Leukotriene B4 Receptor-1 Mediates Intermittent Hypoxia-induced Atherogenesis

Richard C. Li¹, Bodduluri Haribabu², Steven P. Mathis², Jinkwan Kim¹, and David Gozal¹

¹Department of Pediatrics, University of Chicago, Chicago, Illinois; and ²Department of Microbiology and Immunology, University of Louisville, Louisville, Kentucky

Rationale: Obstructive sleep apnea, which is characterized by intermittent hypoxia (IH) during sleep, has emerged as an independent risk factor for cardiovascular disease, including atherosclerosis. Leukotriene B4 (LTB4) production is increased in patients with obstructive sleep apnea and negatively correlates to hypoxic levels during sleep, with continuous positive airway pressure therapy decreasing LTB4 production.

Objectives: Determine the potential role of LTB4 in IH-induced atherosclerosis in a monocyte cellular model and a murine model.

Methods: THP-1 cells were exposed to IH for 3, 6, 24, and 48 hours. Macrophage transformation and foam cell formation were assessed after IH exposures. Apolipopotein E (ApoE)^{-/-} or BLT1^{-/-}/ApoE^{-/-} mice were fed an atherogenic diet and exposed to IH (alternating 21% and 5.7% O_2 from 7 AM to 7 PM each day) for 10 weeks. Atherosclerotic lesion formation in en face aorta was examined by oil red O staining. Measurements and Main Results: IH increased production of LTB4 and the expression of 5-lipoxygenase and leukotriene A4 hydrolase, the key enzymes for producing LTB4. IH was associated with transformation of monocytes to activated macrophages, as evidenced by increased expression of CD14 and CD68. In addition, IH exposures promoted increased cellular cholesterol accumulation and foam cell formation. The LTB4 receptor 1 (BLT1) antagonist U-75302 markedly attenuated IH-induced changes. Furthermore, IH promoted atherosclerotic lesion formation in ApoE $^{-/-}$ mice. IH-induced lesion formation was markedly attenuated in BLT1^{$-/-$}/ApoE^{$-/-$} mice.

Conclusions: BLT1-dependent pathways underlie IH-induced atherogenesis, and may become a potential novel therapeutic target for obstructive sleep apnea–associated cardiovascular disease.

Keywords: obstructive sleep apnea; inflammation; monocyte; atherosclerosis

Obstructive sleep apnea (OSA) is a highly prevalent disorder throughout the lifespan, which may affect up to 2–5% of the population. OSA has emerged as an independent risk factor for cardiovascular diseases, including atherosclerosis (1, 2). Indeed, independent associations between OSA and the evolution of increased carotid artery intima–media thickness and progressive narrowing of the coronary artery lumen have been reported (3– 7). The characteristic disturbances of OSA include intermittent hypoxia (IH), episodic hypercapnia, sleep fragmentation, and increased intrathoracic pressure changes during sleep (8). In

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AT A GLANCE COMMENTARY

Scientific Knowledge on the Subject

Obstructive sleep apnea, which is characterized by intermittent hypoxia (IH) during sleep, has emerged as an independent risk factor for cardiovascular diseases. In recent years, animal models have revealed that IH promotes atherosclerosis. However, the molecular mechanisms underlying IH-induced atherogenesis remain largely undefined.

What This Study Adds to the Field

Leukotriene B4 and its receptor BLT1 play a prominent role in mediating atherogenesis associated with episodic hypoxia during sleep. These findings are consistent with the overarching concept whereby IH triggers inflammatory processes that play a critical role in atherogenesis. Therefore, the leukotriene B4–BLT1 pathway may provide a potential therapeutic target for obstructive sleep apnea–associated atherosclerosis.

recent years, animal models have revealed that IH promotes atherosclerosis (9, 10), systemic hypertension (11), glycemic control disturbances (12, 13), and endothelial dysfunction (14), thereby confirming the contribution of OSA to the metabolic syndrome (15). However, the molecular mechanisms underlying IH-induced atherogenesis remain largely undefined.

It has now become established that atherosclerosis should be viewed as an inflammatory disease (16, 17), such that the mechanistic links between IH and atherosclerosis would likely be mediated by inflammatory processes elicited in the context of IH (18). In the last decade, we established rodent models of IH to enable the exploration of IH-induced pathologic consequences and their underlying mechanisms (19–24). In that context, it became apparent that IH induced prominent oxidative stress and promoted chronic inflammatory processes that could ultimately contribute to IH-associated atherogenesis.

