

# 15(S)-HETE Production in Human Retinal Microvascular Endothelial Cells by Hypoxia: Novel Role for MEK1 in 15(S)-HETE-Induced Angiogenesis

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**PURPOSE.** To examine for the expression of 15-lipoxygenase 1 (15-LOX1) and 15-LOX2 in human retinal microvascular endothelial cells (HRMVECs) and study the role of arachidonic acid metabolites of these enzymes in angiogenesis.

**METHODS.** Quantitative RT-PCR and reverse-phase HPLC analyses were used to determine 15-LOX1/2 expression and their arachidonic acid metabolites in HRMVECs. The role of MEK1 in 15(S)-HETE-induced angiogenesis was studied using HRMVEC migration, tube formation, and basement membrane matrix plug angiogenesis.

**RESULTS.** HRMVECs expressed both 15-LOX1 and 15-LOX2. Hypoxia induced the expression of 15-LOX1 and the production of its arachidonic acid metabolites 15(S)-hydroxyeicosatetraenoic acid (15(S)-HETE) and 12(S)-hydroxyeicosatetraenoic acid (12(S)-HETE). 15(S)-HETE stimulated HRMVEC migration and tube formation as potently as 20 ng/mL fibroblast growth factor-2 (FGF-2). In addition, 15(S)-HETE stimulated the phosphorylation of ERK1/2, JNK1, p38 MAPK, and MEK1 in a time-dependent manner in these cells. Inhibition of MEK1 by pharmacologic and dominant-negative mutant approaches attenuated 15(S)-HETE-induced phosphorylation of ERK1/2 and JNK1 but not p38 MAPK. Blockade of ERK1/2 and JNK1 activation suppressed 15(S)-HETE-induced HRMVEC migration and tube formation and basement membrane matrix plug angiogenesis. Inhibition of p38 MAPK attenuated 15(S)-HETE-induced HRMVEC migration only. Inhibition of MEK1 also blocked 15(S)-HETE-induced HRMVEC migration and tube formation and basement membrane matrix plug angiogenesis.

**CONCLUSIONS.** These results suggest that hypoxia, through the induction of 15-LOX1 expression, leads to the production of 15(S)-HETE in HRMVECs. In addition, 15(S)-HETE, through MEK1-dependent activation of ERK1/2 and JNK1, stimulates the angiogenic differentiation of HRMVECs and basement membrane matrix plug angiogenesis. (*Invest Ophthalmol Vis Sci.* 2007;48:4930–4938) DOI:10.1167/iovs.07-0617

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In addition to its prominent role in normal development and wound healing,<sup>1,2</sup> angiogenesis plays an important role in the progression of several diseases, including atherosclerosis, cancer, and diabetic retinopathy.<sup>3–7</sup> In particular, angiogenesis appears to be a critical factor in proliferative vitreoretinal diseases such as proliferative vitreoretinopathy (PVR) and proliferative diabetic retinopathy (PDR).<sup>6,7</sup> Under hypoxic conditions, several metabolic consequences may take place, including the cessation of oxidative phosphorylation. As a result, the tissues experiencing hypoxia switch to anaerobic metabolism for their energy needs. When the use of ATP becomes greater than its production, adenosine accumulates.<sup>8,9</sup> Similarly, impairment in the blood supply results in the accumulation of metabolic products such as lactate and CO<sub>2</sub> in the interstitial space. Although some of these metabolites dilate blood vessels and thus increase the blood supply to the affected regions in an acute manner,<sup>10,11</sup> under sustained hypoxic conditions they may also act as angiogenic factors.<sup>9</sup> As an adaptive response to low-oxygen tension, the endothelial cells may produce a variety of growth factors, including vascular endothelial growth factor (VEGF),<sup>12</sup> fibroblast growth factor-2 (FGF-2),<sup>13</sup> insulin-like growth factor-1 (IGF-1),<sup>14</sup> and platelet-derived growth factor (PDGF),<sup>15</sup> to stimulate their own growth and new blood vessel formation in the surrounding area. Among these factors, VEGF is the most potent in stimulating angiogenesis in retina under hypoxic conditions.<sup>6</sup>

A large body of data indicates that lipoxygenases play a role in the pathogenesis of various diseases, including cancer and atherosclerosis.<sup>16–19</sup> In delineating the mechanisms underlying the involvement of lipoxygenases in these disease processes, it was reported that 12(S)-hydroxyeicosatetraenoic acid (12(S)-HETE), the 12-lipoxygenase (12-LOX) product of arachidonic acid, influences the growth and motility of cancer cells and vascular smooth muscle cells.<sup>20,21</sup> Similarly, 12(S)-HETE has been shown to stimulate angiogenesis.<sup>22</sup> In regard to the role of other lipoxygenases in the pathogenesis of these diseases, it was reported that 15-LOX1, while inhibiting colorectal cancer cell growth,<sup>23,24</sup> stimulates prostate cancer cell growth.<sup>17,18,25</sup> In contrast, 15-LOX2 has been shown to inhibit prostate cancer growth.<sup>26</sup> Although 15-LOX1 preferentially converts linoleic acid to 13(S)-hydroxy octadecadienoic acid (13(S)-HODE), it also metabolizes arachidonic acid to 15(S)-HETE and 12(S)-HETE.<sup>27</sup> On the other hand, 15-LOX2 converts arachidonic acid exclusively to 15(S)-HETE.<sup>27</sup> In regard to the actions of 15-LOX1/2 metabolites of arachidonic acid/linoleic acid in these cells, only 13(S)-HODE mimicked the effects of 15-LOX1 on the inhibition of colorectal cancer cell growth.<sup>23,24</sup> On the other hand, 15(S)-HETE exhibited proangiogenic activity in endothelial cells<sup>28</sup> and enhanced the growth of some cancer cell types such as erythroid cells.<sup>16</sup> Thus, 15-LOX1/2 and its products, 13(S)-HODE and 15(S)-HETE, exhibited differential effects in different cancer cell types. In addition to these observations, it was reported that on exposure to hypoxia, human neonatal vessels produce levels of 15(S)-HETE higher than normoxic levels.<sup>29</sup>

Because 15(S)-HETE induced growth in some cancer cell types and its production was increased in hypoxia-exposed neonatal vessels, we sought to determine whether this eicosanoid stimulates neovascularization and thereby contributes to the pathogenesis of PDR. In the present study we showed that hypoxia induces the expression of 15-LOX1 and the production of its arachidonic acid metabolites 15(S)-HETE and 12(S)-HETE. In addition, 15(S)-HETE through MEK1-dependent activation of ERK1/2 and JNK1 induced the angiogenic differentiation of HRMVEC and basement membrane matrix (Matrigel; BD Biosciences, Bedford, MA) plug angiogenesis.

