporting Information Figure S4D). Olig2 expression was decreased by BCAS, which was ameliorated by a high dose of noggin cICV (1000 ng/ day) (Figure 6L,N and Supporting Information Figure S4E). Each band of Gfap and Olig2 was normalized with non-specific band stained with ponceau S staining (Figure 6L and Supporting Information Figure S4F). To assess the effect of BMP4 on mature oligodendrocytes, the percent area of Mbp expression was quantitated. The percent area of Mbp was decreased by BCAS, which was ameliorated by noggin cICV at a dosage of 1000 ng/day (Figure 6O,P).

**DISCUSSION**

We previously reported that myelin loss evolved in parallel with shrunken oligodendrocytes in SVD compared with AD (20). Another study revealed that spongiosis, arteriolosclerosis, état criblé and myelin loss were more severe in the SVD cases than the AD cases (13), suggesting different etiologies underlie the white matter attenuation between SVD and AD. In this study, we clarified one of the causative factors underlying the different aspects of white matter changes: namely, BMP4 generated from pericytes. We analyzed expressions of six TGFB superfamily members in the brains of SVD, AD, and age-matched controls, and found BMP4 was distinctly expressed in pericytes of the white matter. BMP4 expression was significantly upregulated in the SVD group, and was negatively correlated with myelin density. Consistent with these findings, long-term OGD in cultured pericytes and in vivo chronic hypoperfusion induced BMP4 upregulation. Given that chronic hypoperfusion during late embryonic stages (36) and neonatal stages (12) increases BMP4 expression and impairs differentiation of OPCs, chronic hypoperfusion may also upregulate BMP4 in the elderly. Recent evidence suggests that Tgfb upregulation contributes to cerebrovascular dysfunction in mice (34), which ascertainment CARASIL pathogenesis in humans through hereditary loss of HtrA1 (18). Therefore, these results suggest upregulation of the TGFB superfamily members is a shared mechanism in the pathogenesis of ischemic cerebrovascular disorders, regardless of whether it is sporadic or hereditary.

Previous studies have shown that Bmps, including Bmp2, 4, 6 and 7, are upregulated in a variety of CNS injury and demyelinating disease animal models (14, 45). In this study, expression of all six
TGFB superfamily members was also observed in macrophages and activated astrocytes within the vicinity of microinfarcts in the SVD group. However, in demyelinated but not infarcted areas of the SVD group, BMP4 was almost exclusively expressed in pericytes.

The density of PDGFRB-positive pericytes in the white matter correlated with myelin attenuation. In previous studies, PDGFRB immunostaining was used to assess microvasculature, showing a reduction of PDGFRB-positive pericytes and a
breakdown of the blood-brain barrier in vascular cognitive impairment and some neurodegenerative diseases, including AD and ALS (46, 51, 55, 56, 59). However, a recent study showed increased expression of PDGFRB immunoreactive pericytes in cerebral microvessels in CADASIL (cerebral autosomal dominant arteriopathy with subcortical infarcts and leukoencephalopathy) compared with similar age controls (11). It has also been suggested that Pdgfrb mRNA is upregulated in pericytes in the infarcted area (37), and Pdgfrb-positive pericytes migrate into the peri-infarct area after stroke, contributing to angiogenesis and vascular remodeling (25). In the current study, Pdgfrb expression was also upregulated by chronic hypoperfusion in the BCAS mice. Given that the PDGFRB is important for proliferation and migration of pericytes (55), PDGFRB may be increased in pericytes at a certain point, as compensation for hypoperfusion.