Leukotriene B4 (LTB4) is an eicosanoid lipid derivative of arachadonic acid, the latter being generated by 5-lipoxygenase (5-LO) (25, 26) and leukotriene A4 hydrolase (LTA4H) (27). LTB4 is a proinflammatory mediator that activates multiple leukocyte subsets, leading to cell recruitment, production of reactive oxygen species, and induction of gene expression (28). LTB4 primarily binds its high-affinity G protein–coupled receptor BLT1, which is highly expressed in circulating peripheral blood leukocytes. BLT2 is another functional receptor subtype for LTB4 that exhibits much lower affinity for its ligand, and is ubiquitously expressed at low levels in many human tissues (29, 30). Mounting evidence has suggested that LTB4 is associated with several chronic inflammatory conditions, including atherosclerosis (28, 31). Indeed, LTB4 is a key mediator of inflammation, and genetic ablation of its high-affinity receptor, BLT1, delays the onset of atherosclerosis, suggesting that the LTB4–BLT1

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Correspondence and requests for reprints should be addressed to Richard C. Li, M.D., Ph.D, Department of Pediatrics, Pritzker School of Medicine, The University of Chicago, The Knapp Center for Biomedical Discovery, RM 4110, 900 E. 57th Street, Chicago, IL 60637. E-mail: rcli@uchicago.edu

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pathway plays a critical role in the development of atherosclerosis (32). Based on these findings, we hypothesized that the LTB4– BLT1 pathway plays a key role in IH-induced atherogenesis. To address our hypothesis, we used both in vitro and in vivo IH models, and explored the LTB4–BLT1 pathway contribution to atherogenesis using both pharmacologic and genetic approaches.

METHODS

Cell Culture and IH Exposures

THP-1 cells were originally obtained from American Tissue Type Cell Collection (ATCC, Manassas, VA) and were cultured in Gibco RPMI 1640 medium. IH exposures were conducted using a custom-designed computer-controlled incubator chamber attached to an external O_{2-} CO2 computer-driven servocontroller (Biospherix, Lacona, NY). This system is able continuously to record the dissolved $O₂$ concentrations in the culture medium and allows for implementation of the desired oxygen concentration profile (see Figure E1 in the online supplement). Cells were exposed to IH or room air (RA) for 3, 6, 24, and 48 hours (see online supplement).

Animals and IH Exposures

Animal care and experimental procedures were performed with approval from animal care committees of the University of Chicago, Chicago, Illinois, and the University of Louisville, Louisville, Kentucky. Apolipopotein E $(ApoE)^{-/-}$, BLT1^{-/-}ApoE^{-/-} mice were fed with a high-cholesterol atherogenic western diet (TD88137; Harlan Laboratories, Indianapolis, IN). Animals were housed in four identical commercially designed chambers under 12-hour light–dark cycles (Biospherix). After a 10-week IH exposure, mice were anesthetized. The heart and aorta were dissected. Ascending aortas were embedded in optimal cutting temperature (OCT) compound and cryosections were prepared for evaluating atherosclerotic lesions by oil red O staining (see online supplement).

Assessment of Atherosclerotic Lesions

Perfusion fixation, preparation of aortas, and quantification of atherosclerotic lesions were performed as previously described (32) (see online supplement).

Intracellular Lipid Analysis in Macrophages

Free cholesterol and total cholesterol were determined by commercial assay systems (Amplex Red Cholesterol Assay kit; Invitrogen, Carlsbad, CA). Cholesterol ester was estimated by subtracting free cholesterol from total cholesterol. Data were normalized to cellular protein content (see online supplement).

Electrophoretic Mobility Shift Assay

Electrophoretic mobility shift assay was performed using a kit from Panomics (Santa Clara, CA) (see online supplement).

Quantitative Real-Time Polymerase Chain Reaction

The mRNA expression (5-LO, LTA4H, MCP-1, CCR2, CD14, CD68, CD36 SRA, and ABCA1 and ApoA-I) was determined by quantitative real-time polymerase chain reaction.

Oil Red O Staining

Cells were differentiated into macrophages in the presence of phorbol myristate acetate (100 ng/ml) for 24 hours. After being washed three times with phosphate-buffered saline, the differentiated macrophages were cultured in RPMI 1640 plus 5% lipoprotein-deficient serum (LPDS) medium containing 50 μ g/ml acetylated low-density lipoprotein (acLDL) and exposed to IH or RA. Cells were fixed in 4% paraformaldehyde and stained with oil red O and hematoxylin.