## MATERIALS AND METHODS

### Reagents

Aprotinin, dithiothreitol, HEPES, leupeptin, phenylmethylsulfonyl fluoride (PMSF), sodium deoxycholate, and sodium orthovanadate were purchased from Sigma Chemical Company (St. Louis, MO). Nordihydroguaiaretic acid (NDGA) was obtained from Aldrich Chemical Company (Milwaukee, WI). Anti-leukocyte 12-LOX antibodies (catalog no. 160304), 5(S)-HETE, 12(S)-HETE, and 15(S)-HETE were bought from Cayman Chemicals (Ann Arbor, MI). Growth factor-reduced basement membrane matrix (Matrigel; catalog no. 354250) was obtained from BD Biosciences (Bedford, MA). Phosphospecific anti-ERK1/2 (catalog no. 9101), anti-JNK1 (catalog no. 9251), anti-p38 MAPK (catalog no. 9216), and anti-MEK1/2 (catalog no. 9121) antibodies were bought from Cell Signaling Technology (Beverly, MA). Anti-ERK2 (catalog no. SC-154), anti-JNK1 (catalog no. SC-474), anti-MEK1 (catalog no. SC-219), anti-p38 MAPK (catalog no. SC-535), and anti- $\beta$ -tubulin (catalog no. SC-9104) antibodies were obtained from Santa Cruz Biotechnology, Inc (Santa Cruz, CA). Anti-15-LOX2 antibodies were bought from Oxford Biomedical Research (Oxford, MI). PD98059 (catalog no. 513000), SP600125 (catalog no. 420119), and SB203580 (catalog no. 559389) were procured from Calbiochem Chemicals (San Diego, CA). [ $^3$ H]-Arachidonic acid (S.A. 60 Ci/mmol) was obtained from Perkin

Elmer (Boston, MA). All the primers were made by IDT (Coralville, IA). All experiments were performed in accordance with the ARVO Statement for the Use of Animals in Ophthalmic and Vision Research and were approved by the Animal Care and Use Committee of the University of Tennessee Health Science Center (Memphis, TN).

### Construction of Adenoviral Vectors

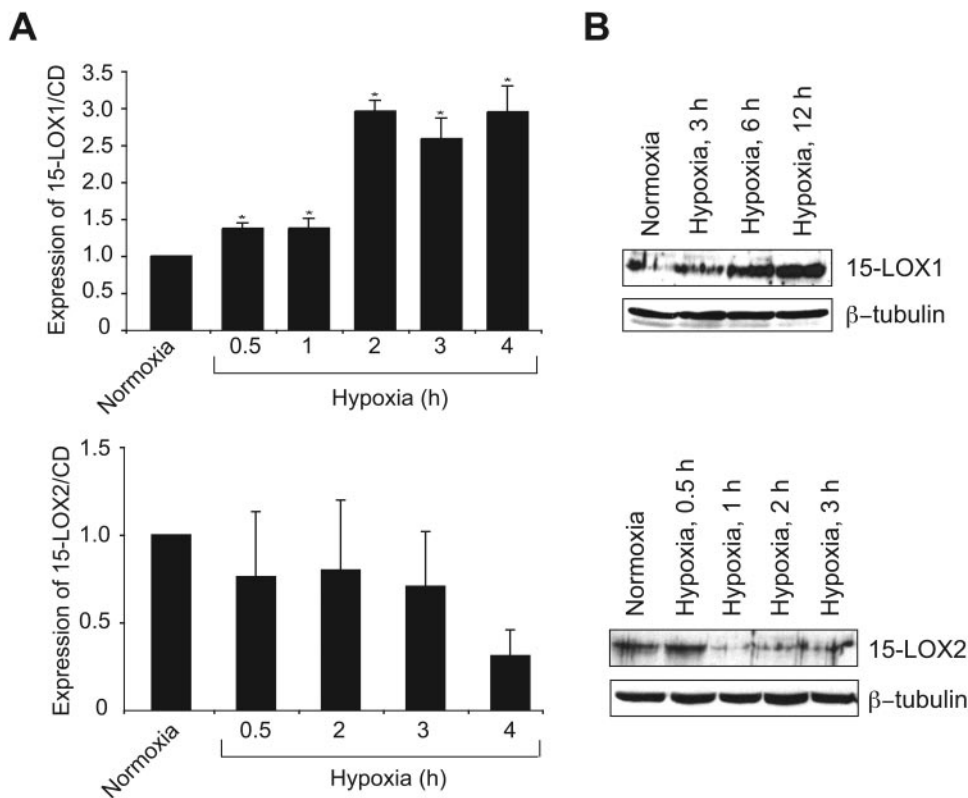
The human dnMEK1 cDNA fragment was released from pMCL-hMMK1-K97M by digestion with *KpnI* and *HindIII*<sup>30</sup> and were cloned into the same sites of pBluescript II SK (+) vector. The human dnMEK1 cDNA was then cloned into *KpnI* and *NotI* sites of entry vector pENTR3C after its release from the pBluescript II SK (+) plasmid by digestion with the same restriction enzymes. The pAd-dnMEK1 construct was obtained by recombination between pdnMEK1-ENTR3C and pAdCMV5DEST (Invitrogen, Carlsbad, CA) and was verified by DNA sequencing. The construction of pAd-GFP was described previously.<sup>31</sup> pAd-GFP and pAd-dnMEK1 were linearized with *PacI* and transfected into HEK293A cells. The resultant adenovirus was further amplified by infection of HEK293A cells and was purified by cesium chloride gradient ultracentrifugation.<sup>32</sup>

### Cell Culture

Human retinal microvascular endothelial cells (HRMVECs; catalog no. ACBRI 181) were purchased from Applied Cell Biology Research Institute (Kirkland, WA). Human dermal microvascular endothelial cells (HDMVECs; catalog no. C-011-5C) were purchased from Cascade Biologics (Portland, OR). Cells were grown in medium 131 containing microvascular growth supplements (MVGs), 10  $\mu$ g/mL gentamicin, and 0.25  $\mu$ g/mL amphotericin B. Cultures were maintained at 37°C in a humidified 95% air/5% CO<sub>2</sub> atmosphere. Cells underwent quiescence by incubation in medium 131 for 24 hours and were used for the experiments unless otherwise indicated.

### Hypoxia

To achieve hypoxia (1% O<sub>2</sub>), a preanalyzed air mixture (5% CO<sub>2</sub>/95% N<sub>2</sub>) was infused into airtight chamber-containing cells (Billups-Rothen-



