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雑誌

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Takaoka M, Uemura S, Kawata H, Imagawa K, Takeda Y, Nakatani K, Naya N, Horii M, Yamano S, Miyamoto Y, Yoshimasa Y , Saito Y.	Inflammatory Response to a cute myocardial infarction augments neointimal hyperplasia after vascular injury in a remote artery	Arterioscler T hromb Vasc Biol.	26	2083-2089	2006
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Makino H, Miyamoto Y, Sawai K, Mori K, Mukoyama M, Nakao K, Yoshimasa Y , Suga S.	Altered gene expressio n related to glomerulo genesis and podocyte s tructure in early diabet ic nephropathy of db/d b mice and its restorat ion by pioglitazone.	<i>Diabetes</i>	55	2747-2756.	2006

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Adrenomedullin/Cyclic AMP Pathway Induces Notch Activation and Differentiation of Arterial Endothelial Cells From Vascular Progenitors

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Objective—The acquisition of arterial or venous identity is highlighted in vascular development. Previously, we have reported an embryonic stem (ES) cell differentiation system that exhibits early vascular development using vascular endothelial growth factor (VEGF) receptor-2 (VEGFR2)-positive cells as common vascular progenitors. In this study, we constructively induced differentiation of arterial and venous endothelial cells (ECs) *in vitro* to elucidate molecular mechanisms of arterial-venous specification.

Methods and Results—ECs were induced from VEGFR2⁺ progenitor cells with various conditions. VEGF was essential to induce ECs. Addition of 8bromo-cAMP or adrenomedullin (AM), an endogenous ligand-elevating cAMP, enhanced VEGF-induced EC differentiation. Whereas VEGF alone mainly induced venous ECs, 8bromo-cAMP (or AM) with VEGF supported substantial induction of arterial ECs. Stimulation of cAMP pathway induced Notch signal activation in ECs. The arterializing effect of VEGF and cAMP was abolished in recombination recognition sequence binding protein at the J κ site deficient ES cells lacking Notch signal activation or in ES cells treated with γ -secretase inhibitor. Nevertheless, forced Notch activation by the constitutively active Notch1 alone did not induce arterial ECs.

Conclusions—Adrenomedullin/cAMP is a novel signaling pathway to activate Notch signaling in differentiating ECs. Coordinated signaling of VEGF, Notch, and cAMP is required to induce arterial ECs from vascular progenitors. (*Arterioscler Thromb Vasc Biol.* 2006;26:1977-1984.)

Key Words: angiogenesis ■ developmental biology ■ embryonic stem cells ■ endothelium ■ vascular biology

Vascular formation is a complicated but well-organized process that involves sprouting, branching, and differential growth of vessels from the primary plexus or existing vessels into a functioning circulation system.¹ During the process, vascular cell specification proceeds in an inseparably coordinated manner.² A transmembrane ligand, ephrinB2, and its receptor, the tyrosine kinase EphB4, are reported as molecular markers for arterial and venous endothelial cells (ECs), respectively.^{3,4} Recently, various molecular markers specific for arterial ECs have been documented such as Delta-like 4 (Dll4), Brnx, Notch1, Activin receptor-like kinase 1 (Alk1), and others.^{5,6} These findings enable the investigation of endothelial specification processes at the cellular and molecular levels being independent of the context of vessel location within the body plan.

The Notch pathway has been highlighted in arterial-venous specification.^{7,8} Notch target genes, Hairy and Enhancer-of-

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split-related basic helix-loop-helix transcription factors, such as *gri* (gridlock) in zebrafish, or *Hey1* and *2* in mammals, are required for arterial vascular development.^{9,10} Arterial-venous specification mechanisms in zebrafish were further demonstrated to be a regulatory signaling cascade of sonic hedgehog-vascular endothelial growth factor (VEGF)-Notch-ephrinB2.⁵ The molecular machinery for arterial-venous specification in mammals, however, is still undergoing investigation.

cAMP is a ubiquitous second messenger produced in cells and is involved in various biological phenomena including cell growth and differentiation.¹¹ Nevertheless, little has been reported for the role of cAMP signaling in vascular development. Adrenomedullin (AM) is a multifunctional polypeptide that was originally isolated from human pheochromocytoma.¹² AM exerts its function by increasing the levels of

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intracellular cAMP through the binding to its receptor complex, calcitonin receptor-like receptor (CRLR), and receptor activity modifying proteins (RAMP)-2 or RAMP-3.¹³ Targeted null mutation of the *AM* gene shows embryonic lethality¹⁴ with aberrant vascular formation and hemorrhage,¹⁵ or extreme hydrops fetalis and cardiovascular abnormalities, including underdeveloped arterial walls,¹⁶ inferring the significance of *AM*/cAMP signaling in vascular development.

Pluripotent embryonic stem (ES) cells are potent materials for both regenerative therapeutic approaches and developmental research. We have developed a novel ES cell differentiation system devoid of embryoid body formation or feeder cells that exhibits early vascular development using VEGF receptor-2 (VEGFR2)-positive cells as common progenitors for vascular cells.^{17,18} We demonstrated that ES cell-derived VEGFR2⁺ cells can differentiate into both ECs and mural cells (MCs) (pericytes and vascular smooth muscle cells) and form mature vascular-like structures *in vitro*.¹⁸ Moreover, transplantation of induced vascular cells can augment the blood flow in tumor angiogenesis.¹⁹ Our ES-derived VEGFR2⁺ cell differentiation system can recapitulate the vascular development processes and dissect the cellular and molecular mechanisms of each developmental step including endothelial differentiation and specification.

In this study, we aimed to specifically induce arterial and venous ECs and elucidate the mechanisms of arterial-venous specification using our ES cell differentiation system. We successfully induced arterial and venous ECs and demonstrated that the *AM*/cAMP pathway is another indispensable signaling pathway in EC differentiation and arterial specification in conjunction with VEGF and Notch by reconstructing the arterial EC differentiation process *in vitro*. Our constructive approach using this ES cell system provides a novel understanding of the cellular and molecular mechanisms of vascular developmental processes.

Methods

Antibodies

Monoclonal antibodies for murine E-cadherin (ECCD2), murine VEGFR2 (AVAS12), and murine VE-cadherin (VECD1) were described previously.¹⁸ Monoclonal antibodies for murine CD31 and CXCR4 were purchased from Pharmingen (San Diego, Calif). MoAb for murine alpha smooth muscle actin (SMA) 1A4 and human estrogen receptor- α (ER α) (F-10) antibody were from Sigma (St Louis, Mo) and Santa Cruz Biotechnology (Santa Cruz, Calif), respectively. Cleaved Notch1 antibody was from Cell Signaling Technology (Beverly, Mass).

Cell Culture

Induction of differentiation of an ES cell line, CCE (gift from Dr Evans), were performed using differentiation medium (alpha minimal essential medium; Gibco, Grand Island, NY) supplemented with 10% fetal calf serum (Equitech-Bio, Kerrville, Tex) and 5×10^{-5} mol/L 2-mercaptoethanol (Gibco) and VEGF165 (R&D System, Minneapolis, Minn) as previously described.^{17,18} Other chemicals, rat *AM* (Peptide Institute, Inc, Osaka, Japan), 8-bromoadenosine-3':5'-cyclic monophosphate sodium salt (8bromo-cAMP) (Nacalai Tesque, Kyoto, Japan), 8-bromoguanosine-3':5'-cyclic monophosphate sodium salt (8bromo-cGMP) (Nacalai Tesque), 3-isobutyl-1-methyl-xanthine (IBMX) (Nacalai Tesque), or γ -secretase inhibitor IX, DAPT (Calbiochem, San Diego, Calif), and iloprost (Cayman

Chemical, Ann Arbor, Mich) were occasionally added to VEGFR2⁺ cell culture.

The recombination recognition sequence binding protein at the *Jk* site (RBP-J^{+/+}), RBP-J^{-/-} and RBP-J^{-/-} D3 ES cell lines have been described previously.²⁰ The ES cell line NERT^{Δ0-7}²¹ was generated by stable introduction of CAG promoter-driven cDNA encoding a fusion protein of a constitutively active part of the intracellular domain of mouse Notch1 and a tamoxifen-sensitive mutant of the hormone binding domain of the human estrogen receptor α (NERT)²² into EB5 ES cells (gift from Dr Niwa). To induce Notch activation, 4-hydroxytamoxifen (OHT) (50 to 500 nmol/L) (Sigma) was added to NERT^{Δ0-7} cell-derived VEGFR2⁺ cells 12 hours after the plating. NERT^{Δ0-7}/Hes-green fluorescent protein (GFP) cells were generated by stable introduction of Hes promoter-driven enhanced GFP (EGFP) gene²³ (gift from Dr Kageyama) into NERT^{Δ0-7} cells.

Flowcytometry and Cell Sorting

Fluorescence-activated cell sorting (FACS) of ES cells was performed as previously described.^{17,18}

Immunocytochemistry

Immunostaining for cultured cells was performed as described.^{18,24} Double immunofluorescent staining for CD31 and ER α was performed using anti-ER α antibody (1:50) and anti-CD31 antibody (1:300) as first antibodies, followed by second antibodies, Alexa Fluor 546-conjugated goat anti-rat IgG (1:500) and Alexa Fluor 488-conjugated goat anti-mouse IgG (1:500) (Molecular Probes, Eugene, Ore). For double staining for EphrinB2 and CD31, the fixed culture slides were incubated with EphB4-human immunoglobulin Fc portion chimeric protein (EphB4-Fc) (1:50; R&D system), followed by peroxidase-conjugated goat IgG fraction to human IgG Fc (1:500; ICN Biomedicals, Inc, Aurora, Ohio). TSA Biotin system (Tyramid signal amplification; PerkinElmer Life Science, Boston, Mass) was used for amplification of the signal for EphB4-Fc staining. EphrinB2⁺ cells were visualized by using streptavidin-Alexa Fluor488-conjugate (Molecular Probes). Phycocerythrin-conjugated anti-CD31 antibody (Pharmingen) and DAPI (Molecular Probes) were added together with streptavidin-conjugated alexa 488. Cleaved intracellular domain of Notch (NICD) staining was performed using TSA Biotin System (PerkinElmer) with cleaved Notch1 antibody (1:300), followed by peroxidase-labeled anti-rabbit IgH (1:250; Vector Laboratories, Burlingame, Calif).

Single-Cell Analysis

Single-cell sorting of VEGFR2⁺ cells using 96-well dishes was performed as previously described.¹⁸ Colonies were stained for EphrinB2 using EphB4-Fc by TSA kit with streptavidin-conjugated horseradish peroxidase, followed by addition of phycocerythrin-conjugated anti-CD31 antibody and DAPI. Numbers of colonies including CD31⁺ cells (EC-including), colonies including EphrinB2⁺ cells (arterial EC-including), and EphrinB2⁺ arterial EC numbers in each arterial EC-including colonies, as well as the total number of colonies that appeared were counted. 1692 VEGFR2⁺ cells were cultured with VEGF alone, and 1128 cells were cultured with VEGF and 8bromo-cAMP. Total colony numbers in every 100 sequential wells, EC-including or arterial EC-including colony numbers in every 10 sequential colonies that appeared, and the arterial EC number in each arterial EC-including colony were statistically evaluated.

Measurement of Intracellular cAMP

After 3 days culture of VEGFR2⁺ cells (2 to 10×10^5 cells), cells were harvested and counted. Intracellular cAMP concentration in total harvested cells was evaluated using cAMP Biotrak Enzyme Immunoassay system kit (Amersham Bioscience). Concentration was normalized by cell number.

In Situ Hybridization

In situ hybridization for CXCR4 was performed as previously described.²⁵

Reverse-Transcription Polymerase Chain Reaction Amplification

Total RNA was isolated from sorted VE-cadherin⁺ ECs induced by VEGF alone, or 8bromo-cAMP and VEGF treatment, using ISOGEN (Nippon Gene, Toyama, Japan). The reverse-transcription polymerase chain reaction was performed as described²⁴ using indicated primers (supplemental Table I, available online at <http://atvb.ahajournals.org>).

Statistical Analysis

Statistical analysis of the data was performed using Student *t* test. *P* < 0.05 was considered significant.