Assessment of LTB4 Production in Culture Medium and Plasma

LTB4 production was measured by LTB4 ELISA kit (Cayman, Ann Arbor, MI).

Data Analysis

All results are reported as mean \pm SEM unless otherwise indicated. Comparisons between the IH and RA groups in the Apo $E^{-/-}$ mice and BLT1⁻ $ApoE^{-/-}$ mice were made by using analysis of variance procedures followed by *post hoc* tests. Statistical significance was assumed at *P* less than 0.05.

RESULTS

IH Induces a Significant Increased Production of LTB4 and Expression of 5-LO and LTA4H

To establish whether LTB4 production and expression of its key enzymes (5-LO and LTA4H) are altered after exposure to IH, THP-1 cells were exposed to IH for 3, 6, 24, and 48 hours. LTB4 concentrations in culture media were significantly increased after 24- and 48-hour IH exposures ($P < 0.05$; Figure 1A). mRNA expression of both 5-LO and LTA4H was increased at 24 and 48 hours of IH. However, the magnitude for IH-induced LTA4H expression was markedly greater than 5-LO, suggesting that increased LTA4H expression may play an important role in IH-induced LTB4 production (* $P < 0.01$; Figure 1B). In

Figure 1. (A) Time course of leukotriene B4 (LTB4) production in culture media after exposure of THP-1 cells to intermittent hypoxia (IH) for 3, 6, 24, and 48 hours. (B) Time course of 5-lipoxygenase (5-LO) and leukotriene A4 hydrolase (LTA4H) mRNA expression in THP-1 cells after exposure to IH for 3, 6, 24, and 48 hours. (C) Time course of BLT1 and BLT2 mRNA expression in THP-1 cells after exposure to IH for 3, 6, 24, and 48 hours. Data are expressed as a percentage of control room air (RA) (mean \pm SE; $n = 8$; $*P < 0.01$ vs. RA).

Figure 2. (A) Time course of DNA binding activity of nuclear factor-kappa B (NF-kB) and hypoxia-inducible factor (HIF)-1 α after exposure to intermittent hypoxia (IH) for 3, 6, and 24 hours. (B) Effect of PD98059 and JSH-23 on 5-lipoxygenase (5-LO) mRNA expression in THP-1 cells after exposure to IH for 24 hours. (C) Effect of PD98059 and JSH-23 on leukotriene A4 hydrolase (LTA4H) mRNA expression in THP-1 cells after exposure to IH for 24 hours. Data are expressed as a percentage of control RA (mean \pm SE; $n = 8$; * $P < 0.01$ vs. RA; $^{#}P < 0.05$ vs. IH).

contrast, mRNA expression of BLT1 was slightly increased only at 1 day of IH exposures ($P < 0.05$), but not thereafter. However, no significant changes were detected in BLT2 mRNA expression after IH exposure ($P > 0.05$; Figure 1C).

Nuclear Factor-kappa B and Mitogen-activated Protein Kinase Are Involved in Regulating IH-induced Expression of 5-LO and LTA4H

To determine whether the IH-induced mRNA expression of 5-LO and LTA4H is regulated by nuclear factor-kappa B (NF- κ B) and hypoxia-inducible factor (HIF)-1 α , DNA binding activity for NF- κ B and HIF-1 α were measured by electrophoretic mobility shift assay after IH exposures for 3, 6, and 24 hours. Increased HIF-1 α binding activity occurred only at 6 hours, and was not sustained after longer IH exposures. In contrast, NF-kB DNA binding activity was markedly increased after exposures to IH starting at 6 hours, and remained elevated even at 24 hours (Figure 2A; $*P <$ 0.01), suggesting that IH activates NF- κ B, but not HIF-1 α in THP-1 cells. To further assess whether NF- κ B and mitogenactivated protein kinase (MAPK) are involved in regulating IH-induced mRNA expression of 5-LO and LTA4H, THP-1 cells were treated with JSH-23 (10 μ M, a NF- κ B inhibitor) and PD98059 (20 μ M, a mitogen-activated kinase/ERK kinase 1 [MEK1] inhibitor) or solvent control. After 30 minutes of treatment, the cells were exposed to IH for 24 hours and mRNA expression of 5-LO and LTA4H was examined.