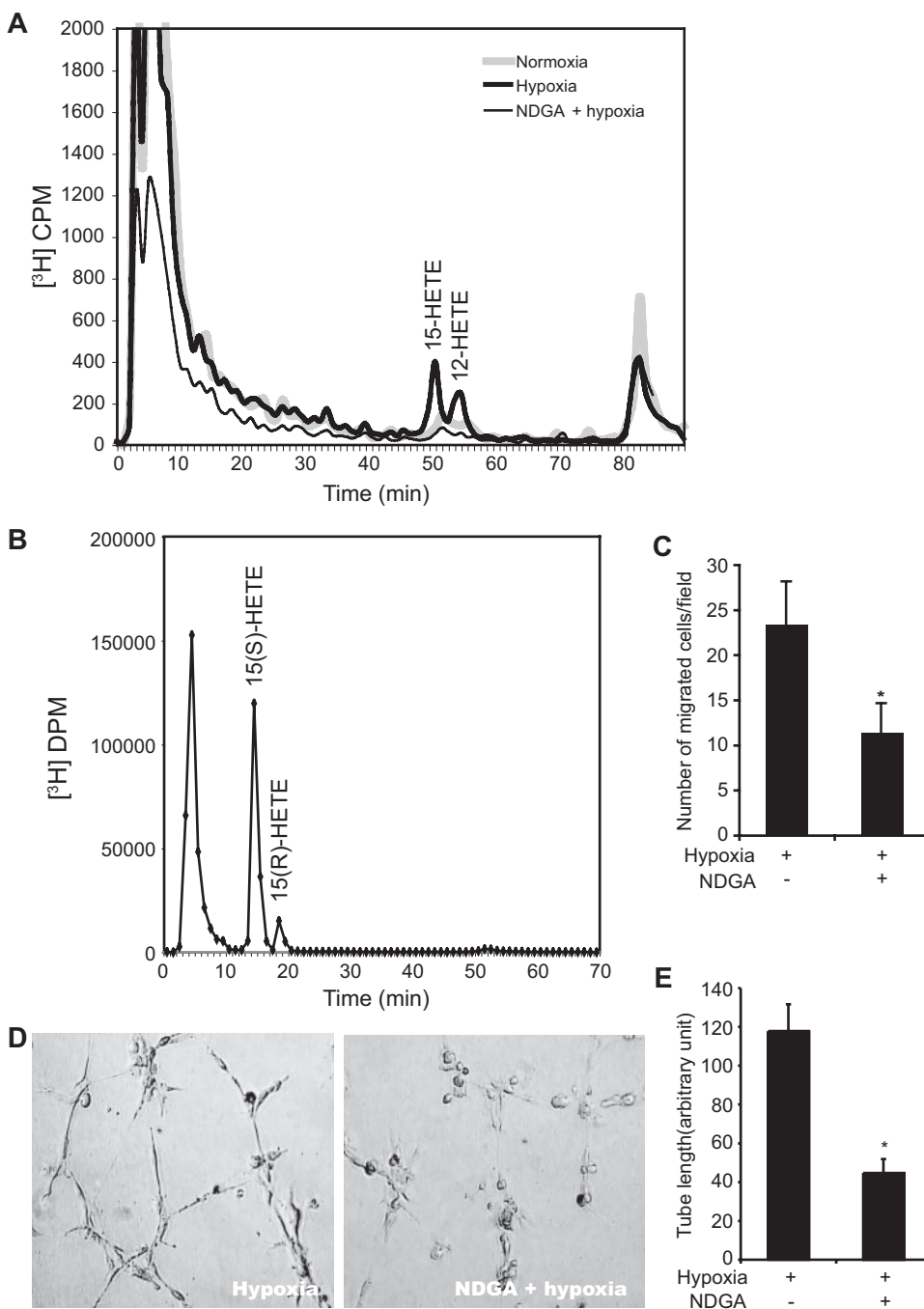
**FIGURE 1.** Differential regulation of 15-LOX1 and 15-LOX2 in HRMVECs by hypoxia. **(A)** An equal amount of total cellular RNA from control and various time points of hypoxia-treated HRMVECs was analyzed for 15-LOX1 and 15-LOX2 mRNA levels by qRT-PCR. **(B)** An equal amount of protein from control and various time points of hypoxia-treated HRMVECs was analyzed by Western blotting for 15-LOX1 and 15-LOX2 protein levels using their specific antibodies. Values are the mean  $\pm$  SD of three independent experiments. \*  $P < 0.01$  versus control.

berg Inc., Delmar, CA) at a flow rate of 3 L/min per 15 minutes. The hypoxic chamber was then placed in a tissue culture incubator at 37°C for the desired length of time.

### Reverse- and Chiral-Phase HPLC Analyses of Arachidonic Acid Metabolites

HRMVECs labeled with [<sup>3</sup>H]-arachidonic acid (0.5 μCi/mL), after quiescence in medium 131, were subjected to hypoxia for the indicated times. The eicosanoids released into the medium were extracted in methanol at a final concentration of 10% (vol/vol). The cell monolayer was rinsed with methanol, and the rinse was added to this mixture. The pH of the mixture was adjusted to 2.5 with 10% (vol/vol) glacial acetic acid, and it was then passed through a solid-phase extraction column (Pre-Sep C18; J. T. Baker, Phillipsburg, NJ). Arachidonic acid metabolites were eluted with 100% methanol followed by evaporation

to dryness. The metabolites were reconstituted in 50 μL of 100% methanol and were analyzed by reverse-phase high-performance liquid chromatography using C18 column (5-μm particle size, 4.6 × 250 mm; Beckman Instruments, Berkeley, CA), as described, using methanol/water/acetic acid with 55:45:0.01 and 75:25:0.01 ratios and 100% methanol as mobile phases at a flow rate of 1.0 mL/min for 40 minutes, 30 minutes, and 20 minutes, respectively.<sup>33,34</sup> For identification of the racemic state of 15-HETE, cells were incubated with 10 μCi [<sup>3</sup>H]-arachidonic acid under hypoxia at 37°C for 30 minutes, and the eicosanoids released into the medium were extracted and analyzed by reverse-phase HPLC. The 15-HETE fractions were collected and subjected to chiral-phase HPLC (Whelk-O1 column) with 4-(3,5-dinitrobenzamido) tetrahydrophenanthrene as a stationary phase and hexane/isopropanol/acetic acid (100:1.5:0.1) as a mobile phase. One-milliliter fractions were collected, and the radioactivity in the fractions was