Results

We first examined the effects of AM and cAMP on EC differentiation from ES cell-derived VEGFR2⁺ progenitor cells. VEGFR2⁺ cells were sorted by FACS and re-cultured for 3 days on type IV collagen-coated dishes in differentiation medium (see Methods) with VEGF (50 ng/mL) and other factors. Double immunostaining of induced cells with an EC marker, CD31, and a MC marker, SMA, revealed that VEGF treatment selectively induced both CD31⁺ ECs and SMA⁺ MCs from VEGFR2⁺ cells as previously reported¹⁸ (Figure 1A). Simultaneous stimulation of cAMP signaling in the presence of VEGF substantially enhanced EC induction from VEGFR2⁺ cells (Figure 1B to 1D). VEGF together with 0.5 mmol/L 8bromo-cAMP resulted in substantial induction of ECs (Figure 1D), whereas 8bromo-cAMP treatment alone exerted almost no effect (data not shown). Another cyclic monophosphate analog, 8bromo-cGMP, showed no effect on VEGF-induced EC induction (data not shown). Addition of 10⁻⁶ mol/L AM also enhanced VEGF-stimulated EC induction, but to a lesser extent than 8bromo-cAMP (Figure 1B). Enhancement of the effect of AM by the simultaneous administration of a phosphodiesterase inhibitor, IBMX, revealed comparable EC induction with 8bromo-cAMP (Figure 1C). We quantitatively evaluated the EC-inducing effects of AM and 8bromo-cAMP using flow cytometry. VEGF treatment induced ECs to ≈30% of total cells. AM increased VEGF-induced ECs up to ≈50%. AM with IBMX or 8bromo-cAMP showed efficient induction of ECs to ≈70% of total cells (Figure 1E). Intracellular concentration of cAMP in the differentiating cells was significantly increased by AM with VEGF (667.6 fmol ± 215.1/10⁶ cells; n=6; *P* < 0.01 versus VEGF alone), or AM and IBMX with VEGF (1142 fmol ± 270.1/10⁶ cells; n=6; *P* < 0.001 versus VEGF alone) than that with VEGF alone (372.2 fmol ± 58.5/10⁶ cells; n=6), and was comparable or lower level with those observed in previous reports using human umbilical vein ECs.²⁶ These results indicated that the AM/cAMP pathway specifically and synergistically enhances the effect of VEGF on EC differentiation from VEGFR2⁺ progenitor cells.

Next, we investigated the features of induced ECs with AM/cAMP treatment with regard to arterial-venous diversity. Arterial ECs were evaluated by ephrinB2 expression, an arterial EC marker, detected by the binding of EphB4-Fc.²⁷

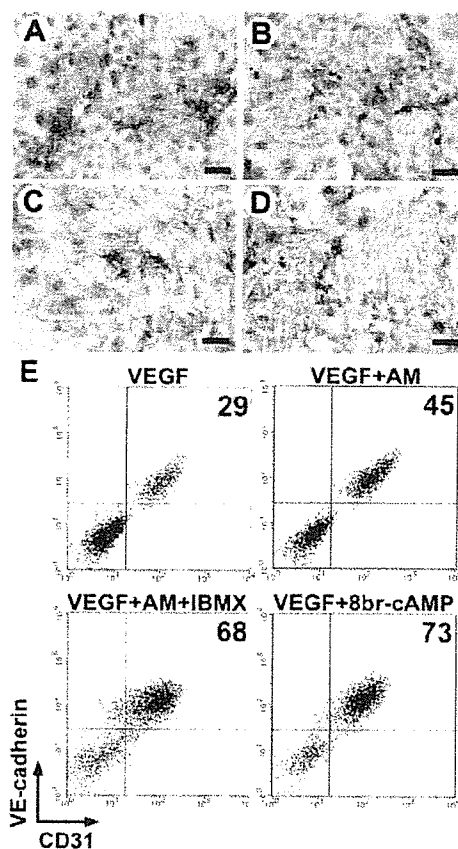


Figure 1. The effect of AM and cAMP on EC induction from VEGFR2⁺ cells. A to D, Double immunostaining of induced ECs and MCs with an EC marker CD31 (purple) and MC marker SMA (brown) after 3 days of culture of VEGFR2⁺ cells on type IV collagen-coated dishes in various conditions. A, VEGF treatment alone (50 ng/mL). CD31⁺ EC sheets and SMA⁺ MCs appear. B, VEGF with 10⁻⁶ mol/L AM. A slight increase of ECs is observed. C, VEGF with 10⁻⁶ mol/L AM and 10⁻⁴ mol/L IBMX. D, VEGF with 0.5 mmol/L 8bromo-cAMP. Remarkable EC induction occurs. Scale bars: 100 μm. E, Flow cytometry of induced cells from VEGFR2⁺ cells with endothelial markers VE-cadherin and CD31. Left upper panel, VEGF treatment alone (50 ng/mL). Right upper panel, VEGF with 10⁻⁶ mol/L AM. Left lower panel, VEGF with 10⁻⁶ mol/L AM and 10⁻⁴ mol/L IBMX. Right lower panel, VEGF with 0.5 mmol/L 8bromo-cAMP. Percentages of VE-cadherin⁺/CD31⁺ ECs of total VEGFR2⁺ cell-derived cells are indicated.

We double-immunostained ECs using anti-CD31 antibody and EphB4-Fc (Figure 2A to 2D). With VEGF treatment alone, very few ephrinB2⁺ arterial ECs were observed among the ECs that appeared, indicating that venous ECs were mainly induced in this condition (Figure 2A). Surprisingly, remarkable appearance of ephrinB2⁺ ECs was clearly observed by the stimulation of cAMP pathway. That is, addition of AM induced ephrinB2⁺ EC appearance (Figure 2B). AM with IBMX, or 8bromo-cAMP together with VEGF, showed substantial induction of ephrinB2⁺ ECs (Figure 2C and 2D). Messenger RNA expression of arterial EC markers, ephrinB2, Dll4, Notch1, Notch4, Alk1, and neuropilin1 (NRP1) were increased in 8bromo-cAMP and VEGF-treated ECs (Figure 2E). In contrast, venous EC markers, COUP-TFII transcription factor²⁸ and NRP2²⁹ mRNA were decreased by

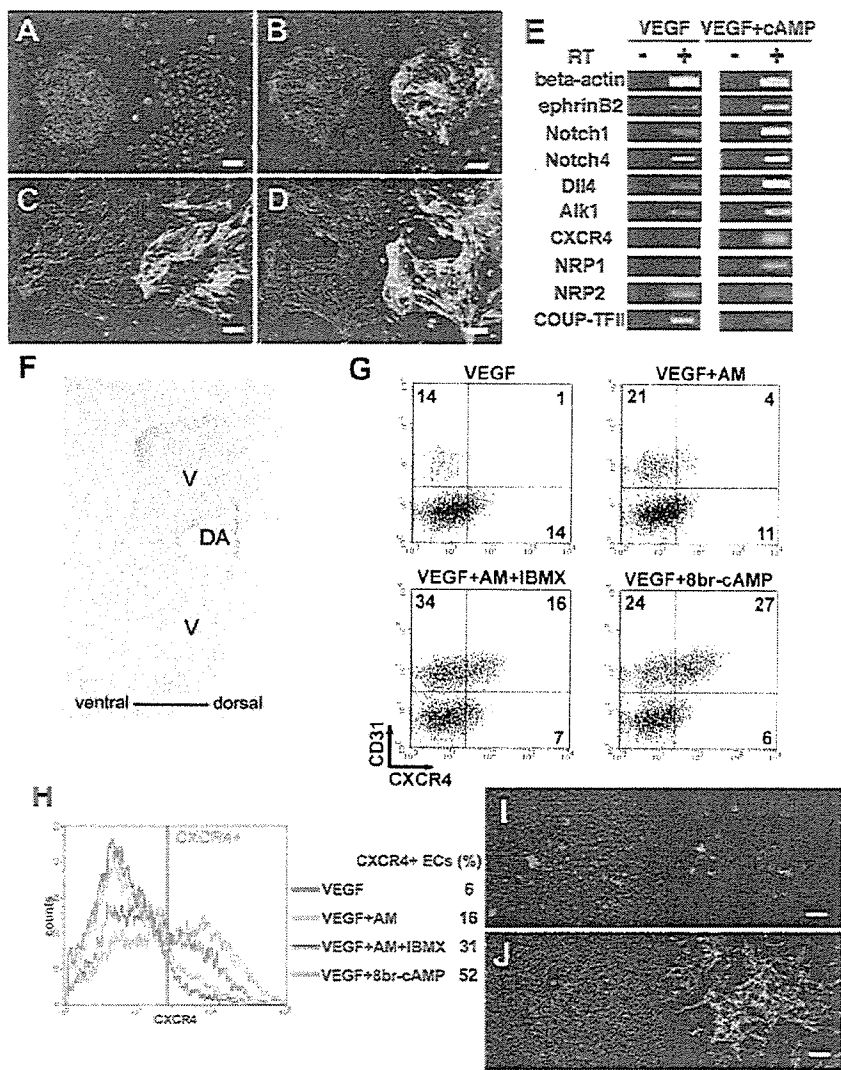


Figure 2. The effect of AM and cAMP on arterial EC induction from VEGFR2⁺ cells. A to D, Double fluorescent staining for CD31 and ephrinB2 after 3 days of culture of VEGFR2⁺ cells. Left panels, CD31 (pan-ECs, red) and DAPI (blue). Right panels, EphB4-Fc (ephrinB2⁺ arterial ECs, green) and DAPI (blue). A, VEGF treatment alone (50 ng/mL). B, VEGF with 10⁻⁶ mol/L AM. C, VEGF with 10⁻⁶ mol/L AM and 10⁻⁴ mol/L IBMX. D, VEGF with 0.5 mmol/L 8bromo-cAMP. Scale bars: 100 μm. E, Reverse-transcription polymerase chain reaction showing mRNA expression of arterial markers (ephrinB2, Notch1, Notch4, Dll4, Alk1, CXCR4, and NRP1) and venous marker (NRP2 and COUP-TFII) in purified ECs induced by VEGF treatment alone or VEGF and 8bromo-cAMP treatment. F, Aortic EC-specific expression of CXCR4 (purple) by in situ hybridization of the isolated aorta-gonadomesonephros (AGM) region in E11.5 mouse embryo. DA indicates dorsal aorta; V, Cardinal veins. G, Flow cytometry for CD31 and CXCR4 expression. Left upper panel, VEGF treatment alone (50 ng/mL). Right upper panel, VEGF with 10⁻⁶ mol/L AM. Left lower panel, VEGF with 10⁻⁶ mol/L AM and 10⁻⁴ mol/L IBMX. Right lower panel, VEGF with 0.5 mmol/L 8bromo-cAMP. H, Expression profile of CXCR4 in CD31⁺ ECs by flow cytometry. VEGF treatment alone (blue line), VEGF with 10⁻⁶ mol/L AM (green line), VEGF with 10⁻⁶ mol/L AM and 10⁻⁴ mol/L IBMX (red line), and VEGF with 0.5 mmol/L 8bromo-cAMP (orange line) are shown. Percentages of CXCR4⁺ arterial ECs in total ECs are indicated. I and J, Gross appearance of ephrinB2⁺ arterial EC induction from VEGFR2⁺ cells (plated at 2 × 10⁶ cells/cm²). Left panels, DAPI (blue). Right panels, EphB4-Fc (ephrinB2⁺ arterial ECs, green). I, VEGF treatment alone (50 ng/mL). J, VEGF with 0.5 mmol/L 8bromo-cAMP. Increase in cell number (DAPI) and substantial arterial EC induction were observed. Scale bars: 400 μm.

8bromo-cAMP and VEGF treatment (Figure 2E). These results indicated that stimulation of cAMP pathway induces arterial ECs.

