

investigation included Thy-1, owing to its characteristics as a cell surface marker expressed on vascular endothelial cells (EC) that are upregulated by the inflammatory cytokines interleukin (IL)-1 β and tumor necrosis factor (TNF)- α .¹⁸ Endostatin is an endogenous angiogenesis inhibitor that is activated by proteases such as those derived from inflammatory cells.^{19,21} The antigen Ki-67 was used as a nuclear marker of cell proliferation.²²

METHODS

PATIENTS AND TREATMENTS

We retrospectively reviewed 38 eyes of 38 consecutive AMD patients in different centers who were surgically treated with full macular translocation surgery with CNVM extraction. All patients had clinically active disease and progression. Fourteen CNV patients without any kind of therapy prior to surgery composed the control CNV group. The clinical characteristics of these 14 patients are summarized in **Table 1**. Twenty-four patients received surgery following intravitreal bevacizumab injection (bevacizumab CNV group). Therapy options, including observation, conventional thermal laser photocoagulation, verteporfin PDT, intravitreal injection of triamcinolone acetonide or anti-VEGF agents, and full macular translocation with 360° retinotomy and CNVM extraction were discussed with the patients. Although submacular removal of CNV was not favored by the randomized submacular surgery trials, submacular surgery trial Group B revealed that submacular surgery for predominantly hemorrhagic CNV did reduce the risk of severe visual acuity loss (loss of ≥ 6 Early Treatment Retinopathy Study lines) and contrast sensitivity loss (≥ 3 segments, using the Pelli-Robson chart at 0.5 m) significantly, in comparison to the group observed without any treatment for 24 months of follow-up.²³ Macular translocation of the retina, a complex but potentially effective surgical method, was reported to be beneficial when CNV was related to submacular hemorrhage²⁴ or RPE tear.²⁵ Furthermore, improvement in distance and near visual acuity as well as in reading speed was detected at the 1-year follow-up of patients with neovascular AMD who were treated with macular translocation.^{26,27}

Intravitreal bevacizumab injection and surgical intervention with full macular translocation surgery were offered when (1) visual acuity was worse than 20/200 (being the minimum visual acuity to recommend the first PDT, according to the Treatment of Age-Related Macular Degeneration with Photodynamic Therapy investigation^{28,29}); (2) CNV was related to retina pigment epithelium (RPE) tear, chorioretinal anastomosis, or massive subretinal/intravitreal hemorrhage; (3) visual deterioration progressed and subretinal hemorrhage and RPE tear appeared following treatment; or (4) reading ability was recently lost, despite improvement in distant visual acuity after previous treatment. The clinical characteristics of the patients treated with intravitreal bevacizumab are summarized in **Table 2**. Reduction of intraoperative bleeding was intended with the preoperative injection of bevacizumab, as suggested in other ocular pathologies.^{30,31}

Each patient gave written informed consent regarding their knowledge of the off-label use and experimental nature of the treatment procedure, and the risks and benefits of all therapeutic options were discussed in detail. The study followed the guidelines of the declaration of Helsinki. Local institutional review board approval was granted for allocation and histological analysis of the specimens.

TISSUE PREPARATION AND IMMUNOHISTOLOGY

Within minutes after surgery, excised CNVM were fixed in 3.7% formalin and embedded in paraffin. After serial sections were de-

Table 1. Clinical Characteristics of Patients Without Any Kind of Preoperative Therapy

Patient, No./Sex/Age, y	Eye	CNV Type
1/F/71	Right	Hemorrhagic
2/F/80	Right	Occult with hemorrhage
3/F/75	Left	Predominantly classic
4/M/75	Right	Classic with hemorrhage
5/F/75	Right	Classic with hemorrhage
6/F/76	Right	Hemorrhagic
7/M/80	Left	Hemorrhagic
8/M/80	Left	Hemorrhagic
9/M/72	Left	Classic with hemorrhage
10/F/73	Right	Hemorrhagic
11/F/74	Left	Occult with hemorrhage
12/F/81	Right	Hemorrhagic
13/M/80	Left	Hemorrhagic
14/M/64	Right	Occult

Abbreviation: CNV, choroidal neovascularization.

paraffinized and rehydrated, antigen retrieval was accomplished by proteolytic digestion with 0.5% protease XXIV (Sigma, St Louis, Missouri) for cytokeratin 18, endostatin, and Thy-1, and by heat treatment in citrate buffer in a pressure cooker under 120°C for 2 minutes for Ki-67, CD45, CD68, ICAM-1, and E-selectin.

Immunohistochemical staining with the primary mouse monoclonal antibodies specific for cytokeratin 18 (Progen, Heidelberg, Germany), ICAM-1 (Novocastra, Newcastle upon Tyne, England), E-selectin (Novocastra), CD45 (Dako, Glostrup, Denmark), and Ki-67 (Clone Ki-55; Dako) was performed using the avidin-biotin complex horseradish peroxidase method, as previously described.³⁴ For E-selectin, cytokeratin 18, and ICAM-1 staining, the brown chromogen 3-diaminobenzidine was replaced with 3-amino-9-ethylcarbazole, a highly sensitive substrate chromogen (Code K3461; Dako).

Immunohistochemical staining for Thy-1, CD68, and endostatin was performed by the alkaline-phosphatase method, according to the manufacturer's instructions (ChemMate Detection Kit, Alkaline Phosphatase/RED, Rabbit/Mouse, K5005; Dako), as previously described,³⁴ using primary antihuman antibodies specific for Thy-1 (mouse anti-CD90 antibody; BD Biosciences Pharmingen, San Jose, California), CD68 (mouse, monoclonal; Dako), and endostatin (rabbit, polyclonal; Dianova GmbH, Hamburg, Germany). Color was developed using chromogen red (ChemMate Detection Kit; Dako). Hematoxylin (Chemmate, Code S2020; Dako) was used for counterstaining.

For negative controls, primary antibodies were substituted by the appropriate normal sera or omitted.

ANALYSIS

Serial sections from a specimen were analyzed by light microscopy.

Immunoreactivity for ICAM-1, E-selectin, CD68, and endostatin was analyzed semiquantitatively and separately in the RPE, EC (in the perivascular area for CD68), and stroma. A grading scheme indicating the degree of labeling was used: numerals 3, 2, 1, and 0 were assigned for intense (70%-100% positive cells), moderate (40%-69% positive cells), weak (1%-39% positive cells), and absent of staining, respectively.

Each specimen was documented under $\times 50$ magnification with a Zeiss microscope (Axioskop; Carl Zeiss, Inc. Oberkochen, Germany) connected to a digital camera (Fujix HC-300Z; Fujifilm, Tokyo, Japan). The area of each specimen was measured using the appropriate hardware and software (Axio-Vision, Version 3.1, Carl Zeiss, Inc. Göttingen, Germany).

Table 2. Clinical Characteristics of Patients Treated With Intravitreal Bevacizumab Before Submacular Surgery

Patient, No./ Sex/Age, y	Eye	CNV Type	No. of Bevacizumab Injections	Time to Surgery From Treatment, d	
				PDT	Intravitreal Bevacizumab
1/F/80	Left	Hemorrhagic	1	-	1
2/F/86	Left	Hemorrhagic	1	-	3
3/M/75	Right	Occult hemorrhagic with RPE tear	1	-	3
4/M/69	Left	Occult	1	-	4
5/M/69	Left	Occult	1	-	4
6/F/71	Right	Hemorrhagic	1	-	6
7/M/83	Right	Hemorrhagic PED	1	-	6
8/M/79	Left	Predominantly classic with CRA	1	-	7
9/M/74	Right	Hemorrhagic	1	-	9
10/F/69	Right	Hemorrhagic with RPE tear	1	-	11
11/F/73	Left	RPE tear	1	-	12
12/F/72	Right	Hemorrhagic occult	1	39	12
13/M/73	Left	RPE tear	1	-	26
14/M/58	Right	Hemorrhagic with RPE tear	1	-	43
15/F/79 ^a	Right	Hemorrhagic with RPE tear	1	343	43
16/F/55	Right	Occult with PED	1	-	44
17/F/81	Right	Hemorrhagic with RPE tear	1	-	46
18/M/79	Left	Predominantly classic	1	64	48
19/M/78	Left	Occult hemorrhagic	5	249	209/167/125/92/50
20/F/76	Left	Hemorrhagic	1	-	53
21/M/79	Right	Occult with RPE tear	2	-	95/55
22/M/75	Right	Occult	1	-	57
23/F/84	Right	Hemorrhagic with PED	2	-	135/85
24/F/75	Right	Predominantly classic	2	-	154

Abbreviations: CNV, choroidal neovascularization; CRA, chorioretinal anastomosis; PED, pigment epithelial detachment; PDT, photodynamic therapy; RPE, retina pigment epithelium.

^aPatient was treated with an intravitreal triamcinolone acetonide injection 342 days before surgery.

All Ki-67-positive nuclei, macrophages, and leukocytes were counted in each specimen. Proliferative activity, density of leukocytes, and macrophage infiltration in each specimen were determined quantitatively by calculating the ratio of the total number of Ki-67 positive nuclei, macrophages, and leukocytes in the CNVM to the total area of the membrane.

Thy-1 expression was determined by the percentage of Thy-1-expressing vessels in the overall vascularization of each membrane.

Intensity of CD68, ICAM-1, E-selectin, and endostatin immunostaining, density of leukocyte and macrophage infiltration, proliferative activity, and Thy-1 expression in the described CNV groups were compared using the Mann-Whitney *U* test. Immunohistologic findings in the bevacizumab CNV group were analyzed based on timing or number of bevacizumab injections, using the Spearman correlation test. A *P* value less than .05 was considered significant.

RESULTS

Immunohistological findings are summarized in **Figure 1**

HISTOPATHOLOGICAL FINDINGS IN THE CONTROL CNV GROUP

Retina pigment epithelial cells immunoreactive for cytokeratin 18 were found in all specimens.

All but 1 membrane was vascularized. Immunohistological stains of the specimen from a patient with massive subretinal hemorrhage (**Figure 2A**) are shown in Figure 2.

Expression of E-selectin was found in the RPE of all CNVM, and intensely in 4 (28.6%) membranes. E-selectin was detected in the EC of 8 CNVM (57.1%) and in the stroma of 9 CNVM (64.3%) (Figures 1A and 2B).

Expression of ICAM-1 was prominent in the RPE; it was intense in 13 CNVM (92.9%). No ICAM-1 was present in the EC of 9 CNVM (64.3%). Intense ICAM-1 expression was detected in the EC of 3 (21.4%) and in the stromal cells of 2 CNVM (14.3%) (Figures 1B and 2C).

Leukocytes detected by CD45 immunostaining were found to be present in all but 1 CNVM. The number of leukocytes in a 1-mm² area of CNVM varied between 0 and 779.6 (median, 106.64 cells/mm²) (Figure 1C). The specimen in Figure 2D has the fourth highest density of leukocyte infiltration (157.39 cells/mm²) in the control CNV group.

The density of macrophages ranged between 183.94 cells/mm² and 2826.73 cells/mm² (median, 882.66 cells/mm²) (Figure 1C). Macrophage infiltration was seen in the RPE cell layer, the perivascular area, and the stroma in 85.7% (n=14), 61.5% (n=13), and 92.8% (n=14) of the specimens. Intense macrophage infiltration was detected in the RPE cell layer of 6 (42.8%), the perivascular area of 2 (14.3%), and the stroma of 4 (28.6%) CNVM (Figures 1D and 2E).

Vessels expressing Thy-1 were detected in 13 of 14 specimens (92.8%), in varying percentages of vessels (median, 80.0%; range, 0%-100%). In 4 (28.6%) CNVM, all vessels displayed Thy-1 (Figure 2F).