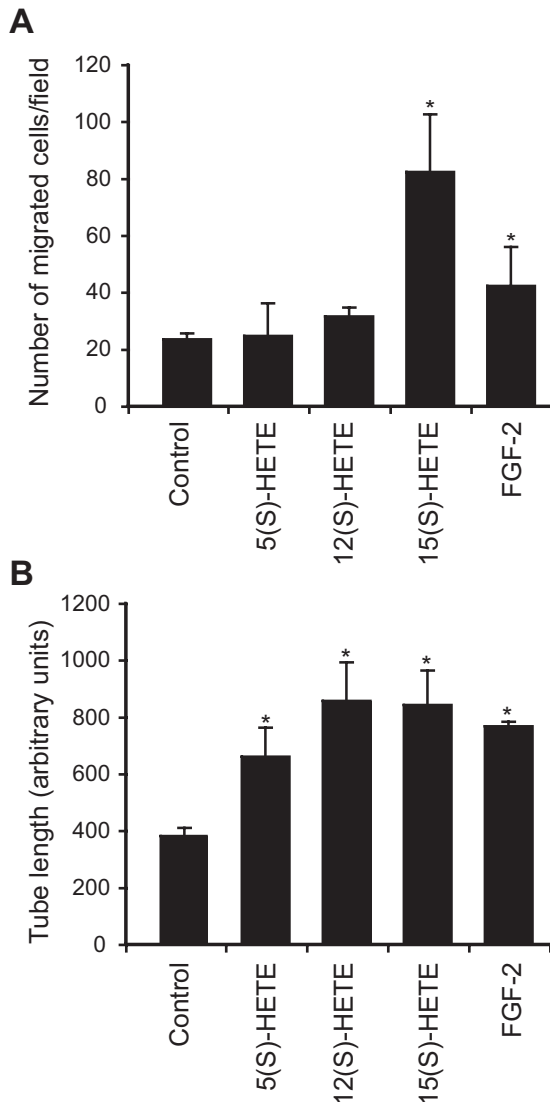


**FIGURE 2.** Hypoxia induced the production of 15(S)-HETE and 12(S)-HETE in HRMVECs. **(A)** HRMVECs that were prelabeled with [<sup>3</sup>H]-AA and that underwent quiescence were kept at normoxia or subjected to hypoxia for 6 hours in the presence and absence of NDGA (25 μM), a potent inhibitor of LOX. At the end of the incubation period, the eicosanoids released into the culture medium were extracted and analyzed by reverse-phase HPLC. **(B)** All the conditions were the same as **(A)** except that cells were subjected to hypoxia for 30 minutes in the presence of 10 μCi [<sup>3</sup>H]-arachidonic acid and 10 μM calcium ionophore A23187. After separation by reverse-phase HPLC, 15-HETE fractions were collected and subjected to chiral-phase HPLC. **(C)** HRMVEC migration was measured using modified Boyden chambers under hypoxia in the presence and absence of 25 μM NDGA. **(D, E)** HRMVEC tube formation was measured in growth factor-reduced basement membrane matrix-coated 24-well plates under hypoxia in the presence and absence of 25 μM NDGA. Representative tube formation **(D)** and quantitative data **(E)**. Values are the mean ± SD of three independent experiments. \**P* < 0.01 versus hypoxia treatment alone.

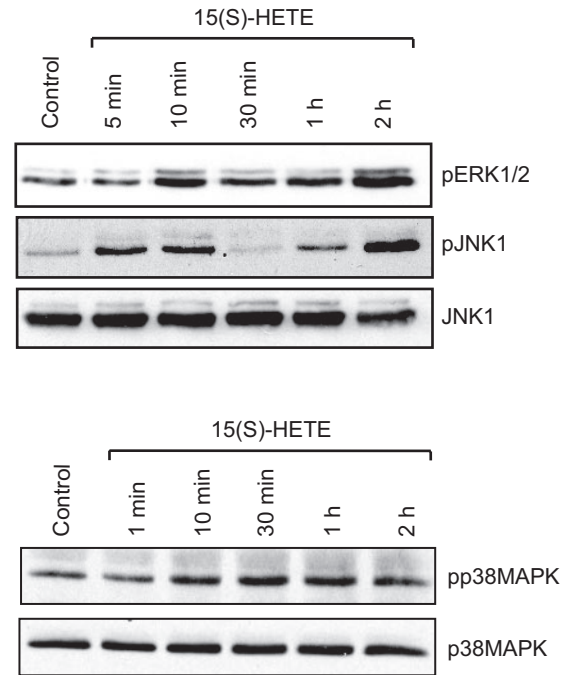
determined by mixing them with full spectrum fluorescent light (Ecolume; Lumiram, Larchmont, NY) scintillation fluid and counting in a liquid scintillation counter.

### Real-Time Quantitative RT-PCR

Total RNA was isolated from control and hypoxia-treated HRMVECs using TRIzol reagent according to the manufacturer's protocol (Sigma Chemical, St. Louis, MO). To carry out the real-time quantitative RT-PCR (qRT-PCR) amplification of 15-LOX-1 and 15-LOX-2, cDNA was generated with 1  $\mu$ g RNA from control and hypoxia-treated HRMVECs using a cDNA RT kit (High-Capacity cDNA Reverse Transcription; catalog no. 4368814; Applied Biosystems, Foster City, CA). Complementary DNA synthesis was performed in 0.2-mL PCR tubes consisting of 100  $\mu$ L of the following reaction mixture: 1  $\mu$ g RNA, 10  $\mu$ L 10 $\times$  random primers, 4  $\mu$ L 25 $\times$  dNTPs (100 mM), 5  $\mu$ L reverse transcriptase (50 U/ $\mu$ L), and 10  $\mu$ L RT buffer; the rest was RNase-free H<sub>2</sub>O.



**FIGURE 3.** Effects of 5(S)-HETE, 12(S)-HETE, 15(S)-HETE, and FGF-2 on HRMVEC migration and tube formation. (A) HRMVEC migration in response to vehicle or 0.1  $\mu$ M of the indicated HETE or 20 ng/mL of FGF-2 was measured by modified Boyden chamber method. (B) HRMVEC tube formation in response to vehicle or 0.1  $\mu$ M of the indicated HETE or 20 ng/mL of FGF-2 was measured in growth factor-reduced basement membrane matrix-coated 24-well plates. Values are the mean  $\pm$  SD of three independent experiments. \*  $P < 0.01$  versus control.



**FIGURE 4.** 15(S)-HETE stimulates phosphorylation of ERK1/2, JNK1, and p38 MAPK in a time-dependent manner in HRMVECs. Quiescent HRMVECs were treated with vehicle or 0.1  $\mu$ M 15(S)-HETE for the indicated times, and cell extracts were prepared. An equal amount of protein from control and each treatment was analyzed by Western blotting for phosphorylated ERK1/2, JNK1, and p38 MAPK levels using their phosphospecific antibodies. As a loading control, the blots were reprobed with anti-JNK1 or anti-p38 MAPK antibodies.

Messenger RNA copy numbers for 15-LOX1, 15-LOX2, and cyclophilin D were determined by qRT-PCR using a fluorescence temperature rapid-air cycle (Light Cycler 480; Roche Applied Science, Indianapolis, IN). Amplifications were performed in a 10- $\mu$ L reaction mixture containing 2  $\mu$ L cDNA solution, 0.1  $\mu$ L Universal Library Probe (10  $\mu$ M), 0.1  $\mu$ L forward primer (20  $\mu$ M), 0.1  $\mu$ L reverse primer (20  $\mu$ M), 5  $\mu$ L LC 480 master mix (2 $\times$ ), and 2.7  $\mu$ L H<sub>2</sub>O. Standard curves were generated for each assay run. All primers for real-time qRT-PCR assays spanned at least one intron, so that any containing DNA did not contribute to the amplifier on which quantification was based. The primers for 15-LOX1 were 5'-AGC CTG ATG GGA AAC TCT TG-3' (forward) and 5'-AGG TGG TGG GGA TCC TGT-3' (reverse), which gave a 68-nt amplified product. The primers for 15-LOX2 were 5'-GAT CTT CAA CTT CCG GAG GAC-3' (forward) and 5'-ACT GGG AGG CGA AGA AGG-3' (reverse), which gave an 80-nt amplified product. The primers for cyclophilin D were 5'-GGA GAC TTC TCA AAT CAG AAT GG-3' (forward) and 5'-ACC CTC CCG ATC ATG CTT-3' (reverse), which gave an amplified product of 96 nt. UPL probes number 42 with sequence 5'-CATCCAGC-3' (catalog no. 04688015001), number 51 with sequence 5'-CCCAGCAG-3' (catalog no. 04688481001), and number 39 with sequence 5'-AGGTGGAG-3' (catalog no. 04687973001) were used as internal standard probes for monitoring the amplification of 15-LOX1, 15-LOX2, and cyclophilin D, respectively. Real-time qRT-PCR consisted of 50 cycles with the following steps: denaturation at 95°C for 30 seconds, annealing at 60°C for 30 seconds, and extension 72°C for 30 seconds. The abundance of 15-LOX1 and 15-LOX 2 was expressed relative to the cyclophilin D levels.