We further attempted to quantitatively evaluate arterial EC induction at the cellular level. CXCR4, a 7-transmembrane G-protein-coupled receptor, is the receptor of CXCL12 (also known as stromal cell-derived factor-1). Recently, CXCR4 has been reported to be expressed in ECs in the superior mesenteric artery, but not in the superior mesenteric vein, and involved in the formation of arteries in the gastrointestinal tract.^{25,30} We examined CXCR4 expression in the mouse embryo by in situ hybridization and found that CXCR4 was detected in ECs of the dorsal aorta but not of cardinal veins in aorta-gonadomesonephros (AGM) region of E11.5 embryos (Figure 2F). In addition, mRNA expression of CXCR4 was increased in 8bromo-cAMP and VEGF-treated ECs together with other arterial EC markers (Figure 2E), indicating that CXCR4 is another arterial EC marker. FACS analysis using an anti-CXCR4 antibody successfully quantified arterial EC induction by AM or 8bromo-cAMP treatment. Most

of ECs induced by VEGF treatment alone (>90% to 95%) were negative for CXCR4. CXCR4⁺/CD31⁺ arterial ECs were induced in the presence of AM together with VEGF. Addition of AM with IBMX, or 8bromo-cAMP further increased CXCR4⁺/CD31⁺ arterial EC appearance (Figure 2G). Overall, 8bromo-cAMP and VEGF treatment induced ≈5- to 10-fold more CXCR4⁺ arterial ECs compared with VEGF treatment alone. AM with VEGF treatment showed slight effect on the arterial EC induction. Simultaneous administration of AM and IBMX with VEGF enhanced the arterializing effect of AM (Figure 2H). These results indicated that cAMP signaling mainly contributes to the arterial EC induction. The maximum percentage of arterial ECs within total ECs was increased to ≈60% by 8bromo-cAMP and VEGF (Figure 3F). Addition of 8bromo-cAMP with VEGF led to an increase in total cell number, total EC number, and arterial EC percentage, resulting in ≈70-times increment of induced arterial EC number than those by VEGF alone (Figure 2I and 2J). Higher doses of VEGF (100 to 200 ng/mL) alone or 8bromo-cAMP (0.5 mmol/L) with VEGF

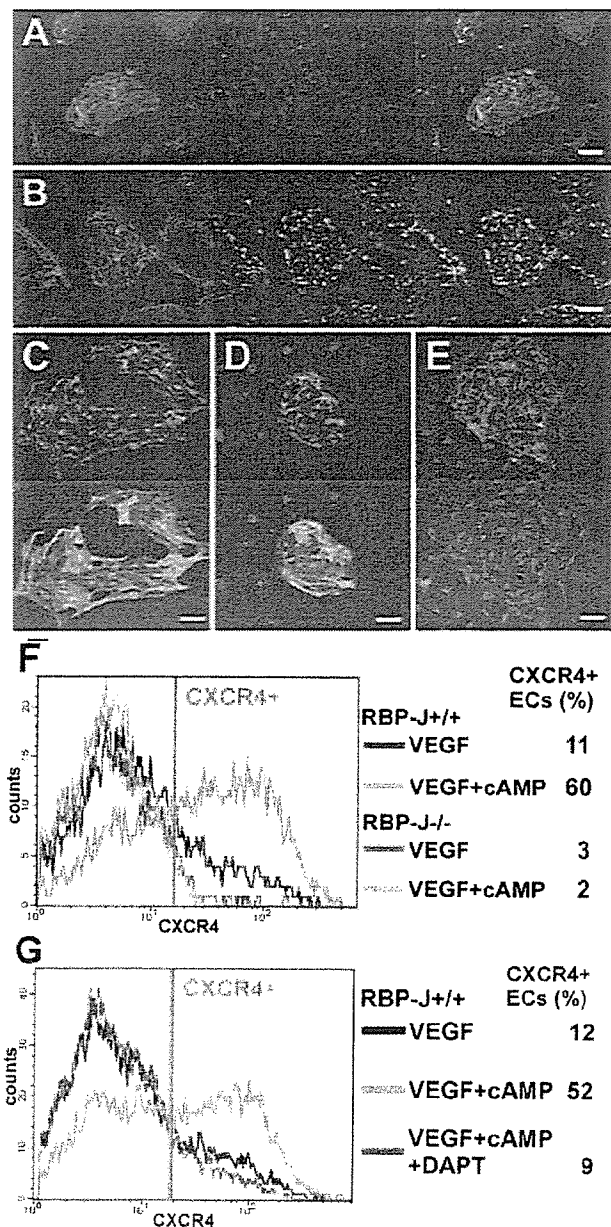


Figure 3. Essential role of Notch signaling in arterial EC induction. A and B, Double fluorescent staining of cleaved Notch intracellular domain (NICD) and CD31 for induced ECs. Left panels, CD31 (pan-ECs, red). Middle panels, Cleaved NICD (green). Right panels, Merged image. A, VEGF treatment alone (50 ng/mL). B, VEGF with 0.5 mmol/L 8bromo-cAMP. Scale Bars: 200 μ m. C to E, Double-fluorescent staining of CD31 and ephrinB2 for ECs induced by VEGF with 8bromo-cAMP using RBP-J-deficient ES cells. Upper panels, CD31 (pan-ECs, red) and DAPI (blue). Lower panels, EphB4-Fc (ephrinB2⁺ arterial ECs, green) and DAPI (blue). C, RBP-J^{+/+} ES cells. D, RBP-J^{+/-} ES cells. E, RBP-J^{-/-} ES cells. Scale bars: 100 μ m. F and G, Expression profile of CXCR4 in CD31⁺ ECs. F, Blue and green lines: RBP-J^{+/+} cells. VEGF treatment alone (blue line), VEGF with 0.5 mmol/L 8bromo-cAMP (green line). Red and orange lines: RBP-J^{-/-} cells. VEGF alone (red line), VEGF with 0.5 mmol/L 8bromo-cAMP (orange line). Percentages of CXCR4⁺ arterial ECs in total ECs are indicated. G, RBP-J^{+/+} cells. VEGF treatment alone (blue line), VEGF with 0.5 mmol/L 8bromo-cAMP (green line). VEGF with 8bromo-cAMP and 2.5 μ mol/L DAPT (red line).

Single-Cell Analysis of VEGFR2⁺ Cell Culture

	VEGF Alone	VEGF With 8bromo-cAMP
Total colony, n (per every 100 sequential wells)	5.62 ± 1.74 (n=16)	16.0 ± 6.06 (n=11)*
EC-including colony, n (per every 10 sequential colonies)	3.40 ± 2.20 (n=15)	7.00 ± 1.70 (n=19)*
AEC-including colony, n (per every 10 sequential colonies)	1.27 ± 1.10 (n=15)	3.63 ± 1.30 (n=19)*
AEC number (per each AEC-including colony)	1.69 ± 0.87 (n=16)	4.51 ± 2.77 (n=76)*

*P < 0.01 vs VEGF alone.
AEC indicates arterial endothelial cell; EC, endothelial cell.

treatment did not show arterial EC induction. Administration of iloprost (10⁻⁷ to 10⁻⁵ mol/L), an analogue of prostaglandin-I2 that elevates intracellular cAMP in mature ECs, showed almost no arterial inducing effect even with VEGF treatment (data not shown). These results indicated that AM/cAMP signaling is a novel potent and specific inducer of arterial ECs from vascular progenitor cells.

To further evaluate the mechanism of AM/cAMP-stimulated arterial EC induction, we performed single-cell culture of VEGFR2⁺ cells. Colonies obtained from single VEGFR2⁺ cells were counted and evaluated by staining for CD31, ephrinB2, and DAPI (Table). VEGF and 8bromo-cAMP treatment significantly increased the total number of colonies that appeared, number of EC-including colonies, and arterial EC-including colonies in appeared colonies, and arterial EC numbers in each arterial EC-including colony than VEGF alone. These results suggest that cAMP increased survival of VEGFR2⁺ progenitor cells, differentiation of ECs and arterial ECs from progenitor cells that survived, and proliferation of arterial ECs. cAMP, thus, should be involved in multi steps of arterial EC differentiation processes.

We then examined the role of Notch signaling in arterial EC induction in this system. Activation of Notch on ligand binding is accompanied by proteolytic processing that releases intracellular domain of Notch (NICD) from the membrane. The NICD then translocates into the nucleus and associates with RBP-J, a DNA-binding protein, to form a transcriptional activator, which turns on transcription of a set of target genes.³¹ First, we examined Notch activation by cAMP treatment with immunostaining of cleaved NICD. Whereas Notch signal was not activated in most of ECs induced by VEGF alone (Figure 3A), administration of 8bromo-cAMP together with VEGF clearly induced nuclear localization of cleaved NICD in ECs, indicating that stimulation of cAMP pathway can activate Notch signaling in differentiating ECs (Figure 3B). cAMP is, thus, found to be a novel signaling pathway that interacts with and activates Notch signaling in EC lineages. Then, we performed a loss-of-function study using RBP-J-deficient ES cells that lack Notch signaling activation.²⁰ VEGFR2⁺ cells derived

from RBP-J^{+/+}, RBP-J^{+/-}, or RBP-J^{-/-} ES cells were sorted and re-cultured with VEGF in the presence of 8bromo-cAMP. Arterial EC induction observed in RBP-J^{+/+} (Figure 3C) or RBP-J^{+/-} ES cells (Figure 3D) was completely abolished in RBP-J^{-/-} ES cells (Figure 3E). FACS analysis using CXCR4 further demonstrated that induction of CXCR4⁺ arterial ECs observed in RBP-J^{+/+} was completely abolished in RBP-J^{-/-} ES cells (Figure 3F). Similarly, administration of γ -secretase inhibitor, DAPT (2.5 μ mol/L), which inhibits proteolytic processing of Notch to activate its signaling, to VEGFR2⁺ cell culture also completely blocked the arterial EC induction (Figure 3G). These results indicate that Notch signaling is essential for arterial EC induction in this ES cell system, and correlates with previous reports in zebrafish^{32,33} and mouse^{34,35} genetic animal models.

Next, we examined the effect of a gain-of-function of Notch in arterial EC induction. We used an ES cell line NERT ^{Δ OP-7},²¹ in which signaling of the activated intracellular domain of murine Notch1 can be regulated using an OHT-inducible system.²² NERT ^{Δ OP-7} ES cell-derived VEGFR2⁺ cells were sorted and re-cultured with VEGF in the presence or absence of OHT. In the absence of OHT, NERT protein was located mainly in the cytoplasm of induced CD31⁺ ECs and other cell types (supplemental Figure IA, available online at <http://atvb.ahajournals.org>). After addition of OHT, NERT protein translocated to the nucleus (supplemental Figure IB). Notch signal activation in VEGF-induced ECs was evaluated by FACS using NERT ^{Δ OP-7}/Hes-GFP cells carrying HES promoter-driven GFP gene (supplemental Figure IC). Addition of 8bromo-cAMP induced endogenous Notch activation in ECs, correlating with our previous results shown in Figure 3A and 3B. OHT treatment showed stronger Notch signal activation through NERT protein than 8bromo-cAMP treatment. Simultaneous stimulation by 8bromo-cAMP and OHT additionally enhanced Notch activation in induced ECs. These results indicate that NERT ^{Δ OP-7} cell system can successfully induce Notch signal activation in differentiating ES cells. NERT ^{Δ OP-7} cell-derived ECs induced by VEGF alone were negative for ephrinB2 (Figure 4A). Unexpectedly, hardly any arterial ECs appeared after Notch activation with OHT, even when co-stimulated with VEGF (Figure 4B). Although ephrin-B2⁺ arterial ECs were successfully induced by VEGF with 8bromo-cAMP (Figure 4C), no apparent effect of OHT was observed on the cAMP-stimulated arterial EC induction with ephrinB2 staining (Figure 4D). FACS analysis further demonstrated that activation of Notch signaling by OHT failed to induce CXCR4⁺ arterial ECs and, moreover, activation of Notch signaling with OHT did not affect, or often reduced, cAMP-induced CXCR4⁺ arterial EC induction (Figure 4E). These results indicate that Notch signal is not sufficient or at least aberrant activation of Notch is not beneficial, for arterial EC induction. This is compatible with the previous *in vivo* study using activated Notch4-transgenic mice in that activation of Notch signaling in embryonic endothelium led to disorganized vascular networks but did not document arterial induction.³⁶

Taken together, VEGF appears essential for EC differentiation from VEGFR2⁺ cells, and venous ECs can be induced by VEGF alone. For arterial EC induction, however, VEGF

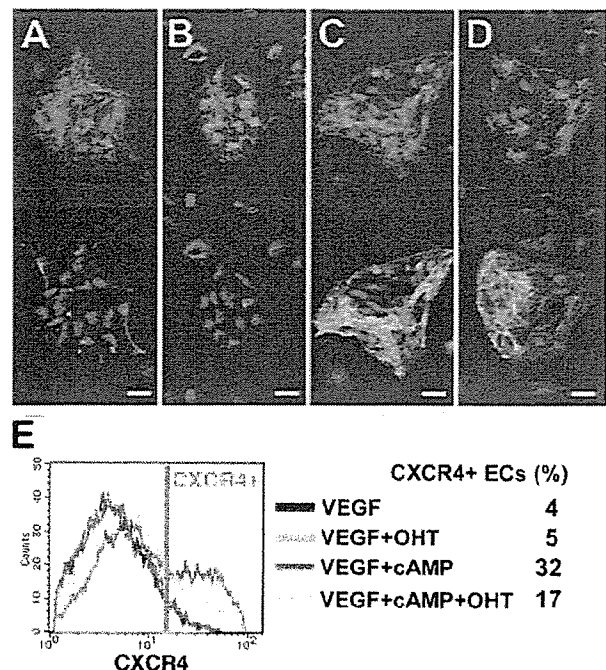


Figure 4. Effects of activated Notch on arterial EC induction from VEGFR2⁺ cells. A-D, Double-fluorescent staining of CD31 and ephrinB2 for induced ECs using NERT ^{Δ OP-7} ES cells. Upper panels, CD31 (pan-ECs, red) and DAPI (blue). Lower panels, EphB4-Fc (ephrinB2⁺ arterial ECs, green) and DAPI (blue). A, VEGF treatment alone (50 ng/mL). B, VEGF and 150 nmol/L OHT. C, VEGF and 0.5 mmol/L 8bromo-cAMP. D, VEGF, 0.5mmol/L 8bromo-cAMP, and 150 nmol/L OHT. Scale bars: 100 μ m. E, Expression profile of CXCR4 in CD31⁺ ECs. VEGF alone (blue line), VEGF and OHT (green line), VEGF and 8bromo-cAMP (red line), and VEGF, 8bromo-cAMP, and OHT (orange line) are shown. Percentages of CXCR4⁺ arterial ECs in total ECs are indicated.

and Notch signaling is essential but not sufficient. AM/cAMP pathway can activate Notch signaling, and is another important signaling to induce arterial ECs. Coordinated signaling of VEGF, Notch, and cAMP is the combination that composes a sufficient condition to constructively induce arterial ECs from vascular progenitor cells.