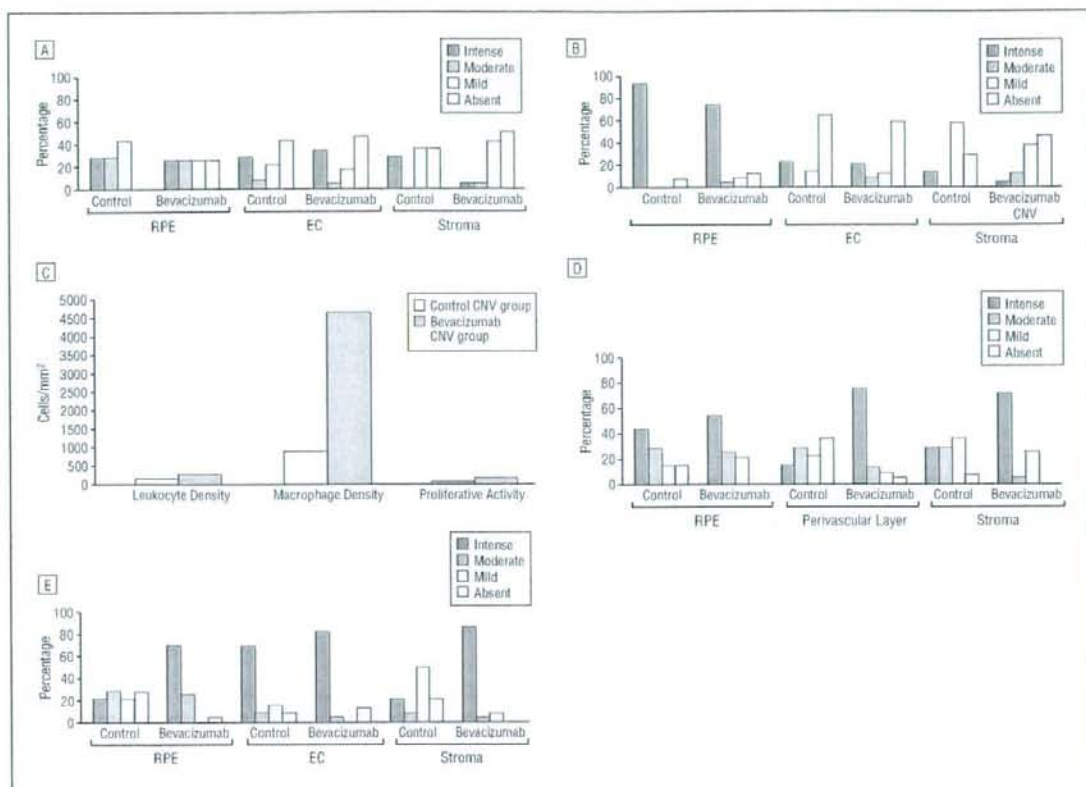


Figure 1. Graphs showing E-selectin (A) and intercellular adhesion molecule (ICAM)-1 (B) immunostaining intensity, density of leukocytes and macrophages (CD68) and proliferative activity (C), CD68 immunoreactivity (D), and intensity of endostatin staining (E) in choroidal neovascular membranes (CNVM) without any kind of previous therapy (control choroidal neovascularization [CNV] group) and CNVM excised after intravitreal injection of bevacizumab (Avastin; Genentech, Inc, South San Francisco, California) (bevacizumab CNV group). Expression of ICAM-1, E-selectin, CD68, and endostatin immunostaining in retina pigment epithelium (RPE), endothelial cells (EC) (perivascular area for CD68), and stroma were evaluated separately and semiquantitatively as Intense (70%-100% positive cells), moderate (40%-69% positive cells), mild (1%-39% positive cells), or absent (A,B,D,E). All Ki-67-positive nuclei, CD45-immunoreactive leukocytes, and CD68-immunopositive macrophages were counted in each specimen. Proliferative activity and density of leukocytes and macrophages were determined separately for each CNVM by the ratio of total number of proliferating cells, leukocytes, and macrophages to the total area of the membrane (mm^2); their median values were shown in (C).

Immunoreactivity to endostatin was detected in the RPE-Bruch membrane complex in 10 (71.4%) membranes. Endostatin was immunonegative in vessels in only 7.7% ($n=1$) of the specimens. Within the stroma, endostatin was expressed by fibroblast-like and inflammatory cells in 11 (78.6%) specimens (Figures 1E and 2G).

Proliferative activity varied between 6.67 and 514.08 Ki-67-expressing nuclei/ mm^2 (median, 34.34 nuclei/ mm^2) (Figure 1C). Proliferating cells belonged mostly to inflammatory infiltrate within the stroma (Figure 2H). In 12 (85.7%) CNVM, proliferative activity was less than 160 Ki-67-expressing nuclei/ mm^2 . The specimen in Figure 2H has the second highest proliferative activity among the control CNV group (511.05 nuclei/ mm^2).

HISTOPATHOLOGICAL FINDINGS IN THE BEVACIZUMAB CNV GROUP

In the bevacizumab CNV group (Figure 3A), the RPE had E-selectin in 18 (75.0%) CNVM. The EC were E-selectin immunonegative in 11 (45.8%) CNVM. Intense E-selectin staining was detected in the RPE of 6 (25%)

and the EC of 8 (33.3%) CNVM. Stromal cells expressed E-selectin in 12 (50.0%) CNVM (Figures 1A and 3B). Expression of E-selectin in the bevacizumab CNV group was comparable to the control CNV group.

Immunoreactivity for ICAM-1 was detected in the RPE of 21 CNVM (87.5%) and intensely in 18 CNVM (75%). Intense ICAM-1 expression was found in the EC of 5 (20.8%) and stromal cells of 1 (4.2%) CNVM (Figures 1B and 3C). Expression of ICAM-1 was not significantly different from the control CNV group.

Leukocytes were found in all specimens (median, 271.61 cells/ mm^2 ; range, 33.50-2351.01 cells/ mm^2) (Figure 3D). The density of leukocytes seemed higher than in the control group, but without statistical significance (Figure 1C, $P=.07$).

Macrophages were present in all CNVM (median, 4661.95 cells/ mm^2 ; range, 321.01-61821.16 cells/ mm^2) (Figures 1C and 3E). The density of macrophages was significantly higher than that of the control CNV group ($P<.001$). Moderate to strong macrophage infiltration was detected within the RPE cell layer and in the perivascular area in 18 (75%) specimens and in the stroma in 17 (70.8%)

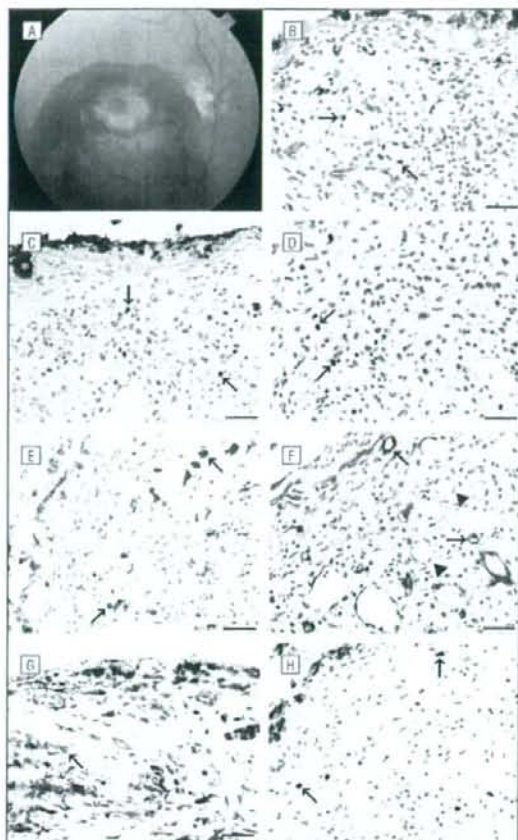


Figure 2. Color fundus (A) and histologic photomicrographs (B-H) of a choroidal neovascularization membrane (CNVM) extracted without any kind of therapy before surgery. Note the massive subretinal hemorrhage (A). Sections were probed with antibodies against E-selectin (B), intercellular adhesion molecule (ICAM)-1 (C), CD45 (common leukocyte antigen; D), CD68 (E, macrophages), Thy-1 (F), endostatin (G), and Ki-67 (H). Sections were stained with 3-amino-9-ethylcarbazole red chromogen (B, C), 3-diaminobenzidine resulting in a brown chromogen (D, H), and red chromogen (E-G). Hematoxylin was used as a counterstain. B, E-selectin expression is prominent in endothelial cells (EC, arrows). ICAM-1 expression (C) was detected in the retina pigment epithelium (RPE, asterisk) and stromal cells (arrows). Leukocytes (D, arrows) and macrophages (E, arrows) were present in perivascular area and stroma. Besides Thy-1-expressing vessels (F, arrows), some Thy-1-immunonegative vessels (arrowheads) were found in the same specimen. Endostatin (G) was found to be weakly present in some vessels (arrow) whereas the RPE (asterisks) was endostatin immunonegative. H, Some proliferating cells (arrows) were present in the CNVM. Scale bars = 50 μ m.

CNVM (Figure 1D). Immunoreactivity of CD68 was significantly more intense in the bevacizumab CNV group in the perivascular area (median, 3) and stroma (median, 3) than in the control CNV group (median, 1; $P < .001$ and median, 2; $P = .03$, respectively).

In the CNVM of all patients treated with bevacizumab, all vessels were lined with EC displaying Thy-1 (median, 100.0%) (Figure 3F). The percentage of Thy-1-expressing vessels was significantly higher in the bevacizumab CNV group ($P < .001$).

The RPE-Bruch membrane complex displayed endostatin in all but 1 CNVM either strongly (70.8%; $n = 17$)

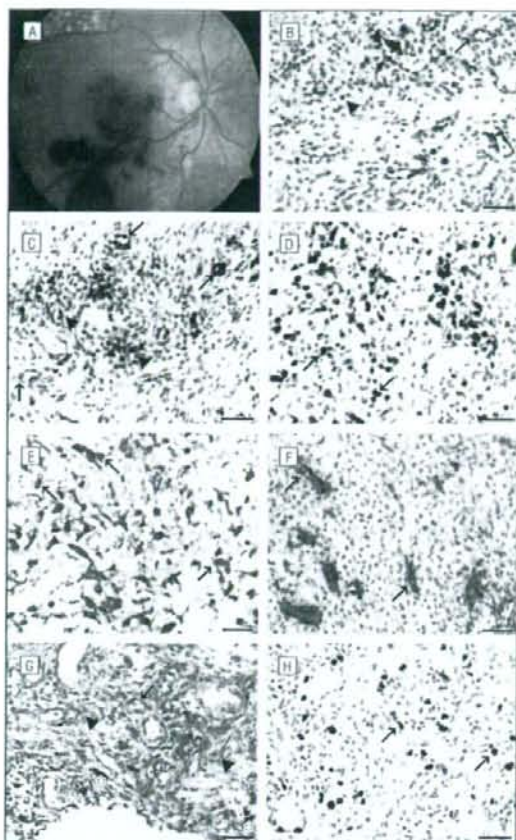


Figure 3. Histologic photomicrographs of the choroidal neovascularization membranes extracted 4 days after intravitreal bevacizumab injection. Note the massive subretinal hemorrhage with pigment epithelium detachment (A). Sections were probed with antibodies against E-selectin (B), intercellular adhesion molecule (ICAM)-1 (C), CD45 (common leukocyte antigen; D), CD68 (macrophages; E), Thy-1 (F), endostatin (G), and Ki-67 (proliferating cells; H). Sections were stained with 3-amino-9-ethylcarbazole red chromogen (B, C), 3-diaminobenzidine resulting in a brown chromogen (D, H), or red chromogen (E-G). Hematoxylin was used as counterstain. Endothelial cells (arrows) and stromal cells (arrowheads) showed E-selectin (B) and ICAM-1 (C). Inflammatory infiltration with many leukocytes (D, arrows) and many macrophages (E, arrows) was detected within the perivascular area and stroma. Expression of Thy-1 was intensely expressed in vascular structures (F, arrows). G, Endostatin was intensely immunopositive in vessels (arrow), stromal cells (arrowheads), and retina pigment epithelium (asterisks). Many cells within the stroma expressed Ki-67 (H, arrows). Scale bars = 50 μ m.

or moderately (25.0%; $n = 6$). Endostatin was intensely expressed in the vessels of 20 CNVM (87.3%). Stromal cells displayed endostatin in all CNVM and strongly in 21 (87.5%) membranes (Figures 1E and 3G). Endostatin was significantly stronger in the RPE-Bruch membrane complex (median, 3; range, 2-3; $P < .001$) and stroma (median, 3; range, 1-3; $P < .001$) of the bevacizumab CNV group than the control CNV group (median, 1.5; range, 0-3 and median, 1; range, 0-3, respectively).