Figure 5. Effects of BMP4 on cultured oligodendrocyte precursor cells. A. Time-lapse images of primary OPCs in proliferation media with or without BMP4. Insets show enlarged images. Bars indicate 100 μm and 50 μm (insets). B. Representative triple-immunofluorescent images for Pdgfra and Gfap, and their merged images with DAPI for nuclear staining. Insets show enlarged images. Bars indicate 50 μm and μm (insets). C, D. Relative mRNA levels of Pdgfra and Gfap in the proliferation assay. (C) BMP4 decreases Pdgfra mRNA levels in a dose-dependent manner (P = 0.062 for 1 ng/ml and **P < 0.001 for 10 ng/ml), which is blocked by noggin (**P = 0.001). Treatment with noggin alone increases Pdgfra mRNA levels (*P = 0.015). (D) BMP4 significantly increases Gfap mRNA expression in a dose-dependent manner (**P = 0.004 for 1 ng/ml and **P < 0.001 for 10 ng/ml), which is blocked by noggin (**P < 0.001). Vertical bars represent mean ± SD. Abbreviations are as follows: Cont = control; Gapdh = glyceraldehyde-3-phosphate dehydrogenase.
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Unlike PDGFRB, ACTA2 expression was not different between the three groups. In terms of tumor angiogenesis, Acta2 served as a mature pericyte marker whereas Pdgfb served as a marker for progenitor perivascular cells with the ability to differentiate into pericytes and regulate vessel stability and vascular survival (50). Therefore, the differences in PDGFRB and ACTA2 expression may be the result of different stages of pericyte maturation; PDGFRB-positive pericytes in the white matter are immature pericytes during angiogenesis following chronic hypoperfusion, and PDGFRB-negative/ACTA2-positive pericytes are mature pericytes in relatively intact white matter. Because BMP4 treatment facilitated pericyte proliferation and Pdgfb mRNA expression, an increase in PDGFRB-positive pericytes in the SVD group might be induced by BMP4 in response to chronic hypoperfusion. BMP4 also facilitated proliferation and tube formation of endothelial cells, which is another key factor for maintenance of vascular integrity. These results suggest that BMP4 induces inherent compensatory angiogenesis against chronic hypoperfusion. Despite the putative compensatory responses, COL4A1-positive vessel density was not significantly different in SVD compared with AD or disease control patients in the cerebral cortex and white matter, suggesting that angiogenesis was not sufficient or compensatory enough to recruit blood vessels and mitigate cerebral hypoperfusion in SVD patients. Because time-specific induction of BMP4 and PDGFRB is necessary for successful angiogenesis, long-lasting hypoperfusion in the aged brain may block appropriate angiogenesis to restore regional blood flow (52).

With regard to myelination, BMP4 overexpression may not be preferable. BMP4 treatment has been shown to induce differentiation of oligodendroglial-astroglial progenitor cells into astrocytes (17, 27). Our results also showed that BMP4 strongly suppressed maturation of primary OPCs and converted OPCs into astrocytes in differentiation media. Compared with 1.2- to 2-fold changes in angiogenesis, the degree of inducing astrogligenesis (49) and suppressing oligodendrogenesis was exponential. Furthermore, when the OPCs were cultured in proliferation media which should maintain an OPC state, BMP4 treatment decreased Pdgfra mRNA expression in a dose-dependent manner. This finding was inconsistent with other studies that demonstrated no effect (16, 57) or even a positive effect (43) of BMP4 on OPC proliferation. However, recent studies have suggested that the effects of BMP4 on stem cell proliferation are dose-dependent; low dose BMP4 increases proliferation, but high dose BMP4 decreases proliferation of mesenchymal stem cells (53) or primordial germ cells (31). Considering that OPC proliferation is also affected by many factors, BMP4 may have dose-dependent and opposite effects on OPC proliferation. Consistent with the results of cell culture, the number of OPCs in SVZ is decreased with higher BMP4 expression of SVD cases with severe astrogligenesi. BCAS mice also showed astrogligenesis and hypomyelination as shown previously (48). As noggin, a receptor antagonist of BMP4, suppressed astrogligenesis and induced oligodendrogenesis in vitro, both high and low doses of noggin cCV decreased the number of astrocytes and high dose of noggin cCV increased the number of oligodendrocytes.

The close relationship between BMP4 and myelin attenuation is also supported by recent studies (3, 15, 26, 43). These data strongly suggest that BMPs are involved in oligovascular pathologies in the brain. Considering a potential anatomical and functional interaction between pericytes and OPCs in the capillary of cerebral white matter, the pericyte-induced BMP4 might directly affect OPC lineage. Because many factors should be expressed in a coordinated manner after cerebral hypoperfusion, time-specific control of BMP expression should be further investigated to protect oligovascular units in ischemic demyelinating diseases.

The main limitation of this study was the lack of clinical and neuropsychometric information on the postmortem brains used. This meant that we could not directly relate our pathological findings to antemortem hypoperfusive and cognitive status. In addition, the lack of comparison of the SVD cases with controls without neurodegenerative pathologies was another limitation considering that PDGFRB positive pericytes may also degenerate in some neurodegenerative diseases (46, 55, 56, 59). The third limitation is that a histopathological study using postmortem human brains does not always uncover the entire cell process that occurs in the ageing
human brain. Although the process was at least partially modeled by both cultured pericytes under OGD and mice subjected to BCAS, future studies should focus on examining dynamic changes of pericyte disruption and BMP4 values in the disease course of SVD using neuroimaging and biomarkers.