Discussion

Our findings provide the first demonstration to our knowledge of arterial and venous EC induction from ES cells by constructively reproducing endothelial differentiation processes *in vitro*. Here we showed that cAMP and AM play specific roles in EC differentiation, especially for arterial EC induction, from VEGFR2⁺ vascular progenitors. We have shown that AM enhances proliferation and migration of cultured ECs and can promote angiogenesis in gel plug assays *in vivo*.³⁷ Recently, AM was reported to enhance angiogenic potency of bone marrow cell transplantation.³⁸ AM should be a novel potent candidate for an endogenous ligand for EC differentiation as well as arterial EC induction.

Our results showed that stimulation of cAMP pathway can activate Notch signaling in EC lineage. To date, little evidence of Notch activation by cAMP pathway has been reported. In neuronal cells, cAMP-response element-binding protein increased expression of presenilin-1, a component of

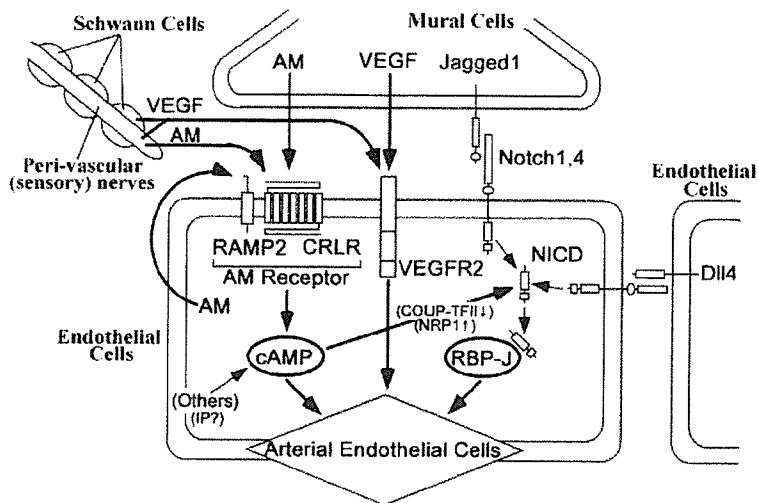


Figure 5. Cellular and molecular mechanisms of arterial EC induction. Putative autocrine/paracrine system for arterial induction in vascular wall. Signals for arterial induction, VEGF, Notch, and AM/cAMP components exist in the vascular wall. VEGFR2, AM receptor complex, RAMP2 and CRLR, and Notch1 and 4 are expressed in ECs. However, their ligands, VEGF, AM, and Jagged1 are expressed in mural cells (MCs). Moreover, AM is expressed in ECs and perivascular nerves. VEGF is produced from peripheral sensory nerves and Schwann cells. Notch ligands, Dll4, and Jagged1 are expressed in arterial ECs. These autocrine/paracrine signals among ECs, MCs, and other perivascular tissues should coordinately regulate the arterial induction and maintenance.

γ -secretase, through transcriptional activation.³⁹ A similar mechanism may contribute in EC and EC progenitors to induce Notch activation. Recently, COUP-TFII has been reported to repress Notch signaling through suppressing NRP1 expression to maintain vein identity.²⁸ Administration of 8bromo-cAMP did not increase mRNA expression of Notch ligands (ie, jagged1, 2, Delata-like1, 3, 4) in surrounding mural cells (data not shown), but suppressed COUP-TFII and increased NRP1 expression in ECs. These results suggest that cAMP pathway may activate Notch signaling through the suppression of COUP-TFII expression. cAMP pathway, thus, may regulate the determination of cell fates between arterial and venous ECs. Although Dll4 and Notch signaling were reported to be growth-suppressive on mature ECs through downregulation of VEGFR2 and NRP1 expression,⁴⁰ forced Notch activation with OHT did not affect on VEGFR2 and NRP1 mRNA expression in differentiating ECs (data not shown). Notch signaling may possess differentiation stage-specific roles in EC differentiation and proliferation. Precise molecular interactions among these pathways should be further investigated to figure out the whole scheme of arterial-venous specification.

In the vascular wall, VEGFR2, Notch1 and 4, and AM receptor complex, CRLR, RAMP-2 and -3, are expressed in ECs.^{5,6} On the other hand, their ligands, VEGF, Jagged1, and AM, are expressed in MCs.^{8,41,42} Dll4 and AM are also expressed in ECs. We confirmed AM mRNA expression in ES cell-derived ECs and MCs, and RAMP-2 and CRLR mRNA in ECs by reverse-transcription polymerase chain reaction analysis. Low-level expression of prostaglandin-I2 receptor mRNA was also observed in ECs (data not shown). Moreover, peripheral sensory nerve and Schwann cell-derived VEGF are reported to be involved in arterial EC induction.⁴³ AM is demonstrated to be expressed in perivascular nerves in the rat mesenteric artery.⁴⁴ The autocrine/paracrine cross-talk of VEGF, Notch, and AM/cAMP signaling between ECs and MCs, and signals from other perivascular tissues, should coordinately regulate vascular development including the induction and maintenance of the arterial structures (Figure 5). Combinatory signaling of VEGF, Notch, and cAMP may mimic these arterial-inducing

machineries in vivo to achieve constructive induction of arterial ECs from vascular progenitor cells in vitro.

Our constructive approach has successfully provided a novel understanding for the mechanisms of arterial EC differentiation. This study, thus, would provide a potent novel strategy as constructive developmental biology to dissect cell differentiation processes and contribute to regenerative medicine.

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Disclosures

None.

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The Neuroprotective and Vasculo-Neuro-Regenerative Roles of Adrenomedullin in Ischemic Brain and Its Therapeutic Potential

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Adrenomedullin (AM) is a vasodilating hormone secreted mainly from vascular wall, and its expression is markedly enhanced after stroke. We have revealed that AM promotes not only vasodilation but also vascular regeneration. In this study, we focused on the roles of AM in the ischemic brain and examined its therapeutic potential. We developed novel AM-transgenic (AM-Tg) mice that overproduce AM in the liver and performed middle cerebral artery occlusion for 20 min (20m-MCAO) to examine the effects of AM on degenerative or regenerative processes in ischemic brain. The infarct area and gliosis after 20m-MCAO was reduced in AM-Tg mice in association with suppression of leukocyte infiltration, oxidative stress, and apoptosis in the ischemic core. In addition, vascular regeneration and subsequent neurogenesis were enhanced in AM-Tg mice, preceded by increase in mobilization

of CD34⁺ mononuclear cells, which can differentiate into endothelial cells. The vasculo-neuro-regenerative actions observed in AM-Tg mice in combination with neuroprotection resulted in improved recovery of motor function. Brain edema was also significantly reduced in AM-Tg mice via suppression of vascular permeability. *In vitro*, AM exerted direct antiapoptotic and neurogenic actions on neuronal cells. Exogenous administration of AM in mice after 20m-MCAO also reduced the infarct area, and promoted vascular regeneration and functional recovery. In summary, this study suggests the neuroprotective and vasculo-neuro-regenerative roles of AM and provides basis for a new strategy to rescue ischemic brain through its multiple hormonal actions. (*Endocrinology* 147: 1642-1653, 2006)

ADRENOMEDULLIN (AM) IS a potent vasodilating peptide comprising 52 amino acids, which was originally isolated from human pheochromocytoma tissues in 1993 as a substance to elevate cAMP concentration in platelets (1). It is secreted mainly from the vascular wall into circulating blood to reduce pre- and post-load on the heart via vasodilation, natriuresis, and suppression of aldosterone release. Intravenous administration of AM to patients with heart failure or pulmonary hypertension has already been initiated and beneficial hemodynamic effects have been reported (2).

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Abbreviations: AM, Adrenomedullin; ANCOVA, analysis of covariance; BP, blood pressure; BrdU, bromodeoxyuridine; CGRP, calcitonin gene-related peptide; diHE, dihydroethidium; GFAP, glial fibrillary acidic protein; LDPL, laser Doppler perfusion imager; MCA, middle cerebral artery; 20m-MCAO, middle cerebral artery occlusion for 20 min; NeuN, neuronal marker; NHNP, normal human neuronal progenitor cells; PAMP, proadrenomedullin N-terminal 20 peptide; PECAM, platelet endothelial cell adhesion molecule; PI3K, phosphatidylinositol-3 kinase; PKA, protein kinase A; ROS, reactive oxygen species; ssDNA, single-strand DNA; Tg, transgenic; Wt, wild type.

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Along with its vasodilating effect, a number of studies have demonstrated various and significant effects of AM on the regulation of vascular structure, including its development, remodeling, and regeneration. Mice lacking the AM gene did not survive their embryonic stage and showed abnormal vasculature with sc hemorrhage (3, 4). Mice overexpressing AM in endothelial cells were revealed to be hypertensive and resistant to vascular remodeling such as neointima formation caused by cuff injury, and atherosclerosis associated with a high-cholesterol diet (5). We have recently established that AM promotes endothelial regeneration in the wound healing assay using cultured endothelial cells and enhances neovascularization *in vivo* into sc implanted gel-plugs in mice (6, 7). We and others (8-11) have further demonstrated that the potentiating action of AM on vascular regeneration is mediated by activation of the phosphatidylinositol-3 kinase (PI3K)-Akt pathway.

Recently, it has been known that AM is secreted from various organs including the heart, lung, kidney, adipose tissues, and central nervous system (12). Moreover, AM expression has been demonstrated to be markedly enhanced by ischemia through the activation of hypoxia-responsive elements in the AM gene via transcription factor hypoxia-inducible factor-1. In the central nervous system, where AM is

mainly expressed in neurons and the endothelium (13), it is reported that transient ischemia boosted AM expression for more than 15 d (14). However, the role of augmented AM has remained unclear for inconsistent previous results: three studies reported neuroprotective effects of AM by demonstrating reduction of infarct size after transient ischemia (15–17), whereas one study detected exacerbation of infarction as a result of AM infusion (14).

In this context, our study presented here focused on the roles of augmented AM in ischemic brain and examined its therapeutic potential. We generated new lines of transgenic mice that overproduce AM (AM-Tg) in the liver that mimics chronic AM administration. After inducing 20-min middle cerebral artery occlusion (20m-MCAO) to produce a nonfatal stroke model in the AM-Tg mice, we observed the long-term effects of AM on the ischemic brain up to postoperative d 56. We examined the mice for the recovery of blood flow in the ischemic region and impaired motor function after stroke, and immunohistochemically examined the ischemic striatum to determine effects of AM on neuronal loss/apoptosis, gliosis, leukocyte infiltration, oxidative stress, vascular regeneration, and neurogenesis after 20m-MCAO. In addition, another stroke model, 2-h middle cerebral artery occlusion (2 h-MCAO), was performed to observe the effect of AM in acute phase of the fatal stroke. *In vitro* studies using neuronal progenitor cells or rat pheochromocytoma PC12 cells were performed to examine direct antiapoptotic and neurogenic

actions of AM on these neuronal cells. Finally, we investigated the effect of exogenous AM administration after 20m-MCAO to determine the appropriate amount and timing of AM treatment after cerebral ischemia.