Proliferative activity (median, 204.33 cells/mm²; range, 3.61-1886.135 cells/mm²) in the bevacizumab CNV group was significantly higher than that in the control CNV group ($P = .007$) (Figures 1C and 3H).

Table 3. Evaluation of Histopathological Findings in Bevacizumab CNV Group With Regard to Application of Bevacizumab and PDT^a

	Based on Timing of Injections		Based on Frequency of Injections		Based on Previous Treatment				P Value
					With PDT		Without PDT		
	P Value	ρ	P Value	ρ	Median	Range	Median	Range	
E-selectin									
RPE	.07	0.38	.62	-0.10	1	1-3	2	0-3	>.99
EC	.39	0.18	.73	0.07	2	1-3	0	0-3	.04
Stroma	.86	-0.04	.93	-0.02	0.5	0-1	0.5	0-3	.86
ICAM-1									
RPE	.81	0.05	.88	0.03	3	0-3	3	0-3	.88
EC	.71	0.08	.76	0.07	0	0-2	0	0-3	.39
Stroma	.81	0.05	.57	-0.12	0.5	0-1	1	0-3	.61
Density of leukocytes	.25	-0.24	.86	-0.04	107.83	50.49-165.12	383.77	33.50-2351.10	.09
Density of macrophages	.64	-0.099	.96	0.01	4085.5	321.01-5078.1	4951.91	1308.57-61821.16	.28
Thy-1	>.99		>.99		100	100-100	100	100-100	>.99
Endostatin									
RPE	.94	0.02	.20	0.27	3	3-3	3	2-3	.28
EC	.35	0.21	.50	0.15	3	2-3	3	0-3	.27
Stroma	.39	0.18	.42	0.17	3	1-3	3	1-3	.38
Proliferative activity	.09	-0.36	.22	-0.26	128.72	3.61-198.73	160.19	25.12-1886.14	.28

Abbreviations: CNV, choroidal neovascularization; CNVM, choroidal neovascular membrane; EC, endothelial cells; ICAM, intercellular adhesion molecule; PDT, photodynamic therapy; RPE, retina pigment epithelium.

^aIntensity of E-selectin, ICAM-1, and endostatin immunostainings in RPE, EC, and stroma. Density of leukocyte and macrophage infiltration, proliferative activity, and Thy-1 expression in bevacizumab CNVM were analyzed based on timing or number of bevacizumab injections with the Spearman correlation test and compared based on previous application of verteporfin PDT using the Mann-Whitney *U* test. *P* < .05 was considered significant.

Immunohistologic findings in the bevacizumab CNV group did not show major differences based on either timing or number of bevacizumab injections (**Table 3**). Based on previous application of PDT, only E-selectin expression showed a major difference, being significantly stronger in the EC of patients in the bevacizumab CNV group treated with PDT previously (*n* = 4) than those in the bevacizumab CNV group without any previous PDT (*n* = 20) (Table 3).

COMMENT

Vascular endothelial growth factor is expressed in the choroid of human eyes,^{35,36} and polarized secretion of VEGF in the RPE is vital for choriocapillaris development and maintenance.³⁷ In AMD, various factors such as free radicals or complement components may upregulate VEGF expression.^{38,39} Vascular endothelial growth factor, chemotactic for inflammatory cells, activates monocytes and macrophages and induces expression of ICAM-1 and E-selectin on vascular EC to facilitate adhesion and migration of inflammatory cells.^{15-18,20,40} Macrophages synthesize the cytokines IL-1 and TNF- α ,^{20,41-43} which enhance ICAM-1 expression in EC and in the RPE,⁴⁴⁻⁴⁶ and induce inflammatory cell infiltration further. There is a closed but amplifying circuit between VEGF and inflammatory cells, because leukocytes and macrophages can produce VEGF themselves and also induce VEGF production in the RPE through IL-1 and TNF- α in human CNVM.^{13,20,47-49} Vascular endothelial growth factor, in turn, stimulates growth and maintenance of the CNVM.^{12,50-52} Because VEGF and inflammation seem to closely interact with each other, VEGF inhibition by bevacizumab

might influence, or possibly inhibit, the inflammatory activity in CNV.

In the specimens treated with bevacizumab, expression of ICAM-1 and E-selectin was comparable to that of the control CNV cases without previous therapy. However, density of leukocytes, and especially of macrophages, was higher in the bevacizumab CNV group than in the control CNV group. Consequently, the percentage of Thy-1-expressing vessels, a mirror of TNF- α and IL-1 activity, was also significantly higher. Macrophages, reported to be present in 60% of the surgically excised CNVM,¹² seem to play a causative role in neovascular AMD pathogenesis.^{48,49} In aged RPE cells, accumulation of oxidation products seems to induce angiogenic and chemotactic factors like monocyte chemoattractant protein 1.⁵³ In healthy human eyes, recruited macrophages may have a physiological role in removing the accumulated subretinal debris. However, excess oxidized lipids in the eyes of patients with AMD were suggested to upregulate the release by macrophages of proteolytic enzymes that damage the Bruch membrane and contribute to CNV development.⁵⁴ Macrophages, in addition to stimulating VEGF, can induce EC proliferation and migration through the release of cytokines, proteases, and growth factors.²⁰ Therefore, experimental studies for potential new antiangiogenic treatment strategies target the complement system,⁵⁵ ICAM,⁵⁶ macrophages, monocytes,^{48,49} and TNF.⁵⁷ These were shown to inhibit leukocyte infiltration and VEGF expression and, therefore, inhibit CNV development or progression in experimental studies.⁴⁸⁻⁴⁹ In experimental models, early intravitreal injection of VEGF-A antibody after CNV induction was shown to inhibit CNV and choroidal macrophage in-

filtration through suppression of monocyte chemoattractant protein 1 expression in the RPE-choroid complex.⁵⁸ Hence, anti-VEGF drugs might be expected to have an anti-inflammatory effect.⁵⁹

Our results were truly unexpected. Inhibition of VEGF with bevacizumab was related to a prominent inflammatory reaction, instead of an anti-inflammatory effect, in comparison to the control CNV group. This might be for several reasons.

First, the Fc portion of bevacizumab in CNV may stimulate a cellular immune response through Fc receptors in immune cells. Bevacizumab was shown to penetrate the retina in an experimental study.⁶⁰ Anatomical defects in the Bruch membrane created by proteolytic enzymes and CNV itself might facilitate the passage of bevacizumab molecules through the RPE to the CNVM in the eyes of patients with AMD.

Second, VEGF acts as a survival factor for EC by inhibiting apoptosis.^{37,61} Deprivation and/or inhibition of VEGF was shown to induce vascular regression through EC apoptosis, especially in neovascular vessels.⁶¹⁻⁶⁵ During vascular regression, macrophages regulate EC and pericyte apoptosis in capillaries. Leukocytes release human neutrophil peptides that are proapoptotic for EC.⁶⁶ Macrophages also recognize and phagocytose apoptotic cells. Macrophage depletion results in EC survival and persistence of functional capillaries.^{67,68} Therefore, increased inflammatory cell infiltration following bevacizumab injection may reflect part of the vascular regression process in CNV, based on debris removal by macrophages.

Third, macrophages might be involved in CNV inhibition. In mice deficient in monocyte chemoattractant protein 1 or its receptor, accumulating complement factors or IgG stimulated VEGF expression in the RPE and, therefore, CNV development.³⁹ In another experimental study, inhibition of macrophage entry into the eye stimulated CNV, whereas direct injection of macrophages inhibited CNV.⁷⁰ Previously, macrophage depletion with liposomal clodronate was shown to inhibit CNV development.^{48,49} However, reduction of neovascularization in these studies was recently suggested to be due to the direct toxicity of liposomes on EC rather than macrophage depletion itself.⁷⁰

Last, inflammatory cells are known to contribute to antiangiogenesis through activation of endogenous angiogenesis inhibitors such as endostatin. Macrophages and leukocytes release proteases such as matrix metalloproteases.²⁰ Proteases cleave endostatin that is bound to collagen XVIII in vascular basement membranes and the Bruch membrane. Through this cleavage, endostatin is released and activated.²¹ In our series, endostatin was significantly more intense in the RPE and stroma of the inflammatory active bevacizumab CNV group than in the control CNV group. This has important consequences because endogenous endostatin was shown to inhibit CNV.¹⁹ Furthermore, endostatin was decreased in the RPE and choroid of human eyes with AMD, and its deficiency was suggested to predispose CNV formation.⁷¹

Vascular endothelial growth factor was shown to be an important mitogenic factor for choroidal EC.^{37,47,72,73}

and also for RPE cells.⁷⁴ However, whether anti-VEGF toxins or antibodies inhibit proliferation of human choroidal and RPE cells is controversial in *in vitro* settings.^{75,77} Recently, we have shown the antiproliferative effect of bevacizumab on EC in *in vitro* studies.^{74,79} Still, *in vitro* results may not reflect the ongoing process *in vivo*. In the bevacizumab CNV group, proliferative activity was significantly higher than in the control CNV group. Proliferating cells were located within the stroma and originated from the inflammatory infiltrate rather than the RPE or EC. In our previous work, proliferative activity was correlated with inflammatory activity, and was significantly higher in inflammatory active specimens than in inflammatory inactive ones.^{34,80}

In our series, timing of the bevacizumab injection prior to surgery varied between 1 and 154 days. Four of the 24 patients were treated with 2 to 5 bevacizumab injections preoperatively. However, immunohistologic findings in the bevacizumab CNV group did not show major differences based on timing or number of injections. Four of the patients were treated with PDT 39 to 343 days before bevacizumab injection. Of the antigens investigated, only E-selectin was found to be significantly more intense in the EC of the bevacizumab CNV patients treated previously with PDT. However, our previous studies³⁴ have shown that immunohistopathological findings dramatically change early (3 days) after PDT when compared with either longer intervals following PDT or the control CNV group. The implications of PDT-bevacizumab combination therapy on inflammation and angiogenesis, therefore, need to be evaluated in a higher number of samples excised early after combination therapy.

To our knowledge, this is the first clinicopathological study evaluating inflammation and proliferation in human CNV treated with an anti-VEGF agent. To interpret the results properly, the following limitations need to be taken into account: first, histopathological specimens reflect only a time window within a dynamic process. Therefore, we selected homogenous control and treatment groups that were comparable in their clinics. Second, in contrast to experimental studies, time between treatment and tissue acquisition varies. Nevertheless, data retained by histopathological examination of human samples may be closer to the truth than any data collected by *in vitro* or *in vivo* experiments.⁸¹

The results of this study, revealing high inflammatory and proliferative activity, were truly unexpected and appear to be, to some extent, paradoxical to the inhibition of angiogenesis. The question is how to interpret this data in view of promising clinical results related to a decrease in disease activity following intravitreal injection of bevacizumab. There is growing knowledge that inflammatory reactions exert not only proangiogenic activities, but are also involved in vascular regression and angiogenesis inhibition. We assume that a similar process may be reflected in our histopathological results. Our results reveal enhanced expression of the endogenous angiogenesis inhibitor endostatin in these inflammatory active bevacizumab CNVM. Still, our understanding of treatment modalities and combination therapies needs to be further evaluated to implement the optimal treatment

strategies and protocols for neovascular AMD. High inflammatory and proliferative activity as well as enhanced endostatin expression in bevacizumab-treated CNV should be considered when combination therapies with bevacizumab are planned.^{11,82,83}

Submitted for Publication: July 19, 2007; final revision received December 17, 2007; accepted December 22, 2007.

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Author Contributions: Olcay Tatar and Salvatore Grisanti had full access to all of the data in the study and take responsibility for the integrity of the data and the accuracy of the data analysis.