Figure 3. (A) Effect of U75302 and LY255283 on 5-lipoxygenase (5-LO) mRNA expression in THP-1 cells after exposure to intermittent hypoxia (IH) for 24 hours. (B) Effect of U75302 and LY255283 on leukotriene A4 hydrolase (LTA4H) mRNA expression in THP-1 cells after exposure to IH for 24 hours. (C) Diagram representing putative signalings of IH-induced leukotriene B4 (LTB4) production and its regulation. Data are expressed as a percentage of control RA (mean \pm SE; n = 8; * P < 0.01 vs. RA; [#] P < 0.05 vs. IH). MAPK $=$ mitogen-activated protein kinase; $NF-\kappa B$ = nuclear factor-kappa B.

IH-induced 5-LO and LTA4H expression were significantly attenuated by either PD98059 or JSH-23 (Figures 2B and 2C; $*P$ < 0.05), indicating that NF-kB and MAPK may play an important role in regulating IH-induced increases in LTB4 production.

BLT1 and BLT2 Potentiate IH-induced 5-LO and LTA4H Expression

To ascertain whether inhibition of BLT1 and BLT2 effectively suppresses IH-induced increases in 5-LO and LTA4H, THP-1 cells were treated with either U75302 (1 μ M, a BLT1 antagonist) or LY255283 (1 μ M, a BLT2 antagonist) (solvent served as control) followed by 24-hour IH exposures, after which mRNA expression of 5-LO and LTA4H expression were assessed. Treatment with U75302 effectively suppressed both 5-LO and LTA4H expression (Figures 3A and $3B$; $^{#}P$ < 0.01). However, LY255283 only slightly suppressed LTA4H expression but not 5-LO expression (Figures 3A and 3B), suggesting that BLT1 may operate as a component of an amplification loop. Once LTB4 production is triggered by IH, LTB4 may bind to its receptors, BLT1 and BLT2, and further promote the production of LTB4 (see diagram in Figure 3C for more detail).

Inhibition of BLT1 or BLT2 Reduces IH-associated Monocyte Recruitment and Macrophage Differentiation

To examine whether IH enhances monocyte recruitment and macrophage differentiation, and whether administration of a BLT1 or a BLT2 blocker reduces these effects, THP-1 cells were exposed to IH with or without U75302 (1 μ M) or LY255283 (1 μ M) treatment. Monocyte recruitment was determined by measuring the expression of MCP-1 and CCR2, whereas macrophage differentiation was determined by assessing the expression of CD14 and CD68. MCP-1, but not CCR2, was significantly increased after IH exposures (Figure 4A; $*P < 0.05$). In addition, IH-increased MCP-1 expression was markedly attenuated by U75302, but not by LY255283 (Figure 4A; $^{*}P < 0.05$), indicating that BLT1 may play a role in mediating IH-induced monocyte recruitment. Furthermore, both CD14 and CD68 were significantly increased after IH exposures (Figure 4B; $*P < 0.05$), suggesting that IH exposures promote monocyte differentiation. In addition,

IH-increased CD14 and CD68 expression was markedly attenuated by either U75302 or LY255283 (Figure 4B; $^{*}P$ < 0.05), indicating that both BLT1 and BLT2 may mediate IH-induced monocyte differentiation.

Inhibition of BLT1 or BLT2 Attenuates IH-induced Cholesterol Accumulation and Foam Cell Formation

To determine further whether IH increases cellular cholesterol accumulation and foam cell formation and the role of BLT1 or BLT2 in these effects, differentiated THP-1 cells were exposed to IH with or without U75302 (1 μ M) or LY255283 (1 μ M) treatment. Cellular cholesterol accumulation was determined using a commercial kit, whereas foam cell formation was determined by staining with oil red O. Intracellular cholesterol accumulation was significantly increased by IH exposures in differentiated THP-1 cells (Figure 5A; $*P < 0.05$). In addition, IH-associated cholesterol accumulation was markedly reduced by U75302, but not by LY255283 (Figure 5A; $^{#}P < 0.05$), indicating that BLT1 is likely involved. Furthermore, foam cell formation was significantly increased after IH exposures in differentiated THP-1 cells (Figure 5B), and this effect was markedly attenuated by either U75302 or LY255283 (Figure 5B), indicating that both BLT1 and BLT2 may be involved.