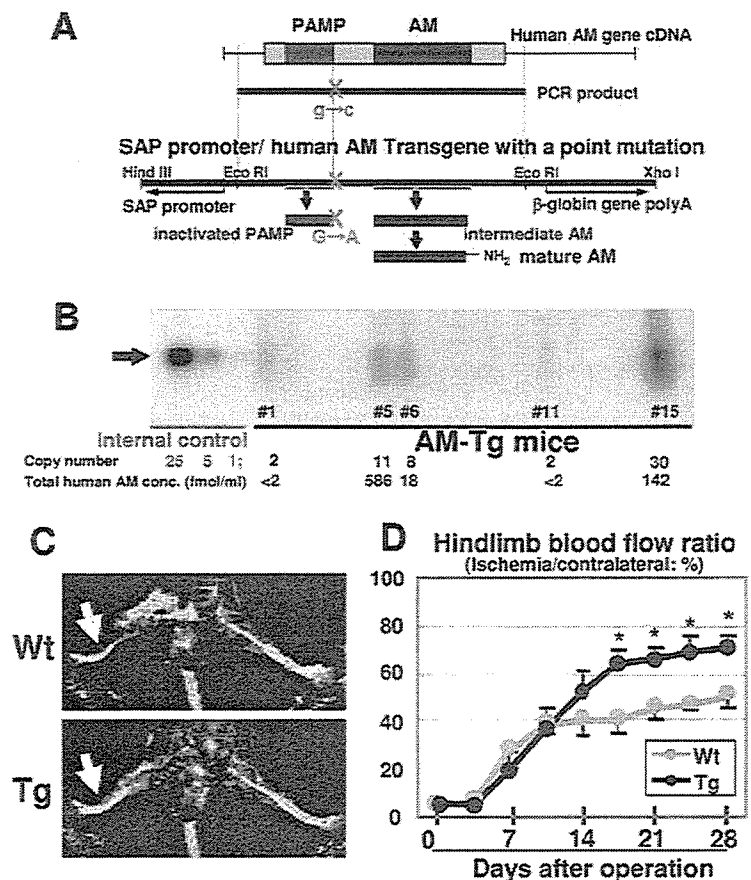
Materials and Methods

Generation of transgenic mice which overproduce human AM but do not overproduce mature proadrenomedullin N-terminal 20 peptide (PAMP)

The AM gene contains coding regions for not only AM but also PAMP, a different vasodilating peptide. Amidation at their carboxyl terminals after their synthesis is needed for both AM and PAMP to exert their biological activity. The bioactive amidated forms are known as mature AM and mature PAMP, respectively. To identify the specific effects of AM, we generated a transgene construct with a point mutation on the PAMP amidation signal in the full-length AM gene cDNA. Guanine was substituted for cytosine on the 3' end of the PAMP coding region so that glycine on the C' terminal of the PAMP product was replaced with alanine. In this way, amidation and maturation of PAMP by peptidylglycine α -hydroxylase and α -hydroxyglycine N-C lyase were inhibited (Fig. 1A). The mutant AM gene cDNA was then inserted into a plasmid containing the human serum amyloid P component promoter, which is widely used to target gene expression specific to the liver. When the product is secreted from the liver, it mimics intravenous administration of the agent. The *Hind*III-*Xho*I fragment of the plasmid was microinjected into the pronucleus of fertilized C57BL/6J mice eggs.

The copy number of transgenes was quantified by means of genomic Southern blotting according to standard procedure. Plasma concentrations of human total AM and mature AM were measured with a commercially available immunoradiometric assay (Cosmic, Tokyo, Japan).

FIG. 1. Generation of transgenic mice which overproduce AM but do not overproduce mature PAMP in the liver and augmented angiogenesis in the transgenic mice after femoral artery occlusion. A, Schematic representation of the transgene construct derived from human AM gene cDNA with a point mutation in the amidation signal of PAMP. B, Southern blot analysis of the tail DNA of the founder mice. Arrow, Blots for the transgene. Internal controls for indicated copies are located in the left three lanes. The line No. indicates the mice in which the transgene was detected by PCR. The copy numbers estimated by densitometry and the plasma concentrations of total human AM in F3 mice of the lines are shown. C, Hindlimb blood flow analyzed by LDPI. Red or white indicates a higher flow than blue or green. Arrows, Comparison of ischemic hindlimbs between Wt and AM-Tg on d 28 after femoral artery ligation. D, Quantitative analysis of the hindlimb blood flow in ischemia. *, $P < 0.05$ for Wt vs. AM-Tg by ANCOVA; $n = 6$.



Human mature PAMP concentration was measured with a recently developed enzyme immunoassay (18). To determine the brain concentration of AM, we used the RIA kits for measurement of human and mouse total AM (Phoenix, Belmont, CA), according to the manufacturer's instruction. Blood pressure (BP) was measured with tail cuff (Softron, Tokyo, Japan). Hindlimb ischemia was induced by ligating the right femoral artery and blood flow of the ischemic limb was estimated with a laser Doppler perfusion imager (LDPI; Moor Instruments Ltd., Devon, UK) to confirm the angiogenic effect of AM-Tg mice. The perfusion ratio (%) was calculated as that of the ipsilateral to the contralateral side. Animal care and experiments were in accordance with the guidelines for animal experiments of Kyoto University.

Induction of stroke by MCAO

We performed nonfatal 20m-MCAO and fatal 2 h-MCAO by the standard *trans*-luminal method, which has been described in various previous reports (19). Briefly, a 8–0 nylon monofilament coated with silicone was inserted from the left common carotid artery via the internal carotid to the base of the left middle cerebral artery (MCA) of 12-wk-old mice anesthetized with 5% halothane and maintained on 1%. After 20 min or 2 h of occlusion, the filament was withdrawn; and the arteries were reperfused, whereas the left common carotid artery was permanently ligated. Occlusion and reperfusion of the MCA was confirmed by means of fiber-shaped laser Doppler perfusion imager (Omegawave, Tokyo, Japan). We observed the mice until postoperative d 56 to examine blood flow in the ischemic region with an LDPI and motor function with a rota-rod exercise test.

Immunohistochemical examination of the ischemic striatum

After the induction of 20m-MCAO, mice were killed on postoperative d 0–56 and the harvested brains were subjected to immunohistochemical examination using a standard procedure described elsewhere (20). We used these primary antibodies: neuronal marker, NeuN (1:200; Chemicon, Temecula, CA); astrocyte marker, glial fibrillary acidic protein (GFAP) (1:400; Chemicon); apoptosis marker, single-strand DNA (ssDNA) (1:50; Dako, Carpinteria, CA); leukocyte marker, CD45 (1:100, PharMingen, San Diego, CA); endothelial marker, platelet endothelial cell adhesion molecule (PECAM)-1 (CD31) (1:100, PharMingen); and a marker for proliferating cells, bromodeoxyuridine (BrdU) (1:50, Molecular Probes, Eugene, OR); to examine infarct area, gliosis, leukocyte infiltration, apoptosis, vascular regeneration and neurogenesis. Briefly, free-floating 30- μ m coronal sections at the level of the anterior commissure were stained and observed with a confocal microscope (LSM5 PASCAL; Carl Zeiss SMT AG, Oberkochen, Germany). The infarct area (mm^2/field) was defined and quantified as the region where loss of NeuN immunoreactivity was observed and gliosis (mm^2/field) as the area stained GFAP in the ischemic striatum at $\times 5$ fields. CD45 or ssDNA-positive cells (cells/mm^2) were quantified to serve as an index of leukocyte infiltration or of apoptosis, respectively, in the ischemic core at $\times 20$ magnification. Capillary density was quantified as the number of PECAM-1-positive cells (cells/mm^2). The vessel counts were performed in the region of ischemic core at 0.5–1.0 mm anterior from the bregma. We prepared two thin sections (6 μ m thickness) per mouse for vessel counting and four representative fields from each section were evaluated for capillary density in the ischemic core. To examine neurogenesis, mice were injected ip with BrdU 50 mg/kg (Sigma-Aldrich Co., St. Louis, MO) twice daily on postoperative d 4–6 and the number of BrdU-NeuN double-positive cells (cells/mm^2), which are generally defined as regenerated neurons, were quantified to serve as an index of neurogenesis. We also examined the production of reactive oxygen species (ROS) *in situ* by using the oxidative fluorescent dye dihydroethidium (dHE; 2×10^{-6} M; Sigma).

Quantification of CD34⁺ mononuclear cells after 20m-MCAO

We counted peripheral CD34⁺ mononuclear cells according to the International Society of Hematotherapy and Graft Engineering (ISHAGE) guidelines (21). Briefly, peripheral blood was taken from the orbital vein and stained with CD34-PE and CD45-FITC monoclonal antibodies (BD PharMingen, San Jose, CA) in a TruCOUNT tube (BD

PharMingen) according to the manufacturer's instruction. After the reaction, CD34⁺-CD45^{dim} cells were quantified as CD34⁺ mononuclear cells by a fluorescence-activated cell sorting machine Aria (BD) by using the ISHAGE sequential gating strategy (21).

Analysis of infarct volume and brain edema after 2 h-MCAO

We performed 2 h-MCAO to examine the effect of AM in the acute phase of fatal stroke. To estimate infarct or edema volume, mice were killed 24 h after the occlusion. The brain was removed and cut into 2 mm-thick slices and immersed in saline containing 2% 2,3,5-triphenyl-tetrazolium chloride for 30 min at 4 C. Infarct or edema volume was calculated as the percentage volume of the contralateral hemisphere with a standard procedure as described elsewhere (22). We estimated Evans Blue leakage in the brain parenchyma as previously reported (23), to serve as an index of vascular permeability *in situ*. Briefly, 0.2 ml of 2.5% Evans Blue solution was injected into mice via a tail vein 10 min before 2 h-MCAO and mice were killed at 24 h after the ischemia. Brain tissues were weighed and homogenized in 50% trichloroacetic acid solution to extract the dye in the supernatant. The tissue content of Evans Blue was estimated from the absorbance of 620 nm.

Estimation of apoptosis and differentiation of neuronal cells

The ratio of apoptotic cells was examined using normal human neuronal progenitor cells (NHNP; Cambrex Bioscience, Walkersville, MD). Cells were plated at a density of 5×10^4 cells/ cm^2 on a laminin-coated 24-well dish and incubated in serum-free neuronal basal medium for 48 h. After the experimental period, the cell number was assessed by 5-mercapto-1-methyltetrazole assay (Nakalai Tesque), and the cells were stained with an anti-ssDNA antibody and nuclear staining propidium iodide to calculate the ratio of apoptotic cells to the total cells in each microscopic image.

Neuronal differentiation was examined as described previously (24), using rat pheochromocytoma PC12 cells (Riken Gene Bank, Tsukuba, Japan). Briefly, the length of the neuronal process (micrometers/cell) was calculated to serve as an index of neuronal differentiation after plating at a density of 10^4 cells/ cm^2 on a collagen I-coated 24-well dish and incubated in 1% serum DMEM for 7 d. The cells were treated with 10^{-5} mol/liter AM or 100 ng/ml nerve growth factor as a positive control, and with the following inhibitors: the two AM antagonists, 10^{-5} mol/liter AM (22–52) and 10^{-5} mol/liter calcitonin gene-related peptide(8–37) [CGRP(8–37)] (Peptide Institute Inc., Osaka, Japan), the two protein kinase A (PKA) inhibitors, 10^{-5} mol/liter adenosine 3P,5P-cyclic monophosphorothioate Rp-isomer (Rp-cAMP) and 10^{-6} mol/liter myristoylated cell-permeable PKA inhibitor peptide sequence (14–22) (PKA Inh), and the two PI3K inhibitors, 10^{-3} mol/liter LY294002 and 10^{-7} mol/liter wortmannin (Calbiochem, San Diego, CA). For endothelial cell coculture experiments, human umbilical vein endothelial cells (HUVEC; Cambrex) were plated into transwell membrane inserts at a density of 10^5 cells/ cm^2 .

Exogenous administration of AM and hydralazine

Recombinant human mature AM dissolved in 0.9% saline was exogenously administered to C57BL/6J wild-type mice (Wt) by means of osmotic pumps (Alzet Model 2002; Alzet Osmotic Pumps Co., Cupertino, CA) at a rate of 50 ng/h, which is estimated to achieve a plasma concentration of 2 fmol/ml (25). To determine appropriate timing to start AM treatment after 20m-MCAO, we implanted the pump ip just after the operation (d 0), or at 24 (d 1) or 72 h (d 3) later. We killed the mice on d 7 for histological examination and the period of the exogenous AM treatment was from d 0, 1, or 3 to d 7. In some experiments, low-dose (0.1 mM) hydralazine was exogenously administered in drinking water.