Funding/Support: This work was supported by grants from the Vision 100 Foundation and the Jung Foundation.

Financial Disclosure: None reported.

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ORIGINAL ARTICLE

Angiotensin II Type 1 Receptor Signaling Contributes to Synaptophysin Degradation and Neuronal Dysfunction in the Diabetic Retina

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OBJECTIVE—Pathogenic mechanisms underlying diabetes-induced retinal dysfunction are not fully understood. The aim of the present study was to show the relationship of the renin-angiotensin system (RAS) with the synaptic vesicle protein synaptophysin and neuronal activity in the diabetic retina.

RESEARCH DESIGN AND METHODS—C57BL/6 mice with streptozotocin-induced diabetes were treated with the angiotensin II type 1 receptor (AT1R) blocker telmisartan or valsartan, and retinal function was analyzed by electroretinography. Retinal production of the RAS components and phosphorylation of ERK (extracellular-signal regulated kinase) were examined by immunoblotting. Retinal mRNA and protein levels of synaptophysin were measured by quantitative RT-PCR and immunoblot analyses, respectively. In vitro, synaptophysin levels were also evaluated using angiotensin II-stimulated PC12D neuronal cells cultured with or without the inhibition of ERK signaling or the ubiquitin-proteasome system (UPS).

RESULTS—Induction of diabetes led to a significant increase in retinal production of angiotensin II and AT1R together with ERK activation in the downstream of AT1R. AT1R blockade significantly reversed diabetes-induced electroretinography changes and reduction of synaptophysin protein, but not mRNA, levels in the diabetic retina. In agreement with the AT1R-mediated post-transcriptional downregulation of synaptophysin in vivo, in vitro application of angiotensin II to PC12D neuronal cells caused the UPS-mediated degradation of synaptophysin protein via AT1R, which proved to be induced by ERK activation.

CONCLUSIONS—These data indicate the first molecular evidence of the RAS-induced synaptophysin degradation and neuronal dysfunction in the diabetic retina, suggesting the possibility of the AT1R blockade as a novel neuroprotective treatment for diabetic retinopathy. *Diabetes* 57:2191–2198, 2008

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Received 11 September 2007 and accepted 12 May 2008.

Published ahead of print at <http://diabetes.diabetesjournals.org> on 16 May 2008. DOI: 10.2337/db07-1281.

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Diabetic retinopathy is a vision-threatening disease with neurodegenerative change due to chronically progressive microangiopathy. The earliest functional disruption clinically detectable is changes in oscillatory potentials (OPs) measured by electroretinography (ERG) (1,2). The cellular source of OPs is regarded as retinal neurons with synapse formation in the inner retina, including bipolar and amacrine cells (3). At present, there is no established neuroprotective treatment for diabetic retinopathy, since molecular mechanisms underlying diabetes-induced retinal neuronal damage remain unclear.

We have recently demonstrated that angiotensin II type 1 receptor (AT1R) signaling contributes to diabetes-induced retinal inflammation such as leukocyte adhesion to the retinal vasculature (4). Angiotensin II functions as a proinflammatory factor to induce the activation of nuclear factor- κ B pathway in microvascular endothelial cells (4). Angiotensin II is a final product of the renin-angiotensin system (RAS) produced from angiotensinogen through enzymatic cascade reactions, and the RAS components required for the generation of angiotensin II are reported to exist in the eye (5–7). Indeed, human surgical samples from eyes with diabetic retinopathy showed a significant increase in angiotensin II levels (8–10). Increasing evidence has suggested the contribution of the RAS to diabetes-induced retinal vascular complications including leukocyte adhesion (4), hyperpermeability (11), and impaired blood flow (12); however, little is known about the pathogenesis of angiotensin II-mediated neuronal dysfunction in the diabetic retina. Although AT1R blockade led to amelioration of hypertension-induced retinal dysfunction that was exacerbated with diabetes (13), no data have been reported that show the direct effect of AT1R signaling on diabetes-induced retinal dysfunction together with underlying molecular mechanisms.

Recently, we revealed the coexpression of AT1R and the synaptic protein synaptophysin in the inner retinal neurons (14), consistent with several previous reports showing synaptic expression of AT1R in the brain (15–18). Synaptophysin, the major synaptic vesicle protein, is a marker of synapses reported to be reduced in the postmortem brains affected by several neurodegenerative diseases (19). Considering that OPs in ERG are originated from inner retinal neurons bearing AT1R, we hypothesize that angiotensin II directly induces synaptophysin dysregulation and visual functional damage represented by ERG changes. In the present article, we report the first evidence showing that AT1R signaling contributes to diabetes-induced retinal dysfunction and

TABLE 1
Systemic data

	Nondiabetes	Diabetes		
		Vehicle	Telmisartan	Valsartan
<i>n</i>	17	17	18	18
Body weight (g)	26.0 ± 1.3	24.2 ± 1.3†	24.8 ± 1.1*	24.3 ± 1.2†
Blood glucose (mg/dl)	156 ± 22	544 ± 64†	539 ± 81†	516 ± 96†

**P* < 0.05, †*P* < 0.01 vs. nondiabetes.

synaptophysin downregulation together with underlying molecular mechanisms.

RESEARCH DESIGN AND METHODS

Induction of diabetes. C57BL/6 mice (Clea, Tokyo, Japan) at the age of 6 weeks were used in diabetes induction. All animal experiments were conducted in accordance with the ARVO (Association for Research in Vision and Ophthalmology) Statement for the Use of Animals in Ophthalmic and Vision Research. Animals received intraperitoneal injections of streptozotocin (Sigma, St. Louis, MO) at the dose of 60 mg/kg body weight for 3 days. Blood glucose concentrations were measured from the tail vein using Mediasafe mini GR-102 (Terumo, Tokyo, Japan). Development of diabetes was defined by blood glucose >250 mg/dl 7 days after the first injection of streptozotocin.

AT1R blockade in vivo. Mice were intraperitoneally injected with the AT1R blocker (ARB) telmisartan or valsartan (U.S. Pharmacopeia, Rockville, MD) at the dose of 5 or 10 mg/kg body weight, respectively, or vehicle (0.25% DMSO in PBS). The ARB treatment started 22 days after the first injection of streptozotocin for 6 consecutive days and continued until the end of the study (4-week diabetes at evaluation). The doses used in each ARB group were determined according to our previous study on diabetes-induced retinal inflammation (4). Telmisartan was a kind gift of Boehringer Ingelheim (Ingelheim, Germany).

ERG analyses. Animals were dark-adapted for 12 h and prepared under dim red illumination. Mice were anesthetized with pentobarbital sodium at the dose of 70 mg/kg body weight and placed on a heating pad that maintained their body temperature at 35–36°C throughout the experiments. The pupils were dilated with a mixed solution of 0.5% tropicamide and 0.5% phenylephrine (Mydrin-P; Santen, Osaka, Japan). The ground electrode was a subcutaneous needle in the tail, and the reference electrode was placed subcutaneously between the eyes. The active contact lens electrodes (Mayo, Inazawa, Japan) were placed on the cornea. Recordings were performed with PowerLab system 2/25 (AD Instruments, New South Wales, Australia). Responses were differentially amplified at the gain of 1,000 times using an AC (alternate current)-coupled bioamplifier ML132 (AD Instruments) and filtered through a bandpass filter ranging from 0.3 to 500 Hz to yield a- and b-waves. OPs were simultaneously recorded using a high-pass filter set to 100 Hz, so that an overall bandpass ranging from 100 to 500 Hz was achieved. Light pulses of 800 cd·s/m² were delivered via Ganzfeld System SG-2002 (LKC Technologies; Gaithersburg, MD). The amplitude and implicit time of the a- and b-waves and OPs were measured and compared among age-matched nondiabetic controls and 4-week diabetic animals treated with vehicle, telmisartan, or valsartan.

In vitro assays. The neuronal cell line PC12D cells were cultured in Dulbecco's modified Eagle's medium (DMEM; Invitrogen, Carlsbad, CA) supplemented with 10% fetal bovine serum and 10% horse serum (Medical & Biological Laboratories, Nagoya, Japan). Cells were incubated with 50 μg/μl of nerve growth factor (Millipore, Billerica, MA) for differentiation to neurons for 2 days. The neuronal cells were applied with the ARB telmisartan (10 μmol/l) or valsartan (25 μmol/l), the proteasome inhibitor MG132 or lactacystin (20 μmol/l) for both; Calbiochem, San Diego, CA), the lysosome inhibitor E64 (50 μmol/l; Sigma), the extracellular signal-regulated kinase (ERK) kinase inhibitor U0128 (10 μmol/l; Cell Signaling Technology, Beverly, MA) or PD98059 (10 μmol/l; Calbiochem), the phosphatidylinositol 3-kinase inhibitor LY294002 (20 μmol/l; Calbiochem) or wortmannin (100 nmol/l; Sigma), the Janus kinase inhibitor AG490 (1, 10, or 100 μmol/l; Calbiochem), or vehicle. After 5 min, the pretreated cells were subsequently stimulated with 100 nmol/l angiotensin II for 30 min. Immediately after the exposure to angiotensin II, the neuronal cells were subjected to quantitative RT-PCR and immunoprecipitation for synaptophysin and to immunoblot analyses for synaptophysin and phosphorylated forms of ERK1/2, Akt (protein kinase B), and signal transducer and activator of transcription (STAT)-3.

AT1R and ERK1/2 knockdown by RNA interference. Transfections were performed using Lipofectamine 2000 (Invitrogen) and the SureSilencing short

hairpin RNA (shRNA) kit (SuperArray Bioscience, Frederick, MD) according to the manufacturers' instructions. Briefly, 4 μg Rat Agtr1a (AT1R), Rat Mapk3 (ERK1), and Rat Mapk1 (ERK2) or control shRNA was incubated with 10 μl of the transfection reagent in 500 μl serum-free medium for 20 min to facilitate complex formation. The resulting mixture was added to PC12D cells cultured in a six-well plate with 2 ml medium for 48 h before differentiation.

Quantitative RT-PCR analyses for synaptophysin. Animals were killed with an overdose of anesthesia. The eyes were immediately enucleated and the retina was carefully isolated. Total RNA was extracted from the retina or PC12D cells using an extraction reagent (TRIzol; Invitrogen), and cDNA was synthesized with SuperScript III reverse transcriptase (Invitrogen). Real-time PCR was performed using MX3000 (Stratagene, La Jolla, CA) with PCR primers for synaptophysin designed by TaqMan Gene Expression Assays (Applied Biosystems, Foster, CA). The mRNA levels were normalized to β-actin as an internal control.

Immunoblot analyses for angiotensin II, AT1R, synaptophysin, and phosphorylated forms of ERK, Akt, and STAT3. The isolated retinas or PC12D cells were placed into lysis buffer (10 mmol/l Tris-HCl [pH 7.6], 100 mmol/l NaCl, 1 mmol/l EDTA, 1% Triton X-100, protease inhibitors). Each sample was separated with SDS-PAGE and electroblotted to polyvinylidene fluoride (PVDF) membranes (Millipore, Bedford, MA). After nonspecific binding was blocked with 4% skim milk, the membranes were incubated at 4°C overnight with a rabbit polyclonal antibody against angiotensin II, AT1R (1:200, 1:100, respectively; Santa Cruz Biotechnology, Santa Cruz, CA), or phosphorylated forms of ERK, Akt, or STAT3 (1:1,000, 1:500, or 1:1,000, respectively; Cell Signaling Technology) or a mouse monoclonal antibody against synaptophysin (1:500; Sigma) or α-tubulin (1:2,000; Sigma). The membranes were then incubated with a horseradish peroxidase-conjugated goat antibody against rabbit or mouse immunoglobulin or with a biotinylated secondary antibody followed by avidin-biotin horseradish peroxidase complexes (Vectastain Elite ABC Kit; Vector, Burlingame, CA). The signals were visualized with chemiluminescence (ECL Blotting Analysis System; Amersham, Arlington Heights, IL), measured by ImageJ software (National Institutes of Health, Bethesda, MD) and normalized to α-tubulin.