### Cell Migration Assay

Cell migration was performed using a modified Boyden chamber method, as described by Nagata et al.<sup>35</sup> Cell culture inserts containing membranes with 10-mm diameter and 8- $\mu$ m pore size (Nalgene Nunc International, Rochester, NY) were placed in a 24-well tissue culture

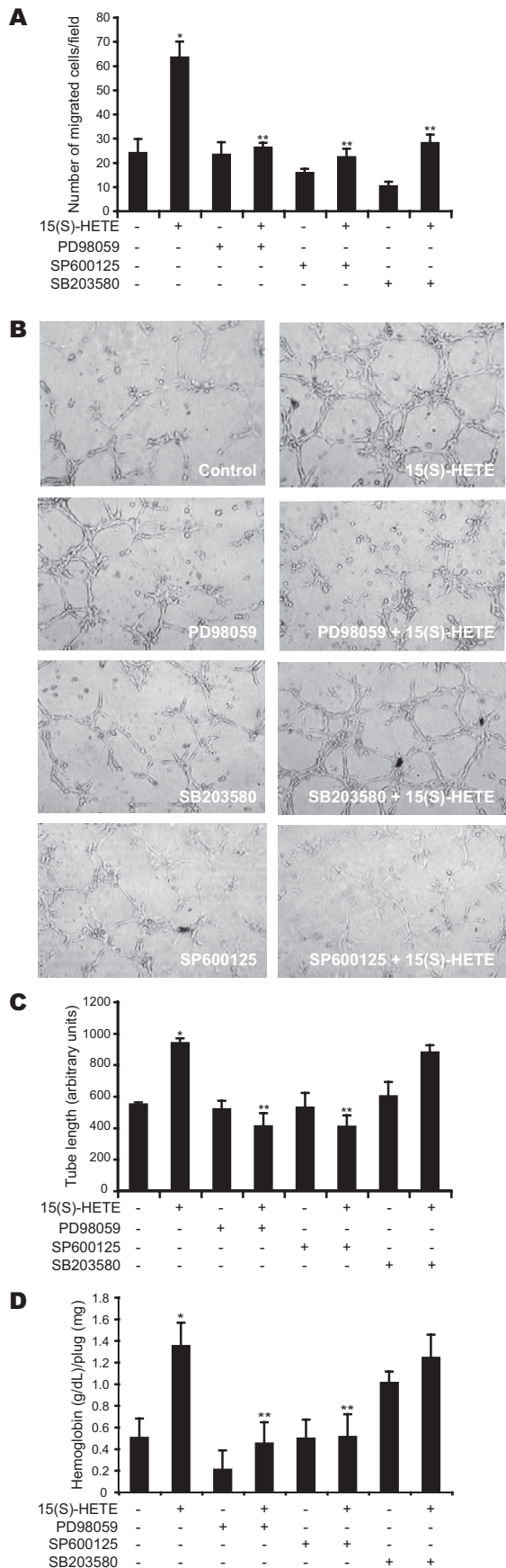


plate (Costar; Corning Incorporated, Corning, NY). The lower surface of the porous membrane was coated with 0.5% gelatin at 4°C overnight and then blocked with 0.1% heat-inactivated BSA at 37°C for 1 hour. HRMVECs underwent quiescence for 24 hours in medium 131, trypsinized, and neutralized with trypsin neutralizer solution. Cells were seeded onto the upper chamber at  $1 \times 10^5$  cells/well. Vehicle or eicosanoid of interest was added to the lower chamber at the indicated concentration. Both the upper and the lower chambers contained medium 131. When the effect of dominant-negative MEK1 was tested on 15(S)-HETE-induced HRMVEC migration, cells were infected first with either Ad-GFP or Ad-dnMEK1 at a multiplicity of infection (MOI) of 80 and underwent quiescence before they were subjected to migration assay. After 8 hours of incubation at 37°C, nonmigrated cells were removed from the upper side of the membrane with cotton swabs, and the cells on the lower surface of the membrane were fixed in methanol for 15 minutes. The membrane was then stained with hematoxylin for 10 minutes and was washed once each with 50% and 100% ethanol. Cells were counted in five randomly selected squares per well under a light microscope (Eclipse 50i; Nikon, Tokyo, Japan) and were presented as number of migrated cells per field.

### Tube Formation Assay

Tube formation assay was performed as described by Nagata et al.<sup>35</sup> Twenty-four-well culture plates (Costar; Corning Incorporated) were coated with growth factor-reduced basement membrane matrix (Matrigel; BD Biosciences) in a total volume of 280  $\mu$ L/well and were allowed to solidify for 30 minutes at 37°C. HRMVECs were trypsinized, neutralized with trypsin neutralizer solution, and resuspended at  $1 \times 10^5$ /mL, and 200  $\mu$ L of this cell suspension was added to each well. Vehicle or eicosanoid of interest, at the indicated concentration, was added to the appropriate well, and the cells were incubated at 37°C for 6 hours. When the effect of dnMEK1 was tested on 15(S)-HETE-induced HRMVEC tube formation, cells were infected first with either Ad-GFP or Ad-dnMEK1 at an MOI of 80 and then underwent quiescence before they were subjected to tube formation. Tube formation was observed under an inverted microscope (Eclipse TS100; Nikon). Images were captured with a charge-coupled device color camera (KP-D20AU; Hitachi, Honshu, Japan) attached to the microscope, and the tube length was measured using the National Institutes of Health Image J 1.31 v Program.

### Western Blot Analysis

After appropriate treatments, HRMVECs were rinsed with cold phosphate-buffered saline (PBS) and lysed in 500  $\mu$ L lysis buffer (PBS, 1% Nonidet P-40, 0.5% sodium deoxycholate, 0.1% SDS, 100  $\mu$ g/mL PMSF, 100  $\mu$ g/mL aprotinin, 1  $\mu$ g/mL leupeptin, and 1 mM sodium orthovanadate) on ice for 20 minutes. Cell lysates were scraped into 1.5-mL

**FIGURE 5.** Effect of blockade of ERK1/2, JNK1, and p38 MAPK on 15(S)-HETE-induced HRMVEC migration and tube formation and basement membrane matrix plug angiogenesis. (A) HRMVEC migration in response to vehicle or 0.1  $\mu$ M 15(S)-HETE in the presence and absence of PD98059 (30  $\mu$ M), SP600125 (10  $\mu$ M), or SB203580 (10  $\mu$ M) was measured by the modified Boyden chamber method. (B, C) HRMVEC tube formation in response to vehicle or 0.1  $\mu$ M 15(S)-HETE in the presence and absence of PD98059 (30  $\mu$ M), SP600125 (10  $\mu$ M), or SB203580 (10  $\mu$ M) was measured in growth factor-reduced basement membrane matrix-coated 24-well plate. Representative tube formation (B) and quantification data (C). (D) C57BL/6 mice were injected subcutaneously with 0.5 mL basement membrane matrix premixed with vehicle or 50  $\mu$ M 15(S)-HETE, with and without PD98059 (50  $\mu$ M), SP600125 (50  $\mu$ M), or SB203580 (50  $\mu$ M). Seven days later, the animals were killed, and the basement membrane matrix plugs were harvested from under the skin and analyzed for hemoglobin with Drabkin reagent. Values are the mean  $\pm$  SD of three independent experiments or four animals. \* $P < 0.01$  versus control. \*\* $P < 0.01$  versus 15(S)-HETE treatment alone.