Statistics

All data were expressed as mean \pm SE. Comparison of means between two groups was performed with Student's *t* test. When more than two groups were compared, ANOVA was used to evaluate significant differences among groups, and if significant differences were confirmed, each difference was further examined by means of multiple comparisons. We

TABLE 1. Plasma concentrations of human AM and systolic BP in Wt and three lines of AM-Tg mice

	Wt	Low conc.	Medium conc.	High conc.
Total AM (fmol/ml)	1.1 ± 0.2	17.6 ± 4.4 ^a	142.2 ± 18.4 ^a	585.5 ± 117.7 ^a
Mature AM (fmol/ml)	0.5 ± 0.4	2.6 ± 0.6 ^a	10.4 ± 2.4 ^a	24.9 ± 4.2 ^a
Systolic BP (mm Hg)	122.7 ± 1.6	113.0 ± 2.5 ^a	113.4 ± 2.6 ^a	109.4 ± 2.5 ^a

conc., Concentration.

^a $P < 0.01$ vs. Wt; $n = 4$ –12.

performed analysis of covariance (ANCOVA) when repeated-measurement had done, specifically, in the rota-rod test and laser Doppler flowmetry. Probability was considered to be statistically significant at $P < 0.05$.

Results

Generation of transgenic mice that overproduce human AM but do not overproduce mature PAMP

We generated seven lines of founder mice carrying the transgene and maintained three of them (lines 5, 6, and 15). Their plasma concentrations of human total AM were 585.5 ± 117.7 , 17.6 ± 4.4 and 142.2 ± 18.4 fmol/ml and the copy numbers of the transgene estimated by Southern blot densitometry analysis were 11, 8, and 30, respectively (Fig. 1B). The physiological concentration of mouse total AM is reportedly 5–10 fmol/ml, so that the transgenic mice were expected to overproduce AM about 100, 3, and 30 times more than endogenous AM. The three lines were designated low (no. 6), medium (no. 15), and high (no. 5) concentration line according to their plasma AM concentration. The high concentration line (no. 5) was used for further study unless

otherwise indicated. The plasma concentration of human mature AM, the bioactive amidated form, increased to 2.6–24.9 fmol/ml in the AM-Tg mice (Table 1). On the other hand, plasma human mature PAMP did not change in AM-Tg mice. The concentration (fmol/ml) was 2.21 ± 0.58 in Wt vs. 2.15 ± 0.35 in AM-Tg ($n = 6$), so that the point mutation on the amidation signal in the PAMP coding region was expected to successfully inhibit maturation of PAMP. There were no apparent differences in overall appearance, behavior, growth or fertility between Wt and AM-Tg mice. The systolic BP in 12-wk-old mice was significantly reduced in all three lines of AM-Tg compared with Wt. The BP (mm Hg) was 122.7 ± 1.6 in Wt vs. 109.4 ± 2.5 – 113.4 ± 2.6 in AM-Tg, depending on the line ($P < 0.05$; $n = 5$; Table 1).

Therapeutic angiogenesis in hindlimb ischemia model was promoted in AM-Tg mice

The recovery of blood flow in the ischemic hindlimb of Wt and AM-Tg mice was compared and was found to have

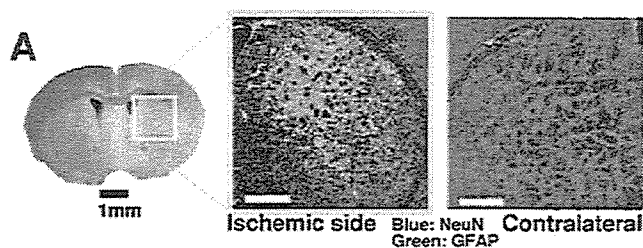
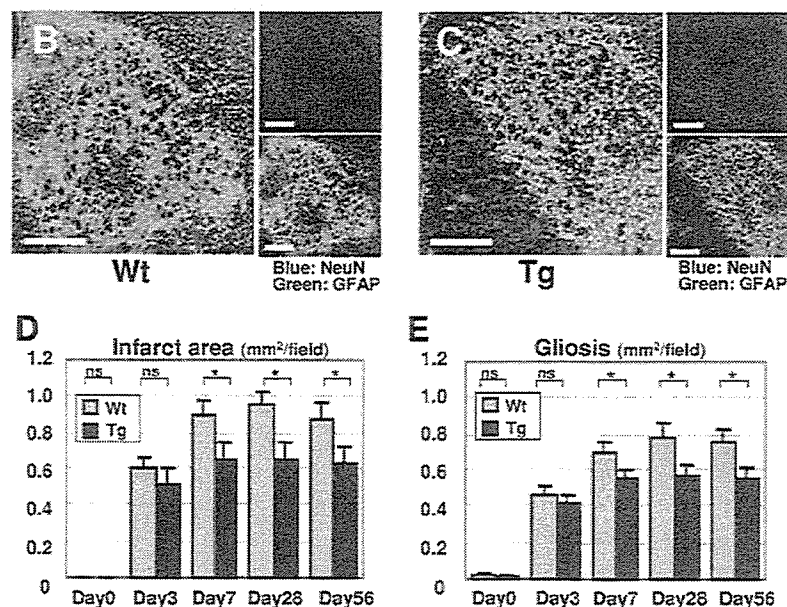


FIG. 2. Effects of AM on infarct area and gliosis after the nonfatal stroke, 20m-MCAO. A, Histological examination of the ischemic striatum. The outlined field was examined for infarct area and gliosis. The ischemic side and contralateral side on d 3 after 20m-MCAO are shown. Scale bar, 500 μ m ($\times 5$ magnification). B and C, Representative images of the ischemic striatum on post-operative d 7 stained for NeuN (blue) and GFAP (green). Infarct area, defined as the region where NeuN immunoreactivity was lost, and gliosis, defined as the area where GFAP immunoreactivity was observed, in Wt (B) and AM-Tg (C) are shown. Scale bar, 500 μ m ($\times 5$ magnification). D and E, Quantitative analysis of the infarct area (D) and gliosis (E) * $P < 0.05$; ns, not significant for Wt vs. AM-Tg; $n = 12$.



significantly improved in AM-Tg mice after postoperative d 17. The hindlimb blood flow ratio on d 28 (ipsilateral/contralateral, %) was 56.6 ± 8.3 in Wt *vs.* 73.8 ± 5.3 in AM-Tg ($P < 0.05$; $n = 6$; Fig. 1, C and D). In this way, promotion of therapeutic angiogenesis by AM was confirmed in AM-Tg mice.

Brain remodeling in ischemic striatum after 20m-MCAO

We investigated the time course of neuronal loss, reactive gliosis, vascular regeneration, and neuronal regeneration; the entire process can be defined as “brain remodeling” after ischemia.

20m-MCAO caused selective loss of NeuN-positive cells and marked reactive gliosis (Fig. 2A) in the ipsilateral striatum within 24 h after the operation; this condition was different from pan-necrosis caused by longer MCAO (*e.g.* 2 h-MCAO). The infarct area, that is, the area of neuronal loss, expanded progressively up to d 7, and then showed gradual increase in size until d 56, whereas gliosis spread in parallel. The expansion of the infarct area in the subacute to chronic phase after mild stroke was compatible with previously reported findings (26). Vascular regeneration in the striatum with enhanced capillary density was obvious after postoperative d 7, and subsequent neurogenesis became obvious after d 28.

The concentrations of the overproduced human AM (fmol/g tissue) in the ischemic brain of AM-Tg mice before

and on postoperative d 1 and 28 after 20m-MCAO were 27.8 ± 10.3 , 87.4 ± 4.0 and 30.3 ± 16.8 , respectively. Those of endogenous mouse AM (fmol/g tissue) were 3.7 ± 2.1 , 7.2 ± 2.5 , and 4.6 ± 3.0 .

Infarct area and gliosis were reduced in AM-Tg mice after 20m-MCAO along with suppression of leukocyte infiltration and ROS production

A significant decrease in infarct area and gliosis was observed in AM-Tg mice (Fig. 2, B–E) after postoperative d 7, but was not obvious on d 3. The infarct area (mm^2/field) on d 56 was 0.88 ± 0.08 in Wt *vs.* 0.64 ± 0.08 in AM-Tg ($P < 0.05$; $n = 12$; Fig. 2D), and gliosis (mm^2/field) on the same day was 0.76 ± 0.08 in Wt and 0.56 ± 0.07 in AM-Tg ($P < 0.05$; $n = 12$; Fig. 2E). Leukocyte infiltration quantified as the number of CD45⁺ cells was significantly suppressed in AM-Tg mice especially from d 3–7. CD45⁺ cells on d 3 ($/\text{mm}^2$) numbered 197.5 ± 16.6 in Wt *vs.* 140.7 ± 14.6 in AM-Tg ($P < 0.05$; $n = 12$; Fig. 3, A, B, and G). *In situ* ROS production detected by immunostaining for diHE, which stained the nucleus of NeuN⁺ or GFAP⁺ cells, was enhanced in Wt compared with that in AM-Tg mice (Fig. 3, C and D). Apoptotic cells quantified as the number of ssDNA⁺ cells in the ischemic core were significantly reduced in the AM-Tg mice on d 3–7. ssDNA⁺ cells ($/\text{mm}^2$) on d 3 numbered 214.8 ± 19.6 in Wt *vs.* 123.2 ± 11.1 in AM-Tg ($P < 0.01$; $n = 12$; Fig. 3, E, F, and H).

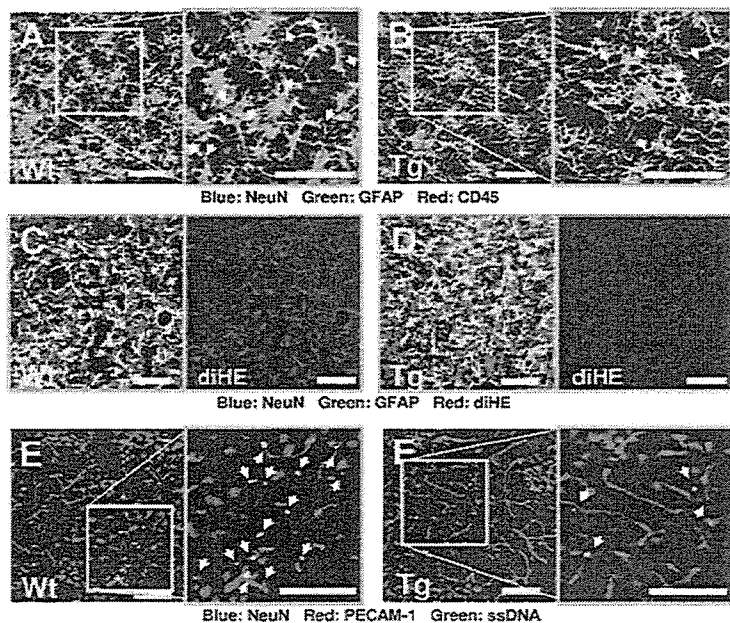
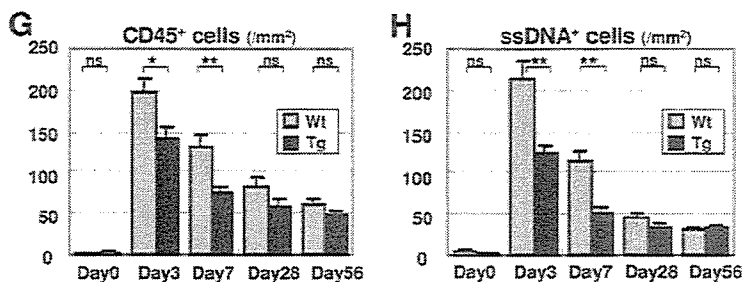


FIG. 3. Effects of AM on leukocyte infiltration, ROS production, and apoptosis in the ischemic brain after 20m-MCAO. A and B, Detection of leukocyte infiltration in the ischemic core on postoperative d 7 by immunostaining for CD45⁺ cells (red) in Wt (A) and AM-Tg (B). Arrows, CD45⁺ cells. C and D, *In situ* detection of ROS in ischemic striatum on postoperative d 7 by immunostaining for diHE (red) in Wt (C) and AM-Tg (D). E and F, Detection of apoptotic cells in the ischemic core on postoperative d 7 by immunostaining for ssDNA⁺ cells (green) in Wt (E) and AM-Tg (F). Arrows, ssDNA⁺ cells. G and H, Quantitative analysis of CD45⁺ cells (G) and ssDNA⁺ cells (H) in the ischemic core. *, $P < 0.05$; **, $P < 0.01$; ns, not significant for Wt *vs.* AM-Tg; $n = 12$. Scale bar, 100 μm ($\times 20$ magnification).