Ubiquitination assays. PC12D cells were transfected with a hemagglutinin-ubiquitin plasmid (provided by Keiji Tanaka, the Tokyo Metropolitan Institute of Medical Science, Tokyo, Japan) as described above at the time of differentiation. Forty-five hours after transfection, the cells were incubated for 3 h with 20 μmol/l MG132. The transfected cells were then pretreated with telmisartan, valsartan, or vehicle and followed by stimulation with angiotensin II as described above. The stimulated cells were placed into the lysis buffer, and the cell lysate was applied with a rabbit polyclonal antibody against synaptophysin (1:25; Sigma). After rocking at 4°C for 1 h, the lysate treated with anti-synaptophysin antibodies were then added with protein G-sepharose beads (Sigma) and incubated overnight at 4°C. After washes of nonspecific binding to beads, the bound protein was solubilized in sample buffer, loaded on an SDS-PAGE, and processed for immunoblotting with a rat monoclonal antibody against hemagglutinin (1:500; Roche Applied Science, Mannheim, Germany) to detect multi-ubiquitinated synaptophysin and with a mouse monoclonal antibody against synaptophysin (1:500; Sigma) to confirm the equal amount of synaptophysin protein applied in each lane.

Statistical analysis. All results were expressed as means ± SD. The values were processed for statistical analyses (one-way ANOVA with Tukey's post hoc test), and differences were considered statistically significant at *P* < 0.05.

RESULTS

Diabetes-induced retinal production of angiotensin II and AT1R led to ERK activation. Mice with streptozotocin-induced diabetes showed a significant (*P* < 0.05) decrease in body weight and a significant (*P* < 0.01) increase in blood glucose, compared with age-matched nondiabetic controls (Table 1). Treatment with telmisar-

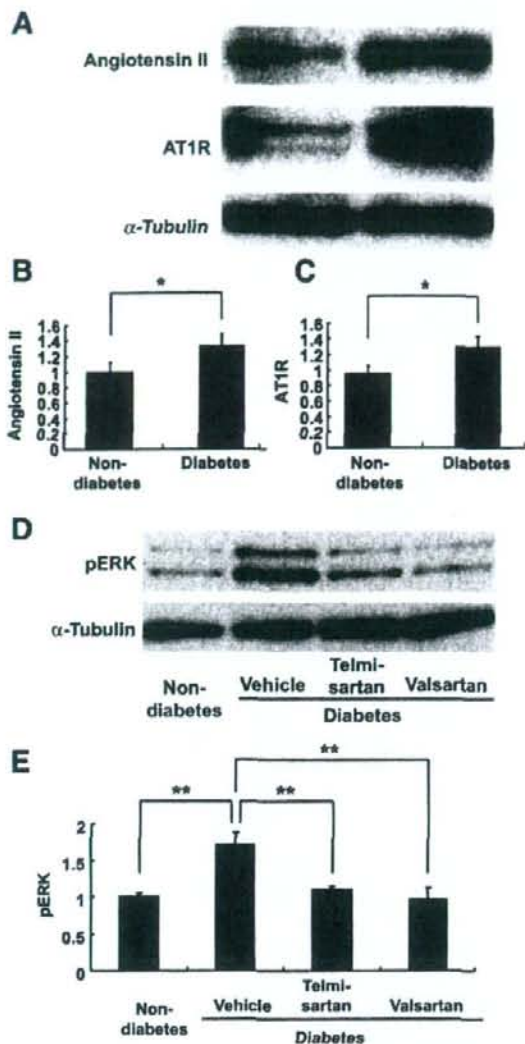


FIG. 1. Diabetes-induced retinal production of angiotensin II and AT1R led to ERK activation. Immunoblot analyses showing the production of the RAS components (A-C) and ERK activation in the diabetic retina (D and E). Retinal levels of angiotensin II and AT1R were significantly higher in 4-week diabetic than in nondiabetic mice (A-C). ERK phosphorylation, elevated in 4-week diabetic mice, was significantly attenuated by AT1R blockade with telmisartan or valsartan (D and E). $n = 4$ for all. * $P < 0.05$.

tan or valsartan of 4-week diabetic mice did not significantly ($P > 0.05$) change these metabolic parameters (Table 1). The retinas from 4-week diabetic mice were subjected to immunoblot analyses to detect the production of the RAS components and its downstream ERK activation (Fig. 1). Retinal levels of angiotensin II and its receptor AT1R were significantly ($P < 0.01$) higher in 4-week diabetic than in nondiabetic animals (Fig. 1A-C). Consistently, ERK phosphorylation, enhanced in 4-week diabetic mice ($P < 0.01$), was significantly ($P < 0.01$) attenuated by AT1R blockade with telmisartan or valsartan to the level equivalent with that in nondiabetic mice

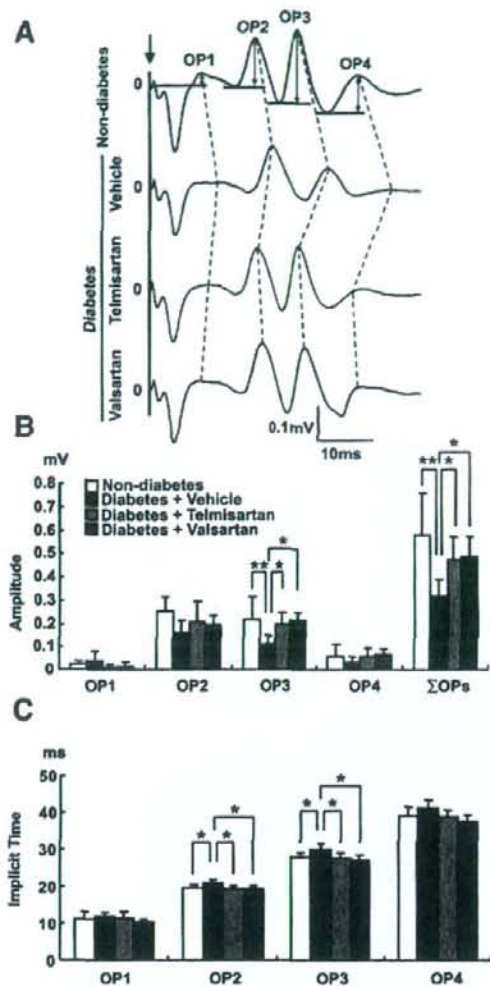


FIG. 2. Diabetes-induced visual dysfunction was suppressed by AT1R blockade. A: Representative wave responses from an individual mouse in each group to one flash. A and B: ERG analyses showing that AT1R blockade with telmisartan or valsartan significantly reversed the reduction of OP3 and total OPs (Σ OPs) amplitude in 4-week diabetic mice. Similarly, the implicit time of OP2-3, prolonged in 4-week diabetic mice, was significantly recovered by AT1R blockade (A and C). $n = 7-9$. * $P < 0.05$, ** $P < 0.01$.

(Fig. 1D and E). In contrast, AT1R blockade did not attenuate the baseline level of retinal pERK in nondiabetic mice (data not shown).

Diabetes-induced visual dysfunction was suppressed by AT1R blockade. To determine the effect of the RAS activation in the diabetic retina (Fig. 1) on visual function, we performed ERG analyses (Fig. 2). OP changes including the reduced amplitude and prolonged implicit time are known to occur in the early stage of murine and human diabetic retinopathy (1,2,20,21). There was no remarkable difference in a- or b-wave among nondiabetes and 4-week diabetes treated with vehicle, telmisartan, and valsartan (data not shown). Importantly, AT1R blockade with telmisartan or valsartan significantly reversed the reduction of

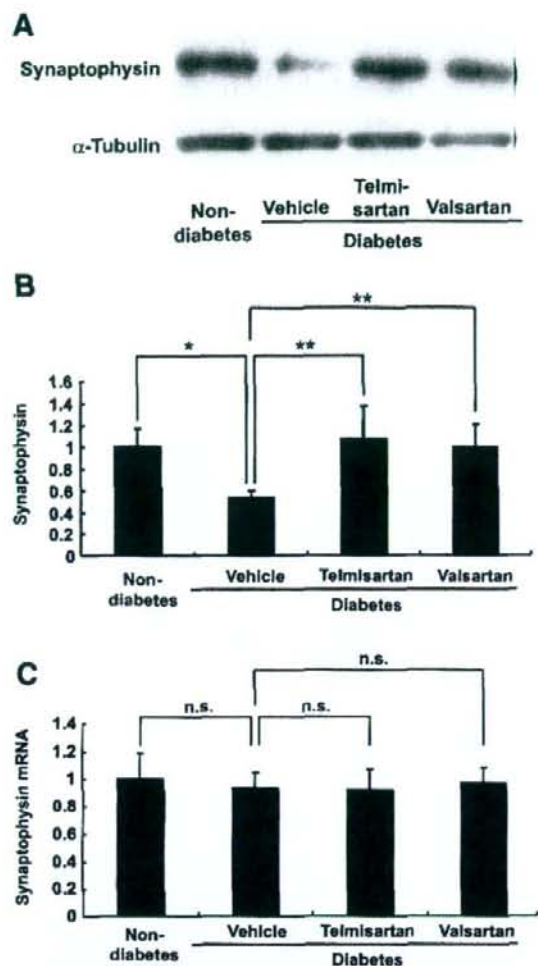


FIG. 3. Posttranscriptional reduction of synaptophysin in the diabetic retina was reversed by AT1R blockade. The retinal levels of synaptophysin protein, reduced by inducing diabetes, were significantly reversed in 4-week diabetic mice by AT1R blockade with telmisartan or valsartan (*A* and *B*). In contrast, mRNA levels of synaptophysin were unaltered in 4-week diabetic mice treated with vehicle, telmisartan, or valsartan compared with age-matched nondiabetic control animals (*C*). $n = 4-8$. * $P < 0.05$, ** $P < 0.01$.

OP3 and total OPs (Σ OPs) amplitude ($P < 0.05$ for both) in 4-week diabetic mice (Fig. 2*A* and *B*). Similarly, the implicit time of OP2 and OP3, prolonged in 4-week diabetic mice, was significantly ($P < 0.05$ for both) recovered by AT1R blockade to the normal level (Fig. 2*A* and *C*). In contrast, AT1R blockade did not alter normal ERG waves in nondiabetic mice (data not shown).

Posttranscriptional reduction of synaptophysin in the diabetic retina was reversed by AT1R blockade. To investigate the molecular mechanisms underlying the AT1R-mediated impairment of ERG responses (Fig. 2), we analyzed the regulation of the synaptic vesicle protein synaptophysin (Fig. 3), which is abundantly expressed in the inner retinal neurons (14), the cellular source of OPs (1,3,22,23). Synaptophysin is indispensable for several

presynaptic functions, including the release of neurotransmitters (24). The retinal levels of synaptophysin protein, reduced by inducing diabetes, were significantly ($P < 0.01$) reversed in 4-week diabetic mice by AT1R blockade with telmisartan or valsartan (Fig. 3*A* and *B*). In contrast, mRNA levels of synaptophysin were unaltered in 4-week diabetic mice treated with vehicle, telmisartan, or valsartan, compared with those in age-matched nondiabetic controls (Fig. 3*C*). The discrepancy between the protein and mRNA levels indicated that the AT1R-mediated decline of synaptophysin in the diabetic retina was posttranscriptionally regulated. In contrast, AT1R blockade did not affect the physiologic level of synaptophysin mRNA or protein in the nondiabetic retina (data not shown).