Eppendorf tubes and cleared by centrifugation at 12,000 rpm for 20 minutes at 4°C. Cell lysates containing equal amounts of protein were resolved by electrophoresis on 0.1% SDS and 10% polyacrylamide gels. Proteins were transferred electrophoretically to a nitrocellulose membrane (Hybond; Amersham Pharmacia Biotech, Piscataway, NJ). After blocking in 10 mM Tris-HCl buffer, pH 8.0, containing 150 mM sodium chloride, 0.1% Tween 20, and 5% (wt/vol) nonfat dry milk, the membrane was treated with appropriate primary antibodies followed by incubation with horseradish peroxidase (HRP)-conjugated secondary antibodies. Antigen-antibody complexes were detected using chemiluminescence reagent kit (GE Health Sciences, Piscataway, NJ).

### Basement Membrane Matrix Plug Angiogenesis Assay

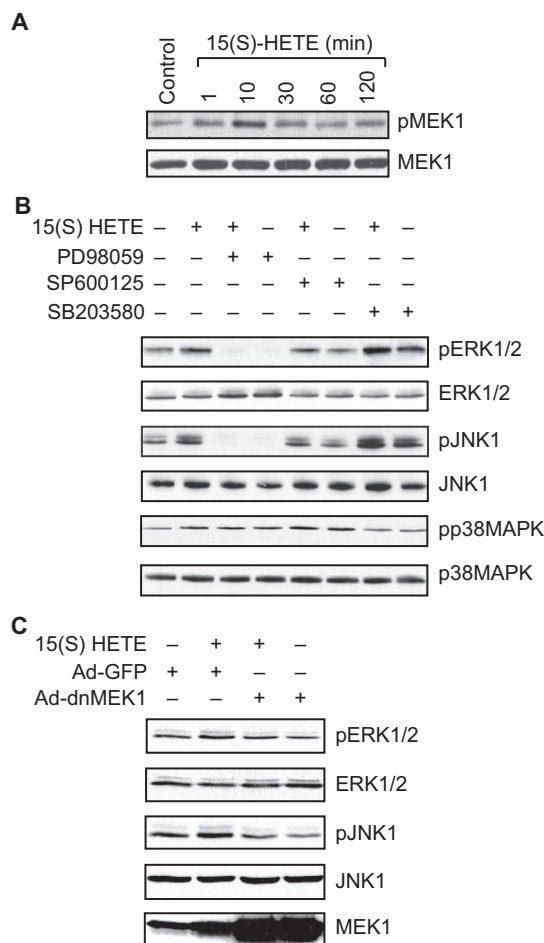
Basement membrane matrix (Matrigel; BD Biosciences) plug assay was performed essentially as described by Medhora et al.<sup>36</sup> C57BL/6 male mice (8 weeks old) were lightly anesthetized with sodium pentobarbital (50 mg/kg, administered intraperitoneally) and were injected subcutaneously with 0.5 mL basement membrane matrix (Matrigel; BD Biosciences) premixed with vehicle or 50  $\mu$ M of 15(S)-HETE along the dorsal midline. The injection was made rapidly with a Becton-Dickinson 26-gauge half-inch needle to ensure the entire content was delivered as a single plug. Whenever the effects of pharmacologic inhibitors, Ad-GFP or Ad-dnMEK1 were tested on 15(S)-HETE-induced angiogenesis, they were added to the basement membrane matrix (Matrigel; Becton-Dickinson) before they were injected into mice. The mice were allowed to recover, and 7 days later, unless otherwise stated, the animals were killed by inhalation of CO<sub>2</sub>, and the basement membrane matrix (Matrigel; Becton-Dickinson) plugs were harvested from under the skin. The plugs were homogenized in 1 mL deionized H<sub>2</sub>O on ice and cleared by centrifugation at 10,000 rpm for 6 minutes at 4°C. The supernatant was collected and used in duplicate to measure hemoglobin content with Drabkin reagent along with hemoglobin standard essentially according to the manufacturer's protocol (Sigma Chemical). Absorbance was read at 540 nm in an ELISA plate reader (Spectra Max 190; Molecular Devices, Eugene, OR). Each group contained four to six mice, and the values were expressed as gram per deciliter hemoglobin per milligram plug.

### Statistical Analysis

All experiments were repeated three times with a similar pattern of results. Data are presented as mean  $\pm$  SD. Treatment effects were analyzed by Student's *t*-test, and *P* < 0.05 was considered statistically significant. In the case of HPLC analysis, RT-PCR, and Western blotting, one representative set of data is shown.

## RESULTS

To understand the role of eicosanoids, particularly 15(S)-HETE, in retinal neovascularization, we first studied the effect of hypoxia on the expression of 15-LOX1/2 using qRT-PCR. Hypoxia induced a time-dependent increase in the expression of 15-LOX1 mRNA and a time-dependent decrease in the expression of 15-LOX2 mRNA (Fig. 1A). To obtain additional evidence for the effect of hypoxia on the expression of 15-LOX1/2, we also measured their protein levels. Western blot analysis of an equal amount of protein from control and various time points of hypoxia-treated HRMVEC using 15-LOX1- and 15-LOX2-specific antibodies showed a time-dependent increase in the levels of 15-LOX1 and a time-dependent decrease in the levels of 15-LOX2 in response to hypoxia (Fig. 1B). 15-LOX1 converts AA to 15(S)-HETE and 12(S)-HETE in a 3:1 ratio. On the other hand, 15-LOX2 metabolizes AA exclusively to 15(S)-HETE.<sup>27</sup> To determine whether hypoxia-induced expression of 15-LOX1 correlates with the production of 15(S)-HETE and 12(S)-HETE, HRMVECs were prelabeled with [<sup>3</sup>H]-AA, subjected to quiescence for 24 hour, and subjected to hypoxia for 6 hours in the



**FIGURE 6.** Effects of PD98059, SP600125, SB203580, and dnMEK-1 on 15(S)-HETE-induced phosphorylation of ERK1/2, JNK1, and p38 MAPK in HRMVECs. (A) Quiescent HRMVECs were treated with vehicle or 0.1  $\mu$ M 15(S)-HETE for the indicated times, and cell extracts were prepared. Equal amounts of protein from control and each treatment were analyzed by Western blotting for phosphorylated MEK1 levels using its phosphospecific antibodies. As a loading control, the same blot was reprobated with its normal antibodies. (B) Quiescent HRMVECs were treated with vehicle or 0.1  $\mu$ M 15(S)-HETE in the presence and absence of PD98059 (30  $\mu$ M), SP600125 (10  $\mu$ M), or SB203580 (10  $\mu$ M) for 10 minutes, and cell extracts were prepared. Equal amounts of protein from control and each treatment were analyzed by Western blotting for phosphorylated ERK1/2, JNK1, and p38 MAPK levels using their phosphospecific antibodies. As a loading control, the same blots were reprobated with their respective normal antibodies. (C) HRMVECs were infected with either Ad-GFP or Ad-dnMEK1 at an MOI of 80, underwent quiescence, and were treated with vehicle or 0.1  $\mu$ M 15(S)-HETE for 10 minutes, and cell extracts were prepared and analyzed for phosphorylated ERK1/2 and JNK1 levels, as described in (B).