Vascular regeneration was augmented in AM-Tg mice after 20m-MCAO associated with increased mobilization of CD34⁺ mononuclear cells

The blood flow in the ischemic brain estimated by LDPI was significantly higher in AM-Tg mice after postoperative d 7 and higher flow was maintained until d 56. The brain blood flow ratio (ipsilateral/contralateral, %) on d 56 was 88.9 ± 2.8 in Wt vs. 97.6 ± 3.0 in AM-Tg ($P < 0.01$ by ANCOVA; $n = 8$; Fig. 4, C, D, and H). We were also able to confirm that capillary density determined as the number of PECAM-1⁺ cells was augmented in AM-Tg mice. The density (/mm²) on d 56 was 468.8 ± 21.8 in Wt vs. 536.6 ± 13.6 in AM-Tg ($P < 0.05$; $n = 8$; Fig. 4I). Thus, the physiological neovascularization in the ischemic core after stroke was augmented in AM-Tg mice. Peripheral CD34⁺ mononuclear cells were physiologically enhanced after 20m-MCAO and further increased in AM-Tg mice on d 3–7. The cells (/ml) on d 3 numbered 1774 ± 272 in Wt vs. 3199 ± 562 in AM-Tg ($P < 0.05$; $n = 6$; Fig. 5, A–C).

Augmented neurogenesis and improved recovery of impaired neurological function were observed in AM-Tg mice after 20m-MCAO

BrdU injection on postoperative d 4–6 proved that most BrdU-positive cells were costained with GFAP (data not shown) and that there were far fewer BrdU-PECAM-1 or BrdU-NeuN double-positive cells. We found that regenerated neurons defined as BrdU-NeuN double-positive cells

were frequently detected adjacent to the vasculature and the number of these cells on d 56 was correlated with capillary density ($P = 0.003$; $n = 12$; Fig. 6, A and B; and Table 2). The cells increased from postoperative d 7–56, and their number was significantly higher in AM-Tg mice. The regenerated neurons (/mm²) on d 56 numbered 20.4 ± 3.9 in Wt vs. 33.9 ± 4.7 in AM-Tg ($P < 0.05$; $n = 12$; Fig. 6C).

Recovery of impaired motor function after 20m-MCAO, quantified as the exercise time on an accelerating rota-rod from the start to collapse down, was significantly better in AM-Tg mice. The exercise time (second) on d 49 was 21.5 ± 1.5 for Wt vs. 27.1 ± 2.0 for AM-Tg ($P < 0.01$ by ANCOVA; $n = 14$; Fig. 6D). To confirm whether vasculogenesis and neurogenesis are the contributing factor to the recovery from the ischemic damage, we analyzed the relation between capillary density, the number of regenerated neuron and the rota-rod result in AM-Tg mice after 20m-MCAO. As shown in Table 2, we found that the capillary density was significantly correlated with the rota-rod exercise time ($P = 0.005$; $n = 24$) and neurogenesis tended to be correlated with it ($P = 0.08$; $n = 12$).

Low-concentration AM-Tg mice also showed reduced infarct area and promoted vascular regeneration

We performed 20m-MCAO, using the low-concentration AM-Tg mice (plasma mature AM, 2.6 ± 0.6 fmol/ml) as well as the high-concentration line (plasma mature AM, 24.9 ± 4.2 fmol/ml) to determine appropriate concentration for AM

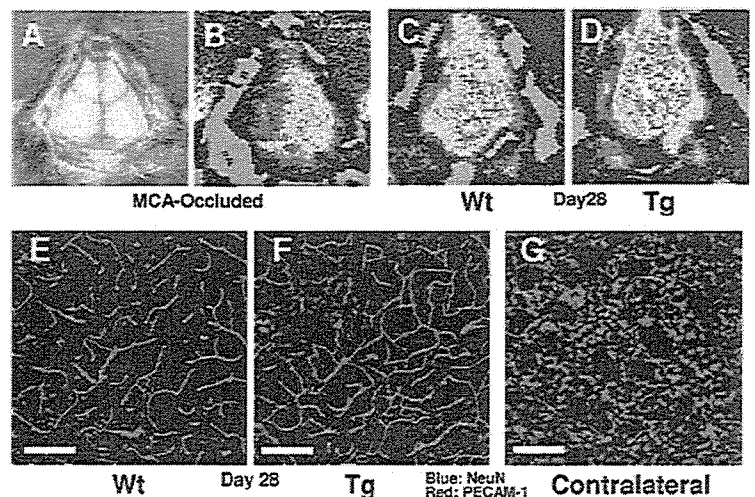
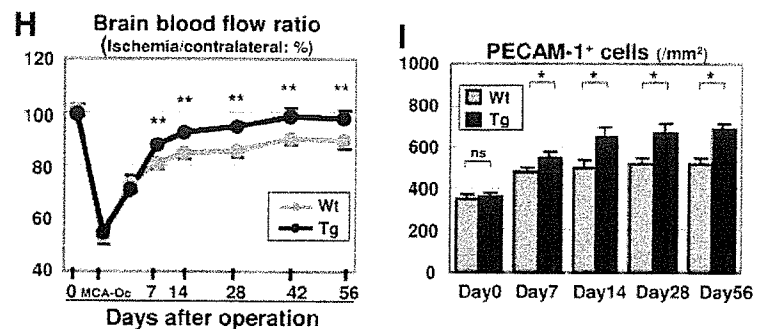


FIG. 4. Effects of AM on vascular regeneration in the ischemic brain after 20m-MCAO. A–D, Analysis of the blood flow in the ischemic brain by LDPI evaluated in mice with the scalp removed (A). Flowmetric analysis of the ischemic brain during MCA-Occlusion (B) and on d 28 after 20m-MCAO in Wt (C) and AM-Tg (D). Red or white indicates higher flow than blue or green. E–G, Histological examination of the vasculature in the ischemic core with PECAM-1 staining. Ischemic striatum on d 28 after 20m-MCAO in Wt (E) and AM-Tg (F), and contralateral nonischemic striatum (G). Scale bar, 100 μ m ($\times 20$ magnification). H, Quantitative analysis of the blood flow in the ischemic brain. Comparison of recovery from ischemia after 20m-MCAO between Wt and AM-Tg. MCA-Oc, blood flow during MCA occlusion; **, $P < 0.01$ for Wt vs. AM-Tg by ANCOVA; $n = 8$. I, Quantitative analysis of capillary density in the ischemic brain. Comparison of time course for increase in capillary density, determined as the number of PECAM-1⁺ cells, between Wt and AM-Tg mice. *, $P < 0.05$; ns, not significant; $n = 8$.



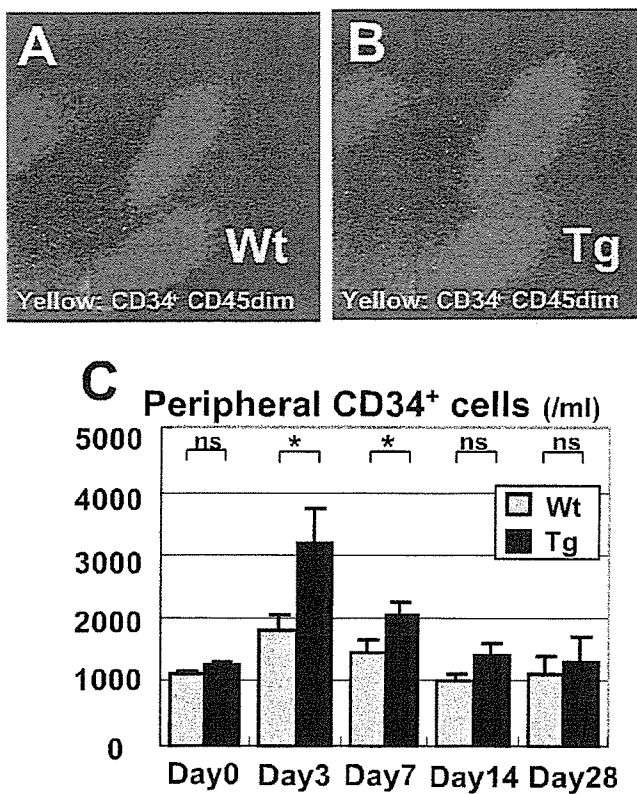


FIG. 5. Effects of AM on mobilization of CD34⁺ mononuclear cells into peripheral blood after 20m-MCAO. A–C, Quantification of CD34⁺ mononuclear cells after 20m-MCAO. Scatter plots for fluorescence-activated cell sorting analysis of the CD34⁺ cells in peripheral blood of Wt (A) and AM-Tg (B) on postoperative d 3. Yellow, CD34⁺-CD45dim mononuclear cells. Comparison of the time course for mobilization of CD34⁺ cells into peripheral blood between Wt and AM-Tg (C). *, $P < 0.05$; ns, not significant; $n = 6$.

treatment. The result showed comparable levels of neuroprotection and vascular regeneration between the low-concentration line and the high-concentration line (Table 3). We further analyzed BP-matched mice by administration of low-dose hydralazine (0.1 mM in drinking water) to exclude the possibility that lower BP observed in AM-Tg mice caused beneficial effects after 20m-MCAO. As shown in Table 3, lower BP alone did not reduce the infarct area nor promote vascular regeneration, although hydralazine administration caused BP reduction comparable to that in AM-Tg mice.

Brain edema was reduced in AM-Tg mice at 24 h after 2 h MCAO

The survival rate of mice after the fatal stroke, 2 h-MCAO, was 0% on d 7. We observed no significant difference in the rate between Wt and AM-Tg mice. The edema volume was reduced in AM-Tg mice 24 h after 2 h-MCAO; although the infarct volume showed no significant difference between them. Edema volume (% volume of contralateral hemisphere) was 13.5 ± 1.2 in Wt vs. 9.7 ± 0.9 in AM-Tg ($P < 0.05$; $n = 9$, Fig. 7C), whereas infarct volume (% volume of contralateral hemisphere) was 39.0 ± 4.9 in Wt vs. 44.5 ± 7.3 in AM-Tg (not significant; $n = 9$; Fig. 7, A and B). As shown in Fig. 7D, we found that Evans Blue leakage into the ischemic

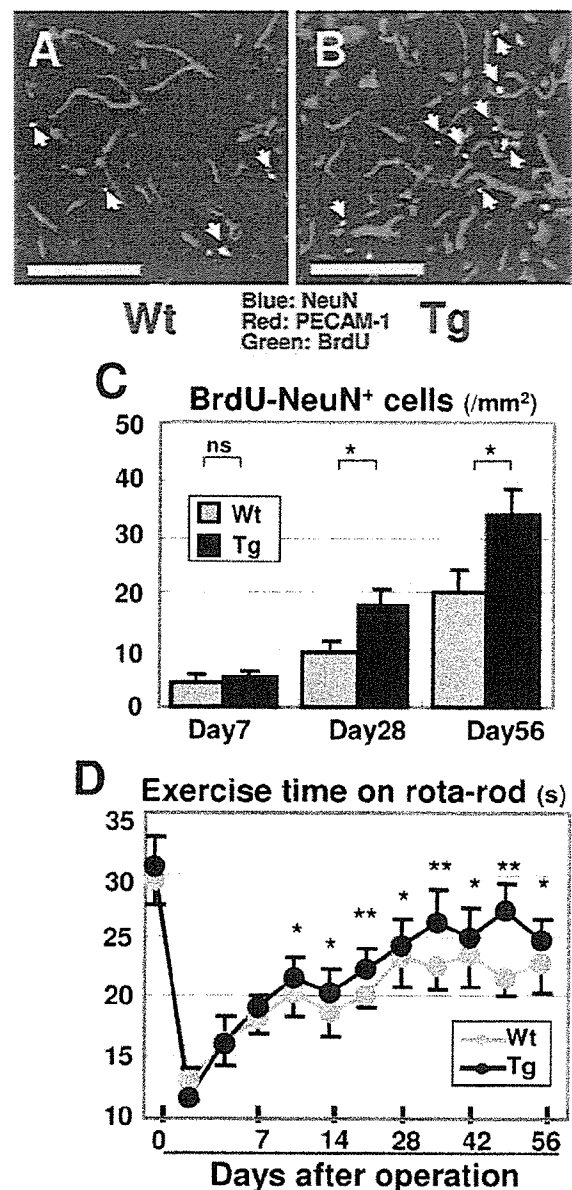


FIG. 6. Effects of AM on neurogenesis and recovery of impaired motor function after 20m-MCAO. A and B, Detection of regenerated neurons on postoperative d 56 by immunostaining for BrdU and NeuN. Arrows, BrdU-NeuN double-positive cells in the ischemic core of Wt (A) and AM-Tg (B). Scale bar, 100 μ m. C, Quantitative analysis of regenerated neurons. *, $P < 0.05$; ns, not significant; $n = 12$. D, Recovery of impaired motor function after 20m-MCAO, quantified as the exercise time on an accelerating rota-rod from the start to collapse down. *, $P < 0.05$; **, $P < 0.01$ for Wt vs. AM-Tg by ANCOVA; $n = 14$.

core was significantly reduced in AM-Tg mice. The content of Evans Blue (ng/g tissue) in the ischemic brain at 24 h after 2 h-MCAO was 239.4 ± 37.3 in Wt vs. 133.9 ± 9.4 in AM-Tg ($P < 0.01$; $n = 4$; Fig. 7E).