AT1R-mediated posttranscriptional reduction of synaptophysin was regulated by the ubiquitin-proteasome system in neuronal cells. To elucidate the molecular mechanisms involving the AT1R-mediated posttranscriptional decrease in synaptophysin in the diabetic retina (Fig. 3), we used the in vitro culture system with PC12D neuronal cells stimulated with angiotensin II (Fig. 4). In agreement with the in vivo data (Fig. 3), in vitro application with angiotensin II significantly ($P < 0.01$) reduced synaptophysin protein (Fig. 4*A* and *B*), but not mRNA (Fig. 4*C*), levels in a posttranscriptional manner, which were significantly ($P < 0.05$ for all) reversed by AT1R blockade with telmisartan, valsartan, or shRNA for AT1R knockdown (Fig. 4*A*, *B*, and *F*). Because synaptic vesicle proteins including synaptophysin have recently proven to be physiologically degraded by the UPS for the maintenance of synaptic plasticity (25-27), we examined the involvement of the ubiquitin-proteasome system (UPS) (Fig. 4*G-I*) with AT1R-mediated posttranscriptional decrease in synaptophysin protein (Fig. 4*A-F*). Multi-ubiquitinated synaptophysin as a high-molecular weight smear, enhanced by stimulation with angiotensin II, was substantially attenuated by AT1R blockade with telmisartan or valsartan (Fig. 4*G*). Application with the proteasome inhibitor MG132 or lactacystin, but not the lysosome inhibitor E64, led to significant ($P < 0.05$) suppression of angiotensin II-induced degradation of synaptophysin (Fig. 4*H* and *I*). In contrast, the baseline levels of synaptophysin in PC12D cells not stimulated with angiotensin II were unaffected with these inhibitors (data not shown).

AT1R-mediated ERK activation was required for synaptophysin degradation in neuronal cells. Because ERK was activated in the downstream of AT1R signaling in the diabetic retina (Fig. 1), we examined the involvement of ERK activation (Fig. 5) with AT1R-mediated degradation of synaptophysin in the neuronal cells (Fig. 4). Consistent with AT1R-mediated ERK activation in the diabetic retina (Fig. 1), angiotensin II administration induced the phosphorylation of ERK in the neuronal cells, which was significantly ($P < 0.05$) inhibited by AT1R blockade with telmisartan or valsartan (Fig. 5*A* and *B*). Inhibition of ERK activation with U0126, PD98059, or shRNA for ERK1/2 knockdown led to significant ($P < 0.05$ for all) suppression of angiotensin II-induced degradation of synaptophysin (Fig. 5*C-H*). In addition to ERK signaling, phosphatidylinositol 3-kinase/Akt and Janus kinase/STAT pathways are known to be in the downstream of AT1R. In the neuronal cells, angiotensin II treatment also increased the phosphorylated forms of Akt and STAT3 (data not shown). In contrast to ERK involvement (Fig. 5*C-H*), however, the phosphatidylinositol 3-kinase inhibitor wortmannin or LY294002 or the Janus kinase inhibitor AG490 did not

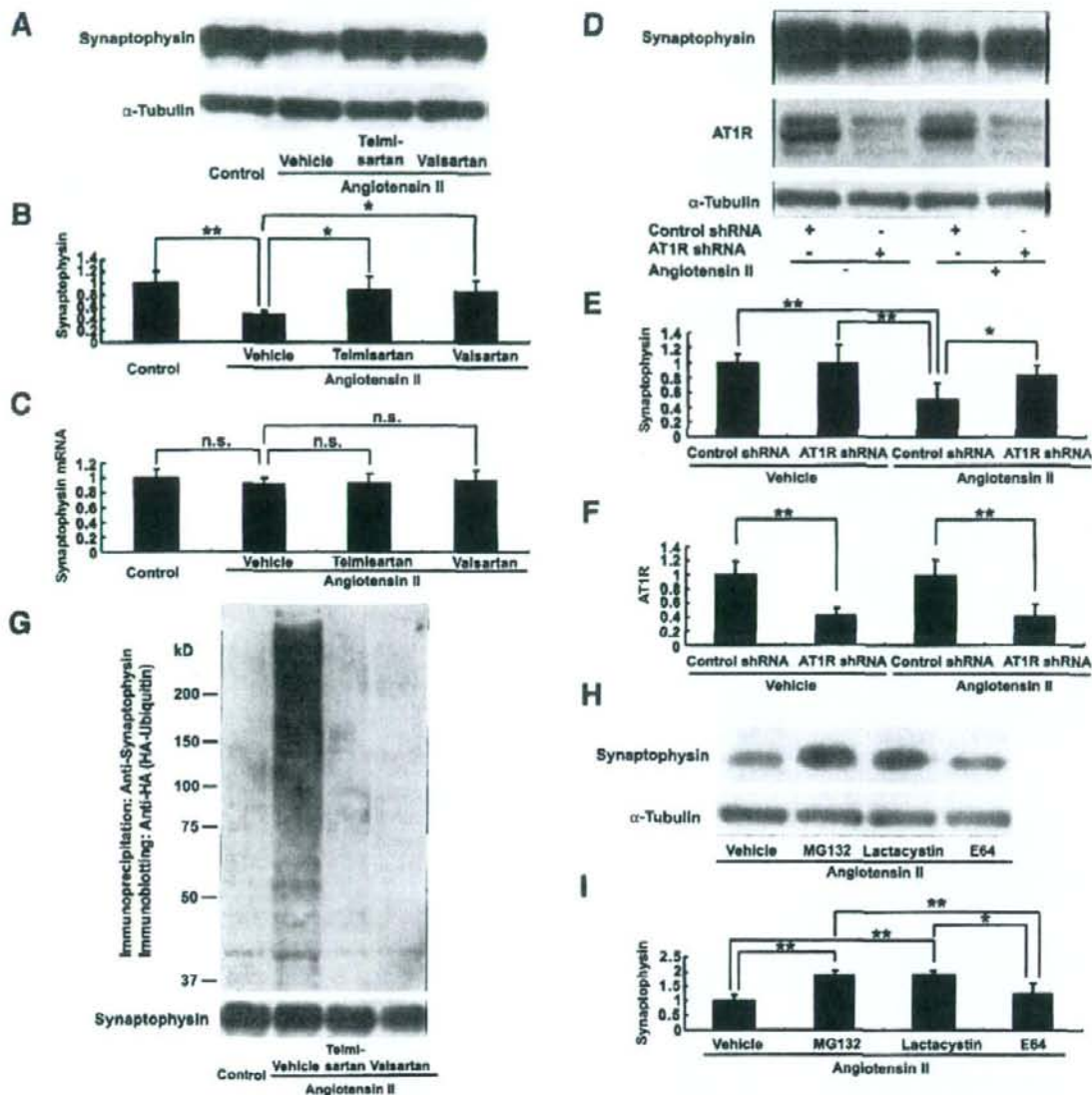


FIG. 4. AT1R-mediated posttranscriptional reduction of synaptophysin was regulated by the UPS in neuronal cells. In vitro application with angiotensin II significantly reduced synaptophysin protein (A and B), but not mRNA (C), levels in a posttranscriptional manner, which were significantly reversed by AT1R blockade with telmisartan or valsartan. The data with the pharmacologic AT1R blockade (A–C) were reproduced via shRNA for AT1R knockdown (D–F). Multi-ubiquitinated synaptophysin as a high-molecular weight smear, enhanced by stimulation with angiotensin II, was substantially attenuated by AT1R blockade (G). Application with the proteasome inhibitor MG132 or lactacytin, but not the lysosome inhibitor E64, led to significant suppression of angiotensin II-induced degradation of synaptophysin (H and I). $n = 4-6$. * $P < 0.05$, ** $P < 0.01$.

reverse angiotensin II-induced degradation of synaptophysin in vitro (data not shown).

DISCUSSION

The present study reveals, for the first time to our knowledge, several important findings concerning the relationship of the RAS with diabetes-related neuronal damage in vivo and in vitro. In the diabetic retina, the RAS components angiotensin II and AT1R were upregulated together with AT1R's downstream ERK activation (Fig. 1). AT1R

blockade with telmisartan or valsartan significantly reversed diabetes-induced OP changes measured by ERG (Fig. 2), known functional abnormalities in human diabetic retinopathy. Moreover, AT1R signaling caused a posttranscriptional decrease in the synaptic vesicle protein synaptophysin in the diabetic retina, which was rescued by the ARB application to diabetic mice (Fig. 3). The protein, but not mRNA, reduction of synaptophysin in neuronal cells was shown to depend on the UPS, which was enhanced via AT1R signaling (Fig. 4). Angiotensin II-induced neuronal ERK

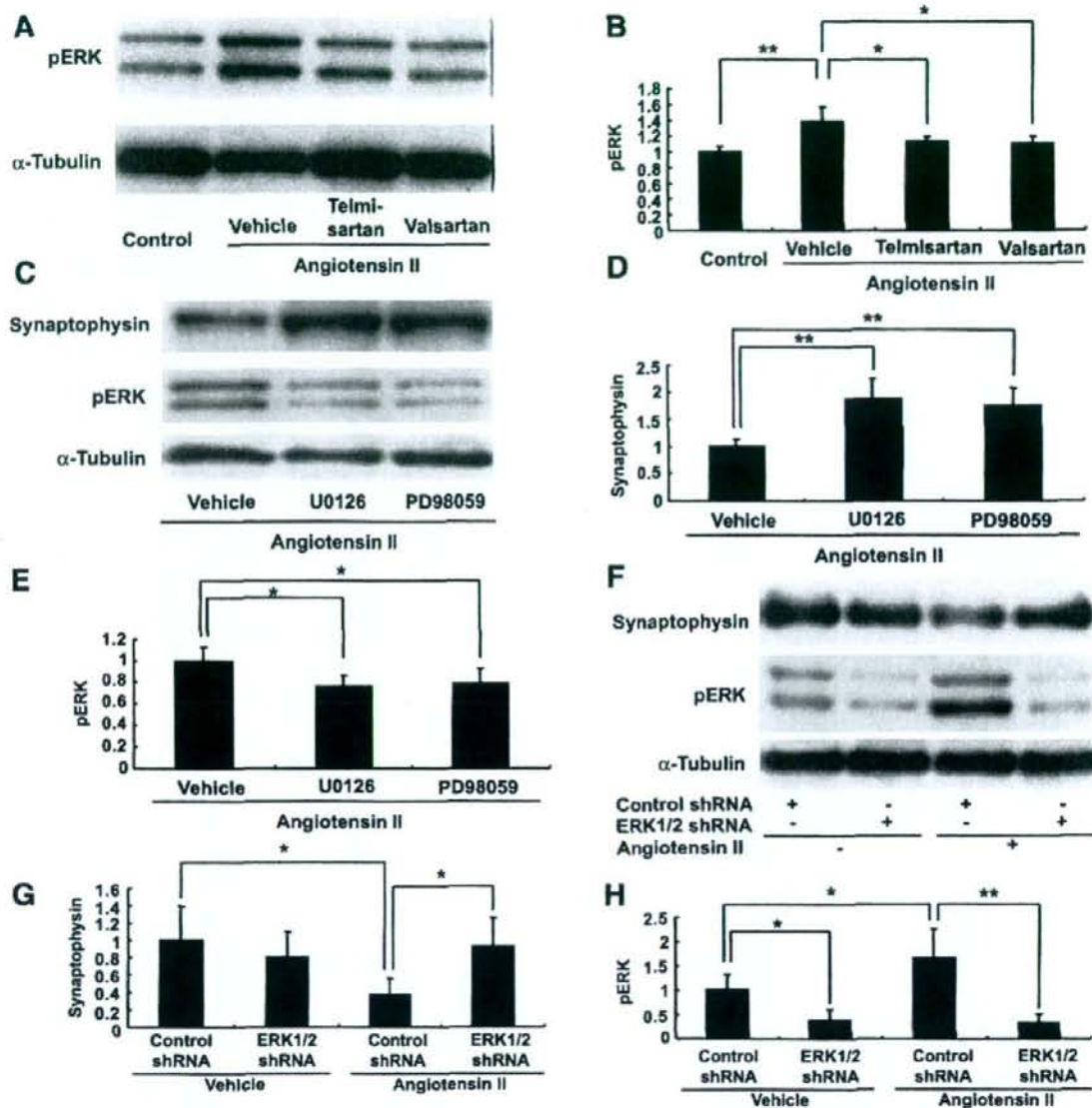


FIG. 5. AT1R-mediated ERK activation was required for synaptophysin degradation in neuronal cells. Angiotensin II administration induced the phosphorylation of ERK in the neuronal cells, which was significantly inhibited by AT1R blockade with telmisartan or valsartan (A and B). Inhibition of ERK activation with U0126 or PD98059 led to significant suppression of angiotensin II-induced decline of synaptophysin (C-E). The data with the pharmacologic ERK inhibition (C-E) were reproduced via shRNA for ERK1/2 knockdown (F-H). $n = 4-6$. * $P < 0.05$, ** $P < 0.01$.

activation, also observed in the diabetic retina (Fig. 1), resulted in the degradation of synaptophysin protein *in vitro* (Fig. 5). The use of two different ARBs throughout the experiments confirmed that the suppression of diabetes-induced retinal dysfunction and synaptophysin degradation is a class effect for ARBs. Additionally, our *in vitro* results were further confirmed via gene-specific shRNA knockdown technique for AT1R and ERK.