Inhibition of BLT1 or BLT2 Attenuates IH-induced Dysregulation of Cellular Cholesterol Transport Proteins

To elicit further the cellular mechanism by which IH promotes cellular cholesterol accumulation in differentiated cells, cholesterol influx was determined by measuring the expression of CD36 and SRA, whereas cholesterol efflux was determined by assessing the expression of ABCA1 and ApoA-I. CD36 and SRA were significantly increased by IH exposure in differentiated THP-1 cells (Figure 6A; $*P <$ 0.05), suggesting that IH exposure enhances influx cholesterol trafficking. In addition, IH-associated CD36 and SRA expression was markedly reduced by both U75302 (1 μ M) and LY255283 (1 μ M) (Figure 6B; $^{#}P$ < 0.05). Furthermore, expression of ApoA-I was slightly increased after IH exposure in differentiated THP-1 cells (Figure 6C; $*P > 0.05$), but no significant changes emerged in the

Figure 5. (A) Effect of U75302 and LY255283 on intracellular cholesterol in THP-1 cells after exposure to intermittent hypoxia (IH) for 24 hours. (B) Microscopy images representing the effect of U75302 and LY255283 on foam cell formation in THP-1 cells after exposure to IH for 24 hours. Data are expressed as a percentage of control RA (mean \pm SE; n = 8; *P < 0.01 vs. RA; $^{#}P < 0.05$ vs. IH).

Figure 6. (A) Effect of U75302 and LY255283 on CD-36 mRNA expression in THP-1 cells after exposure to intermittent hypoxia (IH) for 24 hours. (B) Effect of U75302 and LY255283 on SRA mRNA expression in THP-1 cells after exposure to IH for 24 hours. (C) Effect of U75302 and LY255283 on ABCA1 mRNA expression in THP-1 cells after exposure to IH for 24 hours. (D) Effect U75302 and LY255283 on apolipoprotein A-I mRNA expression in THP-1 cells after exposure to IH for 24 hours. Data are expressed as a percentage of control RA (mean \pm SE; $n = 8$; * $P < 0.01$ vs. RA; [#] $P < 0.05$ vs. IH).

expression of ABCA1 (Figure 6D), suggesting that IH exposures do not seem to alter cholesterol efflux. Therefore, IH exposure promotes cholesterol influx and does not affect efflux, thereby contributing to intracellular cholesterol accumulation.

IH Promotes Atherosclerotic Lesion Formation in Apo $E^{-/-}$ Mice and Deletion of BLT1 Attenuates Atherosclerotic Lesion Formation in BLT1^{$-/-$}/ApoE^{$-/-$} Mice

To examine whether the selected IH profile would increase the production of LTB4 and promote atherosclerotic lesion formation in vivo, as recently shown by Jun and colleagues (10), Apo E^{-1} mice were fed with a high-cholesterol atherogenic western diet, and exposed to IH for 10 weeks. The plasma level of LTB4 production was assessed by an ELISA kit. En face analysis of exposed and control aorta for lesion formation was assessed with Sudan IV staining and by oil red O staining of sections by an investigator who was masked to the treatment group identity. The production of LTB4 in plasma was increased by IH exposure compared with the RA group (data not shown). In the RA-exposed group, there was limited evidence for the presence of lesion formation in either the aorta or the aortic root. In contrast, prominent lesion formations were apparent in both aorta and aortic roots in mice exposed to IH (Figures 7A and 7B), thereby confirming that IH enhances atherogenesis in vivo. To quantitatively assess atherosclerotic lesion formation further, the regions corresponding to atherosclerotic lesions in the aorta and the aortic root were captured and analyzed using image software. In the group exposed to IH, the areas of atherosclerotic lesion in aorta were significantly increased compared with RA controls ($n = 10$ per group) (Figure 7C). To determine further the effect of BLT1 on IH-associated atherosclerotic lesion formation, $BLT1^{-/-}$ and $ApoE^{-/-}$ double knock-out mice ($BLT1^{-/-}$ ApoE^{-/-} mice) were fed the same western diet and exposed to IH as described previously. In BLT1^{$-/-$}ApoE⁻ mice, the atherosclerotic lesions were prominently reduced

Figure 7. (A) En face analysis of aortas from apolipoprotein E (ApoE) $^{-/-}$ mice after RA or intermittent hypoxia (IH) exposure for 10 weeks. (B) Oil red O staining on aorta root from Apo $E^{-/-}$ mice after RA or IH exposure for 10 weeks. (C) Quantitative analysis of aortic lesion formation (mean \pm SE; n = 10; *P < 0.05 vs. ApoE $^{-/-}$ RA; $^{\#}P < 0.05$ vs. ApoE $^{-/-}$ IH).

compared with $ApoE^{-/-}$ mice (Figures 7A and 7B). In addition, the area of atherosclerotic lesion on aorta was significantly attenuated compared with the lesion formation in $ApoE^{-/-}$ mice $(n = 10 \text{ per group})$ (Figure 7C), thereby demonstrating that genetic deletion of BLT1 attenuates IH-associated atherosclerotic lesion formation in vivo, and that BLT1 plays a critical role in IH-associated atherogenesis.