In summary, our study suggests that during chronic hypoperfusion akin to that in SVD, BMP4 is generated in pericytes predominantly in the white matter, providing putative compensatory angiogenesis. However, BMP4 also aggravates white matter damage by inducing astrogliogenesis at the expense of OPC proliferation and maturation. Based on our observations, we propose a putative scheme how these cellular processes could occur (Figure 7). This may explain why white matter is particularly vulnerable to chronic hypoperfusion.

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AUTHOR CONTRIBUTIONS

M.T.U. study design, acquisition and analysis of data, and drafting the manuscript and figures; M.I. study conception and design, handling funding and supervising portions of the study; T.M. design of the study, and supervising and making critical revision of the manuscript for important intellectual content; T.N. data acquisition and analysis, and supervising and making critical revision of the manuscript for important intellectual content; S.K. data acquisition and analysis; K.U. and A.K. supervising and making critical revision of the manuscript for important intellectual content.; R.N.K. and R.T. handling funding, supervising and making critical revision of the manuscript for important intellectual content.

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SUPPORTING INFORMATION

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

Figure S1. Expression of TGFβ1, BMP2, BMP6, BMP7, and BMP9 in white matter. Representative images of TGFβ1, BMP2, BMP6, BMP7, and BMP9 staining in the WM of control, AD, and SVD, respectively. Insets show TGFβ1, BMP2, BMP6, BMP7, or BMP9-positive activated astrocytes with large cell bodies (right panels), TGFβ1 or BMP7-positive pericytes (left panels in TGFβ1 and BMP7), and BMP2-positive endothelial cells (left panels in BMP2), respectively. Bars indicate 100 μm and 10 μm (insets).

Figure S2. Oligodendrocyte precursor cells in the subventricular zone. Representative images for PDGFRα-positive cells surrounding arterioles, venules and capillaries in the subventricular zone. Bars indicate 25 μm.

Figure S3. Effects of BMP4 on oligodendrocyte maturation in the OPC differentiation assay. (A) Time-lapse images of primary OPCs in differentiation media with or without BMP4. Insets show enlarged images. Bars indicate 100 μm and 50 μm (insets). (B) Representative triple-immunofluorescent images for Mbp and Gfap, and their merged images with DAPI for nuclear staining. Insets show enlarged images. Bars indicate 100 μm and 15 μm (insets). (C, D) Relative mRNA levels of Mbp and Gfap in the differentiation assay. Mbp mRNA levels decrease in differentiation media containing BMP4 (**P<0.001 for 1 and 10 ng/ml) in a dose-dependent manner; the effect of BMP4 is reversed by noggin (**P<0.001). Treatment with noggin alone increases Mbp (**P<0.001) (C). BMP4 significantly increases Gfap mRNA expression (**P<0.001 for 1 and 10 ng/ml) in a dose-dependent manner; the effect of BMP4 is canceled out by noggin (**P<0.001) (D). Vertical bars represent mean ± SD. Abbreviation is as follow: Cont, control; Gapdh, glyceraldehyde-3-phosphate dehydrogenase.

Figure S4. Original gel pictures for western blots. The figure shows uncropped western blots displayed in Fig. 6C. (A) Bmp4 precursor and pro-Bmp4 expressions are increased in BCAS mice compared with sham controls. Pdgfrb (B), but not Acta2 (C), expression is increased in BCAS mice compared with sham controls. Each band of Bmp4, Pdgfrb, and Acta2 is normalized to Tubg1. (D) Gfap expressions are increased in BCAS mice compared with sham controls, which are suppressed by noggin cICV (500 ng/day and 1000 ng/day). (E) Olig2 expressions are decreased in BCAS mice compared with sham controls, which are ameliorated by a high dose of noggin cICV (1000 ng/day). (F) Each band of Gfap and Olig2 is normalized to the band at 45 kDa stained with ponceau S. Asterisks show unknown bands. Abbreviations are as follows: B, BCAS; N500, Noggin 500 ng/ day; N1000, Noggin 1000 ng/day; S, Sham operation.

Table S1. Primary antibodies used for immunohistochemistry of human brain tissue.

Table S2. Primers used for RT-PCR.

Table S3. Primary antibodies used for immunocytochemistry.

Table S4. Primary antibodies used for immunohistochemistry of mouse brain tissue.

Table S5. Primary antibodies used for western blot.