In the diabetic retina, the production of angiotensin II and AT1R was upregulated, leading to ERK activation in the downstream of AT1R signaling (Fig. 1). Since ERK is known to be one of AT1R's signaling pathways, the

present data are compatible with clinical (5,7,28,29) and experimental (4) studies showing increased angiotensin II generation in the diabetic eye. The cellular source of angiotensin II was shown to be retinal glial cells by immunohistochemistry for human postmortem eyes (30). In the liver and kidney as well (31-33), diabetes-induced activation of the RAS causes elevated concentration of tissue angiotensin II. The tissue RAS activation in diabetes is explained in part by the findings that *in vitro* stimulation with high glucose enhanced the transcriptional activity of angiotensinogen gene encoding a precursor protein for angiotensin I and II (34,35).

Our ERG analyses demonstrated the reduced amplitude and prolonged implicit time of OPs in diabetic mice, in accordance with clinical and experimental data on attenuated OP changes in early diabetes (1,2,29). Importantly, AT1R blockade significantly improved these functional parameters in the diabetic retina to the normal levels observed in nondiabetic controls (Fig. 2). Reasonably, the present ERG findings, in concert with our recent data on AT1R expression in the inner retinal neurons (14), the cellular origins of OPs, suggest that the functional recovery of the diabetic inner retina was attributable in part to the direct effect of AT1R blockade on the inner retinal neurons.

Several previous studies on neuronal damage in the diabetic retina showed the cellular changes including the apoptosis of ganglion and amacrine cells and the activation of glial cells (36–38). In contrast, the present study revealed, with the molecular changes in the diabetic retina, a significant decrease in synaptophysin (Fig. 3), the synaptic vesicle protein capable of maintaining the function of retinal neurons. Furthermore, AT1R signaling was shown to play a crucial role in the reduction of synaptophysin protein (Fig. 3). Pharmacologic studies using tetrodotoxin and glycine suggested the cellular origins of OPs as neurons with synapse formation in the inner retina including bipolar and amacrine cells (3,22,23), both of which develop a retinal neuronal network contributing to visual function. Synaptophysin, a well-known marker for the presynapse, was shown to decrease with the progression of Parkinson's disease and Alzheimer's disease (19). In the retina as well, pathogenic conditions such as retinal detachment and inflammation caused reduction of synaptophysin (14,39). Mice deficient in synaptophysin exhibited a significant decrease in the number of synaptic vesicles (24). The neuron maintains its viability and activity by interacting with other neurons via synapses; however, synaptophysin knockouts were for the most part functionally compensated possibly because of biological redundancy (24). Although it is reasonable to think about the possibility of cooperative involvement of several other synaptic proteins, the diabetes-induced reduction of synaptophysin (Fig. 3) was attributable, at least in part, to the functional disturbance of neuronal activity in the diabetic retina. The currently observed suppression of synaptophysin decrease by AT1R blockade (Fig. 3) is likely to rescue the activity of the AT1R-bearing inner retinal neurons, leading to the improvement of OP changes (Fig. 2).

The present *in vitro* data (Figs. 4 and 5) further elucidated the detailed molecular mechanisms underlying the AT1R-mediated posttranscriptional decrease in synaptophysin observed *in vivo* (Fig. 3). Angiotensin II directly exerted its bioactivity to the neuronal cells via AT1R, leading to a posttranscriptional decrease in synaptophysin (Fig. 4A–F), which mimicked the *in vivo* data (Fig. 3). Interestingly, angiotensin II/AT1R signaling proved to cause the activation of the UPS, leading to synaptophysin degradation in neuronal cells (Fig. 4G–I). Although the UPS was recently shown to play a role in the degradation of several synaptic proteins (25–27), the present data are the first to reveal the RAS-mediated UPS activation for synaptic protein degradation in neurons. Out of several signaling molecules in the downstream of AT1R, the phosphorylation of ERK, but not Akt or STAT3, was required for synaptophysin degradation (Fig. 5), suggesting that the UPS-dependent synaptophysin degradation is mediated by ERK activation. This is supported by several

recent studies showing that the activation of the ERK pathway promoted the UPS-mediated degradation of the cell cycle protein Cdc25 (40,41) and the cell survival factor Bim (42). In the *Drosophila* eye, ERK signaling was shown to be required for the physiologic activity of Sina (seven in absentia), a key molecule for normal photoreceptor development (43). Interestingly, the mammalian homologues of *Drosophila* Sina, Shiah (seven in absentia homologues), proved to be synaptophysin-binding proteins functioning as E3 ubiquitin ligases to regulate the UPS-mediated degradation of synaptophysin (26). Reasonably, baseline ERK phosphorylation in the normal retina (Fig. 1) is likely to play a physiologic role in the UPS-mediated degradation of synaptophysin for the maintenance of synaptic plasticity (25–27). In the diabetic retina, however, excessive ERK activation mediated by AT1R (Fig. 1) is thought to induce the pathologic decline of synaptophysin (Fig. 3), contributing to neuronal dysfunction demonstrated by ERG (Fig. 2).

In the current study, diabetes-induced retinal dysfunction and synaptophysin downregulation were successfully reversed by AT1R signaling blockade *in vivo* (Figs. 2 and 3). Although future studies are required to determine whether angiotensin II/AT1R signaling leads to diabetes-induced neuronal damage systemically or locally in specific target organs, AT1R blockade has proven to be neuroprotective at least in the diabetic eye. Collectively, our present data show the possibility of AT1R blockade as a novel therapeutic strategy for neuronal dysfunction in vision-threatening diabetic retinopathy.

ACKNOWLEDGMENTS

The authors thank Masao Yoshikawa, Eiichiro Nagasaka (Mayo Company, Inazawa, Japan), and Hirobumi Tada (Yokohama City University, Yokohama, Japan) for expert advice and Haruna Koizumi and Taiga Shioda (Keio University, Tokyo, Japan) for technical assistance. This study was supported by the Japanese Ministry of Education, Culture, Sports, Science and Technology (Grant-in-Aid for Scientific Research [number 19791289] to T.K. and the 21st Century COE Program at Keio University).

This work was presented in part at the 2007 annual meeting of ARVO (Association for Research in Vision and Ophthalmology) in Fort Lauderdale, FL.

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Letters to the Editor

Optical coherence tomographic findings of vitreomacular traction syndrome with macular hole retinal detachment

We report on two patients with the vitreomacular traction (VMT) syndrome whose vitreoretinal interface had a unique morphology. Case 1 was a 70-year-old woman who was referred to treat a macular hole retinal detachment (MHRD) in her right eye. Her visual acuity

was 0.2, and ophthalmoscopy detected a MHRD of approximately three disc diameters (Fig. 1). Her refractive error (spherical equivalent) was +4.25 dioptres and the axial length was 21.02 mm. Optical coherence tomography (OCT) showed a perifoveal vitreous detachment with focal vitreal adhesions to the fovea, an incomplete posterior vitreous detachment (PVD), and two low reflective lines in the vitreous cavity (Fig. 1).

Vitrectomy was performed with phacoemulsification and implantation of an intraocular lens. After a complete PVD was created, the internal limiting membrane (ILM) in the area of the detached retina was peeled off (Fig. 1). The vitreous cavity was

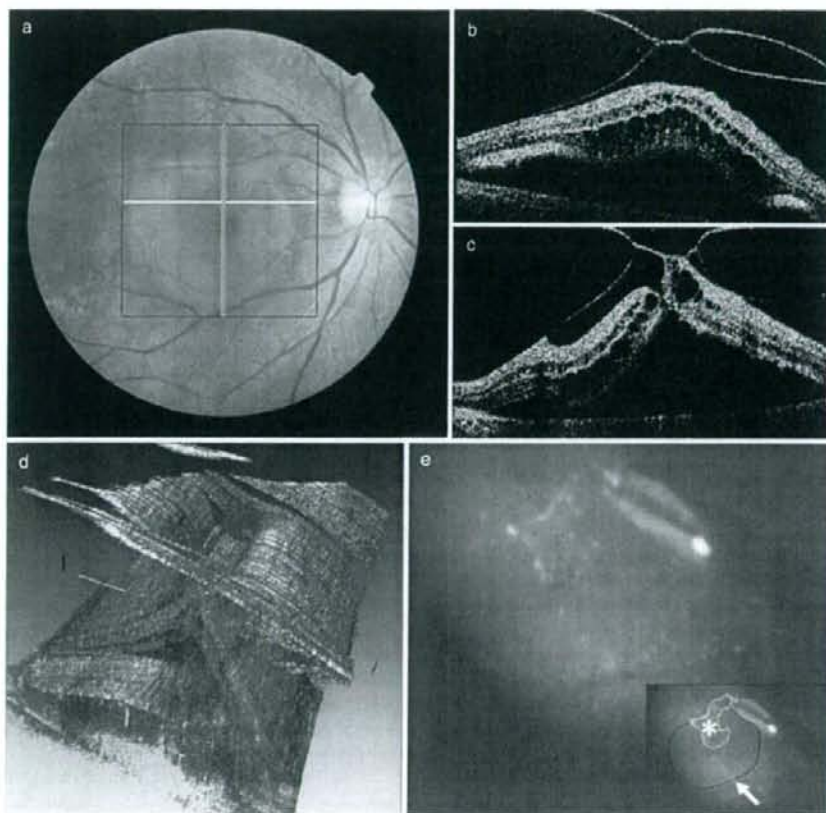


Figure 1 Fundus photograph and optical coherence tomographic (OCT) images showing macular hole retinal detachment in Case 1 (a) Fundus photograph showing a macular hole of approximately three disc diameters and a retinal detachment. The horizontal (yellow) and vertical (green) lines indicate the OCT scanned lines whose images are shown in (b) and (c), respectively. The square represents the scanned area shown in (d) (b) OCT image showing perifoveal vitreous detachment with the detached posterior vitreous cortex comprised of two low reflective lines (double-layered membrane) in the vitreous cavity. (c) OCT image showing perifoveal vitreous detachment with a macular hole and a focal vitreous adhesion to the fovea. (d) A three-dimensional OCT image showing the focal adhesion of the posterior vitreous surface causing it to form an inverted tent-like configuration over the macula and tent-like configuration of the macula. (e) Intraoperative photograph showing that the internal limiting membrane (ILM) was peeled off after removal of double-layered posterior vitreous membrane. Inset: The asterisk (*) indicates peeled ILM with triamcinolone crystals, and an arrow indicates the area where the ILM had been peeled off.