DISCUSSION

In this study, we found that LTB4 production and the expression of 5-LO and LTA4H, the key enzymes involved in LTB4 biosynthesis, were induced by IH in a time-dependent fashion. In contrast, BLT1 and BLT2 were globally not affected. However, IH-induced 5-LO and LTA4H expression was reduced by a BLT1 antagonist and by a NF-kB blocker. IH was associated with increased transformation of monocytes to activated macrophages, and promoted increased cellular cholesterol accumulation and foam cell formation, all of which were markedly attenuated by BLT1 blockade. Furthermore, IH promoted in *vivo* atherosclerotic lesion formation in $\text{ApoE}^{-/-}$ mice, whereas genetic deletion of BLT1 abrogated this effect. Therefore, LTB4-BLT1 pathways play an important causative role in IH-induced atherogenesis and may provide a potential novel therapeutic target for OSA-associated atherosclerosis.

Current findings shed some light on the complex interplay between IH-induced production of LTB4 and its regulation. Our original assumption posited that IH may increase the production of LTB4 or lead to up-regulation of LTB4 receptors. Indeed, although robust LTB4 production and up-regulation of the two LTB4 biosynthetic enzymes emerged, the expression of LTB4 receptors, namely BLT1 and BLT2, remained unchanged. A possible explanation for the discrepant effects of IH on LTB4 and its receptors could be a down-regulatory effect on the receptors caused by increased LTB4 bioavailability or alternatively could represent post-translational modifications of LTB4 receptors linked to adaptation processes. Elevated circulating LTB4 concentrations have been reported in untreated patients with OSA, and continuous positive airway pressure treatment effectively reduced LTB4 plasma levels. Interestingly, LTB4 concentrations were correlated with luminal diameter, but not with intima–media thickness (33). However, our study is the first to provide direct evidence on the role of IH in inducing LTB4 production. These observations were somewhat anticipated because expression of 5-LO, 5-lipoxygenaseactivating protein (FLAP), and LTA4H can be modified by a variety of stressors, including ischemia and hypoxia (34–37). Thus, IH emerges as an important modulator of LTB4 biosynthesis, and can therefore induce local or systemic inflammatory processes, ultimately contributing to IH-associated atherogenesis. To elucidate potential signaling pathways involved in IH-induced regulation of 5-LO and LTA4H further, we explored DNA binding activity changes in HIF-1 α and NF- κ B, and pharmacologic manipulation of MAP kinases. Interestingly, IH induced DNA binding activity of NF-kB at 6 and 24 hours. In contrast, IH only induced a slight and transient increase in $HIF-1\alpha$ DNA binding activity at 6 hours. In addition, both MAPK and NF-kB blockers partially inhibited IH-induced 5-LO and LTA4H expression. Because activation of BLT1 and BLT2 may accelerate LTB4 synthesis (38), we explored whether either BLT1 or BLT2 may contribute to IH-induced 5-LO and LTA4H expression and increased LTB4 production. Treatment with a BLT1 antagonist effectively blocked both IH-induced 5-LO and LTA4H expression, whereas a BLT2 inhibitor was almost void of any effect, thereby suggesting that BLT1 is the candidate receptor involved in IH-regulated LTB4 production, at least in part by activating a LTB4–BLT1 amplification loop.