AM exerted direct antiapoptotic and neurodifferentiating effects on neuronal cells in vitro

After 48 h incubation of NHNP under serum-free apoptotic conditions, in which the number of the cells had decreased

TABLE 2. Significant correlation between the regenerative elements and apoptosis, neurogenesis, and functional recovery after 20m-MCAO

X	Y	Regression line	P
Capillary density (% field)	Apoptotic cells (/mm ²)	Y = -2.3X+37	0.01
Capillary density (% field)	Regenerated neuron (/mm ²)	Y = 3.2X-21	0.003
Capillary density (% field)	Rota-rod result (sec)	Y = 1.3X+9	0.005
Regenerated neuron (/mm ²)	Rota-rod result (sec)	Y = 0.3X+19	0.08

n = 12–24.

to half, the viable cell number was increased in the AM 10⁻⁸ mol/liter-treated group to 38.8 ± 7.1% over the control (*P* < 0.01; n = 4; Fig. 8C). The ratio of ssDNA⁺ cells to total cells (%) was 9.8 ± 1.9 in Wt *vs.* 4.0 ± 0.6 in the AM 10⁻⁸ mol/liter-treated group (*P* < 0.05; n = 4; Fig. 8, A, B, and D).

After 7-d incubation of PC12 cells under differentiation condition, both the cell number and the length of neuronal process increased dose dependently as a result of AM treatment (*P* < 0.01; n = 6; Fig. 8, E and I). Coculture with endothelial cells also increased the cell number and the length of neuronal process. The effect of AM was canceled by AM blockers, PKA inhibitors, and PI3K inhibitors (Table 4).

Exogenous administration of AM reduced infarct area, promoted vascular regeneration, and improved neurological function after 20m-MCAO

We further examined the effects of exogenous infusion of mature AM by means of an osmotic pump in the amount reported to achieve a plasma concentration of 2 fmol/ml. Implantation of the pump just after the operation resulted in increase in the blood flow and reduction of the infarct area on postoperative d 7 to a comparable level to those in AM-Tg mice. Moreover, the treatment started at 24 h after the operation (d 1) showed almost the same therapeutic effect. However, the implantation at 72 h after the operation (d 3) failed to reveal any significant effect (Fig. 9, A and B). The rota-rod exercise time was significantly improved in the AM-treated group. The exercise time (second) on d 7 was 17.0 ± 1.5 in vehicle group *vs.* 18.1 ± 2.0 in AM-treated group (n = 6 for vehicle group and 12 for AM-treated group; *P* < 0.05 by ANCOVA).

Discussion

In the present study, we generated novel transgenic mice that overproduce AM in their liver without overproduction of mature PAMP and investigated the roles of AM in degeneration or regeneration processes after brain ischemia, which can be defined as brain remodeling, as summarized in

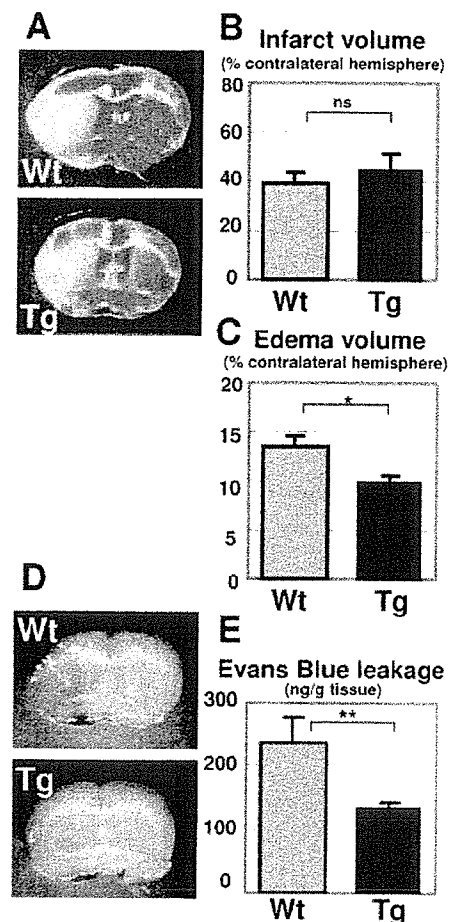
TABLE 3. Comparison of the effects on neuroprotection and vascular regeneration after 20m-MCAO between Wt control mice, hydralazine-administrated mice, and the low and high concentration lines of AM-Tg

Mice	Infarct area (mm ² /field)	Brain blood flow (% Contralateral)	Systolic BP (mm Hg)
Control	0.90 ± 0.09	80.8 ± 2.3	120.1 ± 2.2
Hydralazine	0.94 ± 0.17 ^{ns}	79.6 ± 2.6 ^{ns}	101.0 ± 3.9 ^a
Low-conc. AM-Tg	0.58 ± 0.12 ^b	88.4 ± 2.9 ^b	105.1 ± 1.8 ^a
High-conc. AM-Tg	0.67 ± 0.09 ^b	86.3 ± 2.0 ^b	106.4 ± 3.5 ^a

conc., Concentration.

^a *P* < 0.01; ^b *P* < 0.05; ns, not significant *vs.* control; n = 6.

Fig. 10. Brain edema in acute phase, neuronal loss and gliosis in subacute to chronic phase after 20m-MCAO were reduced in AM-Tg mice. Furthermore, vascular regeneration, mobilization of CD34⁺ mononuclear cells and subsequent neurogenesis were enhanced in them. These effects resulted in improved recovery of motor function after the nonfatal stroke. AM was also found to exert direct antiapoptotic and neuro-differentiating effects on neuronal cells *in vitro*. Exogenous administration of AM in mice after 20m-MCAO also

**FIG. 7.** Effects of AM on infarct size and brain edema in the fatal stroke, 2 h-MCAO. A, Comparison of infarct size between Wt and AM-Tg with 2,3,5-triphenyltetrazolium chloride staining at 4.0 mm from the frontal pole. White area represents infarction. B and C, Infarct (B) and edema (C) volumes quantified 24 h after the operation of 2 h-MCAO. *, *P* < 0.05; ns, not significant for Wt and AM-Tg; n = 9. D, Representative image of *in situ* Evans Blue leakage into the ischemic core at 24 h after 2 h-MCAO. E, Quantification of Evans Blue in the ischemic brain. **, *P* < 0.01; n = 4.

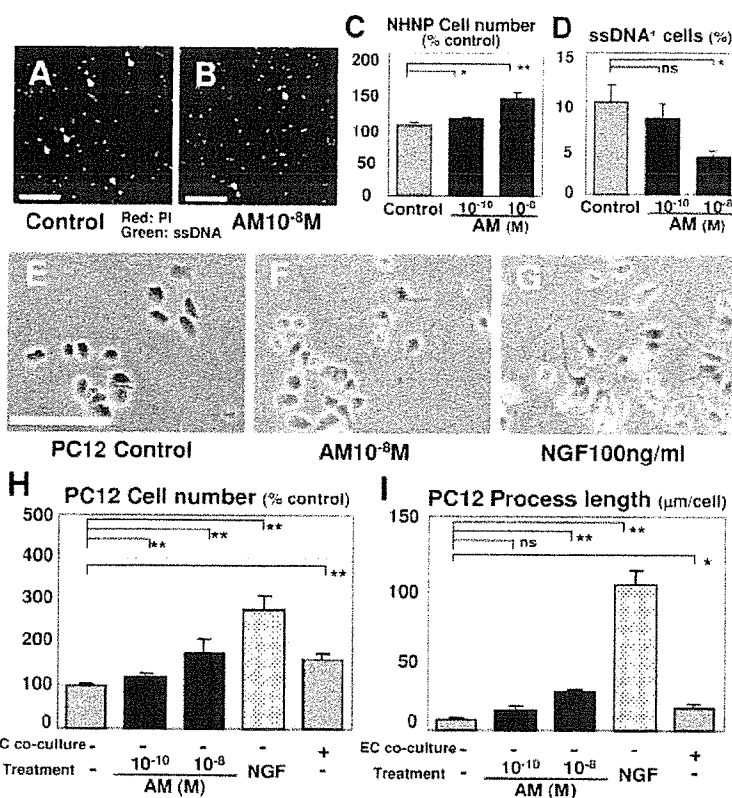


FIG. 8. Effects of AM *in vitro* on apoptosis of NHNP neuronal progenitor cells and neuronal differentiation of PC12 cells. A–D, *In vitro* analysis of apoptotic NHNP after incubation with (B) or without (A) AM. NHNP cell number (C) and the ratio of ssDNA⁺ cells to total cells (D) after 48 h incubation. *, $P < 0.05$; **, $P < 0.01$; ns, not significant *vs.* control; $n = 4$; scale bar, 100 μm . E–G, Effects of AM on neuronal differentiation of PC12 cells evaluated by the length of neuronal process. Microscopic examination of PC12 cells after incubation for 7 d (E). AM (F) or nerve growth factor (G) was added to the culture medium. Quantification of cell number (H) and the length of neuronal process (I). *, $P < 0.05$; **, $P < 0.01$; ns, not significant; $n = 6$; scale bar, 100 μm .

reduced the infarct area, and promoted vascular regeneration and functional recovery.

Stroke causes two different types of neuronal death: necrosis and apoptosis. Acute neuronal loss, which is completed within a few days after ischemic damage, is necrotic, whereas delayed neuronal loss, which may start several days after transient ischemia, is considered to be apoptotic (27, 28). Many studies have found that treatments that reduce inflammation or oxidative stress are beneficial for the prevention of apoptotic neuronal loss (29, 30).

In this study, we demonstrated that AM exerts neuroprotective actions in the ischemic brain. A significant reduction in neuronal loss in AM-Tg mice after 20m-MCAO became obvious after postoperative d 7, but was not obvious before d 3. A significant decrease in ssDNA-positive cells inside and

on the border of the ischemic area was observed in AM-Tg mice in association with a reduction in CD45⁺ cells and *in situ* ROS production in the subacute phase. AM is therefore assumed to reduce delayed neuronal loss through suppression of the apoptotic process. Furthermore, we confirmed that AM directly suppresses apoptosis of neuronal progenitor cells *in vitro*. These findings suggest that AM exerts neuroprotective effects on the ischemic brain by reducing apoptotic neuronal loss through both its direct antiapoptotic action on neurons and indirect effect via antiinflammation and anti-ROS production. Consistent with the findings in this study, several recent reports have provided evidences for the organ-protective effects of AM against inflammation and oxidative stress (31–33). In addition, we found significant negative correlation between capillary density and apoptotic cells in the same section on postoperative d 7 after 20m-MCAO. Moreover, the infarct area kept expanding between d 7–28 in Wt mice, whereas AM-Tg mice did not show the increase in size in this period. These findings suggest that the increased blood flow in AM-Tg mice was one of the causes of neuroprotection after 20m-MCAO, although we suppose that multiple actions of AM, as described above, could also contribute for neuroprotection.

Increased vascularity is reported to be associated with improved neurological recovery in human patients with stroke (34). This implies that physiological vascular regeneration in the ischemic brain constitutes a beneficial response for the recovery of impaired neurological function. Moreover, neurogenesis after stroke even in adulthood has been demonstrated to occur in a place surrounded by the vascu-

TABLE 4. Effects of AM-antagonists, PKA inhibitors, and PI3K inhibitors on AM-induced neural differentiation of PC12 cells

	Process length (μm/cell)
PC12	6.8 ± 1.7
+AM (10 ⁻⁸ mol/liter)	23.6 ± 4.0 ^a
+AM+AM(22–52) (10 ⁻⁵ mol/liter)	11.8 ± 3.4 ^b
+AM+CGRP(8–37) (10 ⁻⁵ mol/liter)	14.8 ± 1.9 ^c
+AM+Rp-cAMP (10 ⁻⁵ mol/liter)	10.2 ± 2.7 ^b
+AM+PKA Inh (10 ⁻⁶ mol/liter)	7.2 ± 2.3 ^b
+AM+LY294002 (10 ⁻⁵ mol/liter)	4.6 ± 1.6 ^b
+AM+wortmannin (10 ⁻⁷ mol/liter)	5.4 ± 1.1 ^b
PC12-EC coculture	20.7 ± 2.1 ^a

EC, Endothelial cell.

^a $P < 0.01$ *vs.* PC12 without AM; ^b $P < 0.01$ *vs.* PC12 with AM (10⁻⁸ mol/liter); ^c $P < 0.05$; $n = 8$.