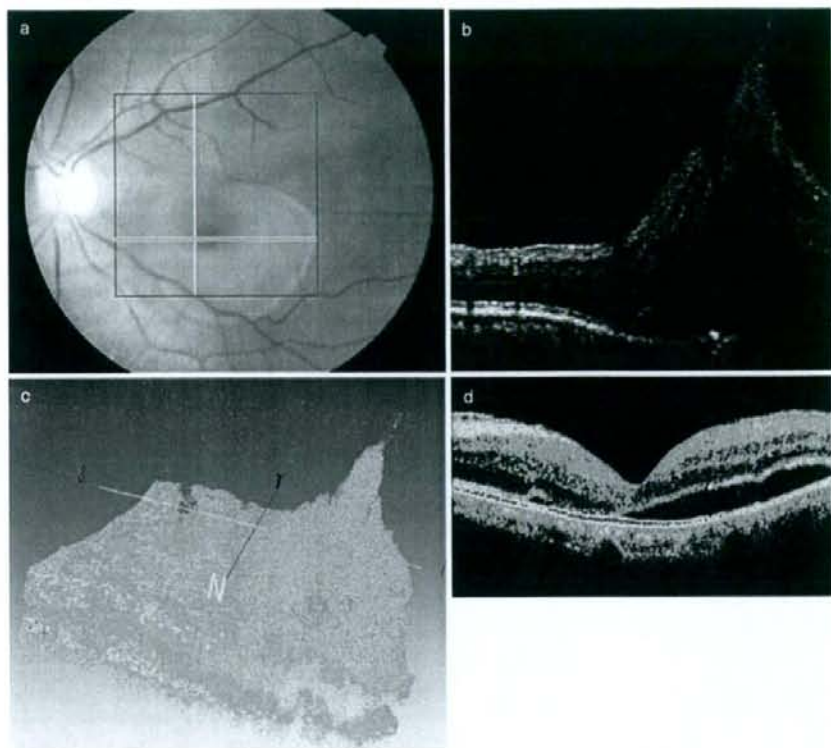


Figure 2. Fundus photograph and optical coherence tomographic (OCT) images showing macular hole retinal detachment (MHIRD) in Case 2. (a) Fundus photograph showing a MHIRD of approximately 2×3 disc diameters. The detached macular area is deviated inferiorly. The vertical (yellow) and horizontal (green) lines indicate the OCT scanned lines which are shown in (b) and (d) (after surgery), respectively. The square represents the OCT scanned area shown in (c). (b) OCT image showing focal attachment of the posterior vitreous to the perifovea that appears to exert a strong anterior-posterior tractional force. (c) A three-dimensional OCT image showing a focal adhesion of the posterior vitreous surface to a region inferior to the fovea causing a strong anterior-posterior tractional force and asymmetrical retinal detachment with its top deviated inferiorly. N = nasal, T = temporal. (d) One month postoperative OCT shows improved macular configuration with residual subretinal fluid.

filled with room air to tamponade the retina. Postoperatively, the retina was reattached, and the MH was closed. The visual acuity improved to 0.9 12 months after the vitrectomy (Fig. 1).

Case 2 was an 84-year-old man who was referred to treat a MHIRD in his left eye. His visual acuity was 0.1, and ophthalmoscopy showed an inferiorly-deviated MHIRD of approximately 2×3 disc diameters (Fig. 2). The left eye was pseudophakic and the axial length measured before MHIRD was 23.17 mm. OCT showed a focal attachment of the posterior vitreous to the perifovea that appeared to cause a strong anterior-posterior tractional force (Fig. 2). Vitrectomy was successfully performed with the same procedure as in Case 1. The retinal and foveal configuration improved although some subretinal fluid remained. The visual acuity improved to 0.7 1 month after surgery (Fig. 2).

The VMT syndrome is characterized by an incomplete PVD with focal vitreomacular adhesions that distort the appearance and depress the function of the macula.⁴ A persistent traction can lead to progressive macular oedema, thickening of the retina, and/or a RD.¹⁵ In addition, a MHIRD can occasionally develop even in non-myopic eyes as in our cases.

Morphological assessments of the vitreoretinal interface by OCT have led to a better understanding of how the anatomical change such as posterior staphyloma and vitreous traction can lead to MHIRD or tractional retinoschisis in highly myopic eyes. Even in non-myopic eyes that do not have such an anatomical pathology, VMT may cause a MHIRD.⁶ But the rarity of these cases has prevented a comprehensive investigations on the morphology of the vitreomacular interface.

Our three-dimensional OCT study clearly illustrated the extensive morphological changes at the vitreomacular interface which played a role in development of MHIRD in these two non-myopic eyes. A double-layered membrane attached to the fovea was considered to cause an unusual anterior-posterior traction thereby leading to the MHIRD in Case 1. Because histology was not available, the exact composition of the membrane cannot be made but the fact that the ILM was peeled off after removal of the membrane suggests that it was not part of the ILM. It may represent an epiretinal membrane (ERM) that was still attached to the detached posterior hyaloid membrane (PHM). Because the ERM was also firmly attached to the retina, traction by the detached PHM on the

ERM may have resulted in the RD. Further histological investigations will help in determining whether this membrane contained contractile elements or whether it was a vitreous schisis.

In Case 2, the off-centred vitreous adhesion probably caused an unusual anterior-posterior tractional force to develop the inferiorly deviated MHRD. Three-dimensional OCT in combination with intraoperative findings may be valuable ways to monitor the progress of the VMT syndrome and to clarify the mechanism of the secondary macular changes.

ACKNOWLEDGEMENTS

Support of this study was provided by Research Grants on Sensory and Communicative Disorders from the Ministry of Health, Labor and Welfare, Japan. No author has a proprietary interest in any material or method mentioned.

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Received 16 December 2007; accepted 24 April 2008

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Optical coherence tomography in relentless placoid chorioretinitis

Relentless placoid chorioretinitis (RPC) was recently described¹ as a unique entity with atypical clinical findings and clinical course to both serpinginous choroidopathy and acute posterior multifocal placoid pigmented epitheliopathy (APMPPE). We report here on the changes in optical coherence tomographic (OCT) features in a patient suffering from RPC.

A 17-year-old healthy boy presented to the Ophthalmology Department at Hadassah University Hospital with an abrupt loss of left eye vision of one-day duration.

On examination visual acuity was 6/6 right eye and finger counting 2 m left eye. He had normal intraocular pressure and normal anterior segment examination. On funduscopy, faint subretinal lesions were noted in the macula and in the mid-peripheral retina of the right eye (Fig. 1a). In the left eye, an elevated yellowish macular lesion was noted (Fig. 1b). No lesions were seen elsewhere. Fluorescein angiography of the right eye showed areas of late hyperfluorescence in the macular area and in the mid-peripheral retina (Fig. 2a), while in the left eye, blocked fluorescence was noted in the macular area in the early phase with late hyperfluorescence (Fig. 2b). OCT was normal in the right eye, but in the left eye subretinal fluid with small area of pigment epithelial detachment was detected in the macular area (Fig. 3a). Also, hyper-reflectivity of the nuclear retinal layers was noted, namely, the ganglion cell layer, the inner and the outer nuclear layers. Full-field FERG, EOG and colour vision were normal. Systemic medical work-up for infectious and inflammatory aetiology, including complete blood count, erythrocyte sedimentation rate, C-reactive protein, kidney function tests, liver function tests and serology for herpes viruses, syphilis, HIV and Lyme disease, was normal. Antinuclear antibodies were not detected, and angiotensin-converting enzyme level and chest X-ray were normal.

Ten days later, the patient presented with spontaneous improvement in left eye vision to 6/10. On examination, anterior segments were normal, +1 vitreous cells were noted with bilateral placoid pigmented chorioretinal scars and active fluffy diffuse subretinal lesions in the mid and far periphery. There was resolution of the left macular yellow lesion with fine pigmentary changes instead. The lesions looked similar to those seen with APMPPE and there was posterior pole involvement. Prednisolone (1 mg/kg/day) was started with a tapering regimen. During the ensuing months and despite the steroid therapy, active lesions were continuously appearing with

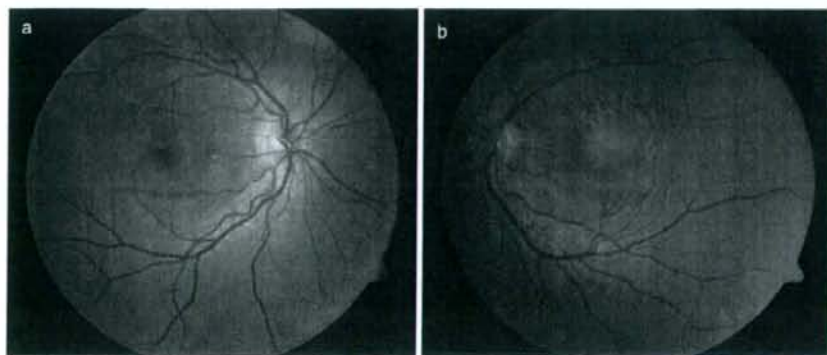


Figure 1 (a) Colour fundus photography of the right eye showing faint subretinal lesions in the macular area temporal to the fovea and in the mid-peripheral retina. (b) Colour fundus photography of the left eye showing an elevated yellowish macular lesion.

Diagnosis/Therapy in Ophthalmology

Frosted branch angiitis associated with streptococcal infection: optical coherence tomography as a follow-up tool

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Acta Ophthalmol.

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Journal compilation © 2008 Acta Ophthalmol

doi: 10.1111/j.1755-3768.2008.01335.x

A 7-year-old boy presented with decreased vision and conjunctivitis in both eyes. He had a flu-like episode with fever about 2 weeks earlier, and the skin of both palms had peeled off. His visual acuity (VA) was 0.15 OD and 0.1 OS. Slit-lamp examination showed conjunctival and ciliary injection, cells in the anterior chamber, flare and fine keratic precipitates bilaterally. Ophthalmoscopy revealed diffuse, white perivascular sheathing of the arteries and veins, tortuous and dilated veins, scattered retinal haemorrhages and papilloedema in both eyes (Fig. 1A). Fluorescein angiography demonstrated extensive dye leakage from almost all vessels (Fig. 1B) and both optic discs were hyperfluorescent in the late phase. Indocyanine green angiography (ICGA) showed a mottled background pattern, staining of the vessels especially in the posterior pole, and hypofluorescence of the discs with blurred margins (Fig. 1C). Optical coherence tomography (OCT Model 2000; Carl Zeiss Meditec, Inc., Dublin, CA, USA) showed a clear serous detachment and oedema in the macula (Fig. 1D, E). Goldmann perimetry showed an enlarged blind spot. The amplitudes of the a- and b-waves and the oscillatory potentials of the electroretinograms (ERGs) elicited by

a bright flash were significantly reduced (Fig. 1F).

Systemic examinations disclosed no abnormal findings except for high titres of anti-streptolysin O (ASO, 1830x) and anti-streptokinase (ASK, 5120x).

From these findings, the subject was diagnosed with frosted branch angiitis (FBA). He was treated with topical 0.1% betamethasone and 1.0% atropine along with i.v. prednisolone (580 mg/day) for 3 days and subsequently oral prednisolone for 15 days (total 165 mg). Three days later, the aqueous inflammation and the perivascular sheathing were markedly reduced. Thereafter, OCT and ophthalmoscopy of the macular area showed a resolution of the oedema followed by the appearance of a star-shaped figure (Fig. 1G, H). The subject's ERGs were markedly improved but were still not normal (Fig. 1I). The serous detachment (Fig. 1J, K) and finally exudates disappeared.

Despite the rapid improvement in the appearance of the retina, the patient's VA improved much more slowly and attained 1.0 only after 3 months. Fundus angiography at that time demonstrated the complete resolution of all abnormal findings.

The increase in the number of cases diagnosed as FBA raises questions

about whether this disorder represents a distinct clinical syndrome or merely a clinical sign of eyes with inflammatory conditions. From their descriptions in the literature, cases diagnosed as FBA are characterized by bilateral, perivascular retinal blood vessel sheathing, retinal haemorrhages and oedema of different degrees, panuveitis and papilloedema (Walker et al. 2004). Our case had diffuse angiitis involving the vessels of the macular and peripheral retina and the choroid. Anterior uveitis and papilloedema were also present. The abnormal ICGA findings suggested that the choroid was also altered, which probably accounts for the decreased a-waves on ERG. The reduced b-waves indicated alterations to the inner retina. Whether these findings are also found in other chorioretinal inflammatory diseases is yet to be determined. The prevalence of FBA is reported to be higher in Asians than in White populations (Walker et al. 2004).

Despite the marked panophthalmic inflammation, the eye responded well to steroids. Our subject's VA recovered to normal levels, but some FBA subjects suffer a permanent loss of vision caused by macular scarring (Kleiner et al. 1988; Walker et al. 2004).