The critical role of LTB4 in the early stages of atherogenesis is now well established (31, 32, 39–43). LTB4 is a potent chemoattractant that facilitates recruitment of monocytes to the inflammatory sites (28). Recruitment of monocytes and leukocyte invasion of the arterial wall are critical steps in the development of atherogenesis (43–45). We have previously shown that LTB4 promotes atherosclerosis by chemoattracting monocytes, providing an amplification loop for monocyte chemotaxis via CCL2 production, and by converting monocytes to foam cells via enhanced expression of CD36 and fatty acid accumulation (32) . Current findings strongly support several critical roles for involvement of the LTB4–BLT1 pathway in IH-associated atherogenesis: (1) IH initiates monocyte recruitment as evidenced by increased MCP-1 and CCR2; (2) IH promotes macrophage differentiation, as shown by increased specific macrophage biomarkers in IH-exposed THP-1 cells; (3) IH up-regulated several key elements (e.g., CD36, SRA) that have well-established roles in cholesterol accumulation and foam cell formation, and may further play a critical role in the development of atherosclerosis; and (4) the atherogenic events were attenuated by a BLT1 antagonist (U75302). Although the multiple regulatory mechanisms underlying the expression of these individual genes are unknown, the coordinated regulation of all of these molecules through the activation of BLT1 by IH-induced LTB4 production clearly revealed a putative initiating and integrated step in atherogenesis.

In vivo studies have also consistently supported the view that the LTB4–BLT1 pathway plays a major role in the pathogenesis and progression of atherosclerosis (32, 40–42, 46, 47). ApoE knockout (Apo $E^{-/-}$) and low density lipoproteins (LDL) receptors knockout mice treated with BLT1 antagonists exhibited reduced lipid accumulation, monocyte infiltration, and smaller atheromata (48). Moreover, levels of adhesion molecules were reduced in mice treated with BLT1 antagonists (48). In rat basophilic cells, LTB4 promotes conversion of monocytes to foam cells through enhanced expression of scavenger receptors (CD36) and subsequent fatty acid accumulation (32). The cumulative evidence from aforementioned studies suggests that the effects of LTB4 in chemotaxis and foam cell formation are mediated by BLT1 and BLT2 receptors (32, 41–43, 48–50). In addition, expression of the 5-LO pathway (FLAP) is increased in atherosclerotic lesions at various stages of development in human aorta and coronary and carotid arteries (47, 51, 52). Furthermore, human genetic studies have shown that a promoter variant of 5-LO is associated with an increase in carotid intima–media thickness in healthy subjects, and certain FLAP haplotypes have been linked to an almost twofold increase in the risk of myocardial infarction or stroke (53). Although studies have explored the role of IH in OSA-associated atherosclerosis (10), its underlying mechanisms were not specifically sought. Here, we not only confirm that IH indeed promotes atherosclerosis, but further demonstrate that genetic deletion of BLT1 greatly attenuates IH-associated atherogenesis. It is possible that IH induces atherosclerosis via mechanisms other than BLT1, but the double knockout is partially protected because of the proatherogenic properties of BLT1, which are independent of IH. Therefore, we exposed $ApoE^ BLT1^{-/-}$ mice to either RA or IH and compared the lesion formation under those conditions. As illustrated in Figure 7C, the lesion formation in Apo $E^{-/-}BLT1^{-/-}$ mice exposed to IH was not significantly increased compared with the RA group. These results further support the notion that IH-induced atherogenesis is mediated through a LTB4–BLT1 dependent pathway. It is also well established that dyslipidemia plays an important role in the development of atherosclerosis. Therefore, IH-induced hyperlipidemia could also contribute to IH-associated atherogenesis (10). Of note, the biosynthesis of LTB4 is regulated by cholesterol (54)

and inhibition of leukotriene may reduce plasma triglyceride (TG) levels (55). However, it remains unclear whether IH-induced hyperlipidemia affects the biosynthesis of LTB4 and downstream atherogenesis. As a methodologic comment, we should point out that IH exposure only provides one aspect of OSA, because sleep fragmentation, episodic hypercapnia, and upper airway collapse are absent in this murine model. Also, the IH profile used here represents a standardization of exposures aiming to reflect severe disease, and as such may not completely represent the variable hypoxic profile characteristics seen in patients with OSA. Because new animal models are being developed to incorporate other elements of OSA (56, 52), assessment of the role played by the LTB4–BLT1 pathway in some or all of the constitutive components of OSA needs to be confirmed.

In summary, we have shown that LTB4 and its receptor BLT1 play a prominent role in the pathophysiologic mechanisms mediating atherogenesis associated with the typical pattern of episodic hypoxia associated with sleep-disordered breathing. These findings are consistent with the overarching concept whereby IH triggers the initiation of an inflammatory process, which is involved in atherogenesis. Thus, the LTB4–BLT1 pathway may provide a potential novel therapeutic target for OSA-associated atherosclerosis.

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