presence and absence of NDGA, a potent inhibitor of LOXs.<sup>37</sup> The eicosanoids released into the medium were extracted and analyzed by reverse-phase HPLC. Hypoxia induced the production of 15-HETE and 12-HETE in an approximately 2:1 ratio in HRMVECs. NDGA, at a concentration of 25  $\mu$ M, completely suppressed their production (Fig. 2A). Although 15(S)-HETE/12(S)-HETE is formed by enzymatic actions of 15-LOX1/2 on AA, 15(R)-HETE is formed by nonenzymatic conversion of AA. Therefore, to find the source of production of 15-HETE, quiescent HRMVECs were exposed to hypoxia for 30 minutes in the presence of 10  $\mu$ Ci [<sup>3</sup>H]-AA and 10  $\mu$ M calcium ionophore A23187, and the eicosanoids released into the medium were

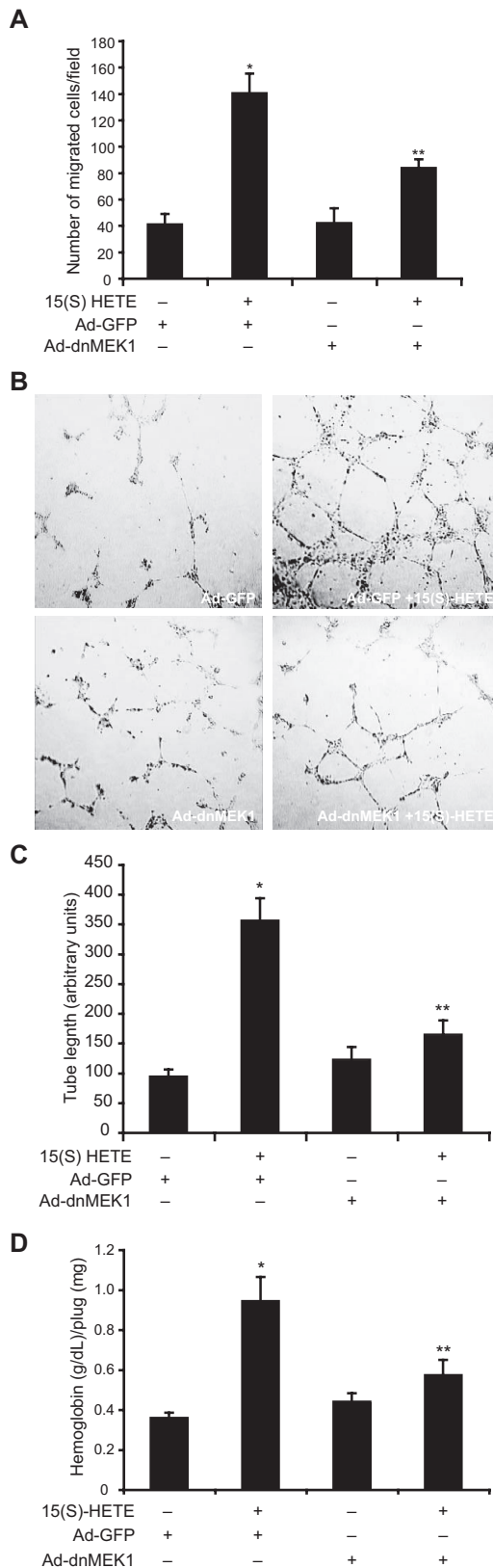
extracted and analyzed by reverse-phase HPLC. The eluate of the peak, with a retention time of 50 minutes, was collected and subjected to chiral-phase HPLC. The chiral-phase HPLC analysis of 50-minute reverse-phase HPLC fractions indicated the presence of 15(S)-HETE to a major level (90%) and 15(R)-

HETE to a minor level (10%; Fig. 2B). This result confirms the enzymatic production of 15(S)-HETE by HRMVECs in response to hypoxia. To understand the potential role of the 15-LOX1 to 15(S)-HETE axis in hypoxia-induced angiogenesis, we studied the effect of NDGA on hypoxia-induced HRMVEC migration and tube formation. As shown in Figure 2C-E, NDGA inhibited hypoxia-induced HRMVEC migration and tube formation.

To understand the relationship between HETEs and retinal neovascularization, we next tested the effect of 5(S)-HETE, 12(S)-HETE, and 15(S)-HETE on HRMVEC migration and tube formation under normoxia. Among these three HETEs, only 15(S)-HETE increased HRMVEC migration to a significant level (threefold vs. control). Furthermore, the chemotactic effect of 15(S)-HETE was higher than that caused by 20 ng/mL FGF-2 (Fig. 3A). Although all three HETEs induced HRMVEC tube formation, the effect of 12(S)- and 15(S)-HETEs was slightly higher than that of 5(S)-HETE and equal to that of 20 ng/mL FGF-2 (Fig. 3B).

Because human neonatal vessels and HRMVECs produced higher levels of 15(S)-HETE under hypoxic than normoxic conditions, we next focused on the elucidation of mechanisms underlying its angiogenic effects. For these studies, we examined the role of extracellular signal-regulated kinases 1/2 (ERK1/2), Jun N-terminal kinase 1 (JNK1), and p38 mitogen-activated protein kinase (p38 MAPK) groups of MAPKs. Western blot analysis of an equal amount of protein from control and various times of 15(S)-HETE-treated HRMVECs showed a time-dependent increase in the phosphorylation of ERK1/2, JNK1, and p38 MAPK (Fig. 4). A biphasic increase in the phosphorylation of ERK1/2 was observed, with a first peak around 10 minutes after 15(S)-HETE treatment, followed by a second peak at 2 hours. Peak phosphorylation of JNK1 occurred at 10 minutes, followed by a decrease at 30 minutes and a gradual increase thereafter. On the other hand, the phosphorylation of p38 MAPK was increased at 30 minutes, and this effect was sustained for at least 2 hours (Fig. 4). PD98059, a potent inhibitor of MEK1/2,<sup>38</sup> at a concentration of 30  $\mu$ M completely suppressed 15(S)-HETE-induced HRMVEC migration and tube formation. Similarly, SP600125, a potent and specific inhibitor of JNK,<sup>39</sup> also blocked 15(S)-HETE-induced HRMVEC migration and tube formation. In contrast, SB203580, a potent inhibitor of p38 MAPK,<sup>40</sup> attenuated 15(S)-HETE-induced migration only but had no effect on tube formation (Figs. 5A-C). Interestingly, only PD98059 and SP600125, but not SB203580, inhibited 15(S)-HETE-induced basement membrane matrix (Matrigel; BD Biosciences) plug angiogenesis (Fig. 5D).

To test whether 15(S)-HETE stimulates ERK1/2 and JNK1 through a common mechanism, we next studied the role of MEK1 and found that 15(S)-HETE stimulated MEK1 phosphorylation in a time-dependent manner, with a maximum increase at 10 minutes (Fig. 6A). PD98059, an inhibitor of MEK1/2,



**FIGURE 7.** Adenoviral-mediated expression of dominant-negative MEK1 suppresses 15(S)-HETE-induced HRMVEC migration and tube formation and basement membrane matrix plug angiogenesis. HRMVECs were infected with Ad-GFP or Ad-dnMEK1 at an MOI of 80, underwent quiescence, and were subjected to vehicle or 0.1  $\mu$ M of 15(S)-HETE-induced migration (A) and tube formation (B, C) as described in the Figure 3 legend. (D) C57BL/6 mice were injected subcutaneously with 0.5 mL basement membrane matrix premixed with vehicle or 50  $\mu$ M 15(S)-HETE, with and without adenovirus expressing either GFP ( $1 \times 10^8$  pfu) or dnMEK1 ( $1 \times 10^8$  pfu). One week later, the animals were killed, and the basement membrane matrix plugs were harvested from under the skin and were analyzed for hemoglobin with Drabkin reagent. Values are the mean  $\pm$  SD of three independent experiments or four animals. \* $P < 0.01$  versus control. \*\* $P < 0.01$  versus 15(S)-HETE treatment alone.

blocked ERK1/2 phosphorylation and attenuated JNK1 phosphorylation but had no effect on p38 MAPK phosphorylation induced by 15(S)-HETE (Fig. 6B). PD98059 by itself enhanced the basal phosphorylation of p38 MAPK. SP600125 inhibited 15(S)-HETE-induced JNK1 phosphorylation but either had no effect or enhanced the phosphorylation of ERK1/2 and p38 MAPK. On the other hand, SB203580 suppressed the phosphorylation of p38 MAPK only. These results strongly suggest that both ERK1/2 and JNK1 are activated by MEK1/2 in response to 15(S)-HETE.

To substantiate this finding further, we tested the effect of dnMEK1. Adenovirus-mediated expression of dnMEK1 completely blocked ERK1/2 and JNK1 phosphorylation induced by 15(S)-HETE (Fig. 6C). To understand the role of MEK1 in 15(S)-HETE-induced angiogenesis, we tested the effect of dnMEK1 on HRMVEC migration, tube formation, and basement membrane matrix (Matrigel; BD Biosciences) plug angiogenesis. Adenovirus-mediated expression of dnMEK1 suppressed all of them (Figs. 7A-D).

## DISCUSSION

The present study yielded several important findings. HRMVECs expressed both 15-LOX1 and 15-LOX2. Hypoxia induced the expression of 15-LOX1 and the production of its arachidonic acid metabolites 15(S)- and 12(S)-HETEs. 15(S)-HETE stimulated HRMVEC migration and tube formation as potently as 20 ng/mL of FGF-2. 15(S)-HETE stimulated ERK1/2, JNK1, and p38 MAPK phosphorylation in a time-dependent manner in HRMVECs. 15(S)-HETE-induced HRMVEC migration and tube formation, and basement membrane matrix (Matrigel; BD Biosciences) plug angiogenesis required ERK1/2 and JNK1 activation. 15(S)-HETE stimulated MEK1 phosphorylation in a time-dependent manner in HRMVECs. Inhibition of MEK1 by pharmacologic treatments or transduction of its dominant-negative mutant suppressed the 15(S)-HETE-induced ERK1/2 and JNK1 phosphorylation in HRMVECs and the migration and tube formation of these cells as well as basement membrane matrix (Matrigel; BD Biosciences) plug angiogenesis.

Evidence indicates that 15-LOX1 and 15-LOX2 play opposite roles in the regulation of cell growth; the latter is predominantly associated with proapoptotic signaling.<sup>17,18,23-26</sup> Similarly, 15(S)-HETE, which could be produced by the actions of these enzymes on arachidonic acid, plays a role in the regulation of cell growth and apoptosis.<sup>16,41,42</sup> It was demonstrated that human neonatal vessels, on exposure to hypoxia, produce hydroxyacids, including 15(S)-HETE.<sup>29</sup> In addition, 15(S)-HETE enhances HRMVEC migration.<sup>28</sup> The present finding that the expression of 15-LOX1 is induced by hypoxia in HRMVECs, resulting in the production of 15(S)-HETE, implies that 15-LOX1-15(S)-HETE signaling is involved in the angiogenic differentiation of these cells. This assumption was further supported by the observation that 15(S)-HETE is the most potent of the three HETEs tested in the stimulation of HRMVEC migration and tube formation, and it was as effective as 20 ng/mL FGF-2 in inducing these angiogenic events in HRMVECs. Given that exposure of human neonatal vessels to hypoxia led to increased production of 15(S)-HETE and that it has the ability to stimulate angiogenesis,<sup>43</sup> it is possible that this eicosanoid may be involved in the pathogenesis of retinal neovascularization. Because 15-LOX1 also converts arachidonic acid to 12(S)-HETE and this eicosanoid is produced in HRMVECs in response to hypoxia, a role for 12(S)-HETE in retinal neovascularization cannot be ruled out. In fact, it was demonstrated that 12(S)-HETE induces angiogenesis and prostate tumor growth.<sup>44</sup>

Previously, we have shown that arachidonic acid stimulates ERK1/2, JNK1, and p38 MAPK in vascular smooth muscle cells

(VSMCs), responses that require its conversion through the LOX pathway.<sup>33,45,46</sup> Furthermore, arachidonic acid-induced VSMC migration required activation of ERKs, JNKs, and p38 MAPK.<sup>46</sup> In this study, we showed that 15(S)-HETE stimulates all three major groups of MAPKs—ERK1/2, JNK1, and p38 MAPK—in HRMVECs. Although a role for all three major groups of MAPKs in the regulation of cell migration has been demonstrated<sup>47-50</sup> and 15(S)-HETE-induced HRMVECs migration appeared to be dependent on the activation of ERK1/2, JNK1, and p38 MAPK, only the former two groups of MAPKs seemed to be involved in angiogenesis induced by this lipid molecule. Other studies have shown that ERKs and JNKs played roles in angiogenesis.<sup>51</sup>

A large body of data suggests that MEK1/2 (also known as MKK1/2) plays a role in the activation of ERK1/2.<sup>48</sup> Similarly, MKK4/7 and MKK3/6 have been reported to be involved in the phosphorylation and activation of JNKs and p38 MAPK, respectively.<sup>52</sup> It is interesting to note that 15(S)-HETE-induced ERK1/2 and JNK1 phosphorylation in HRMVECs is mediated by MEK1. Furthermore, blocking MEK1 activation by pharmacologic treatments or transduction of its dominant-negative mutant inhibited 15(S)-HETE-induced HRMVEC migration, tube formation, and basement membrane matrix (Matrigel; BD Biosciences) plug angiogenesis. Based on these observations, it is possible that MEK1 mediates 15(S)-HETE-induced HRMVEC angiogenic events by targeting both ERK1/2 and JNK1. A role for MAPK in serine phosphorylation and activation of STATs, particularly STAT-1 and STAT-3, has been demonstrated.<sup>53</sup> Recently, we have shown that STAT-3 is involved in 15(S)-HETE-induced VEGF expression and angiogenesis.<sup>54</sup> Based on these observations, it is possible that MEK1, by targeting ERK1/2 and JNK1, may facilitate serine phosphorylation and activation of STAT-3, leading to VEGF expression and angiogenesis. Further studies are required to address whether this is the mechanism involved in 15(S)-HETE-induced retinal angiogenesis.

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