

Original Article

Increased expression and co-localization of ACE, angiotensin II AT₁ receptors and inducible nitric oxide synthase in atherosclerotic human coronary arteries

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Abstract: Using immunohistochemistry and quantitative *in vitro* autoradiography, the present study was undertaken to examine whether co-expression of pro-atherosclerotic factors, ACE, the AT₁ receptor, and iNOS, is increased in early and advanced atherosclerotic lesions of human coronary arteries. In normal coronary arteries, ACE and eNOS were strongly co-expressed in endothelial cells (ECs), whereas the AT₁ receptor was expressed in medial smooth muscle cells (SMCs). By contrast, iNOS was not expressed in ECs and SMCs. In early atherosclerotic lesions and atheromatous plaques, ACE, the AT₁ receptor and iNOS immunostaining were primarily co-localized in infiltrated macrophages and SMCs adjacent to macrophages. eNOS expression was lower in ECs than in normal arteries, and absent in accumulated macrophages and SMCs. In fibrosclerotic plaques, ACE, the AT₁ receptor, and iNOS immunostaining were still positive in macrophages as well as new microvessels within the plaques. Interestingly, SMCs in vasa vasorum of the adventitia in atheromatous and fibrosclerotic plaques were also strongly positive for AT₁ receptor and iNOS, while ECs of the vasa vasorum were positive for ACE and eNOS. The present study demonstrates that multiple pro-atherosclerotic factors ACE, AT₁ receptor and iNOS are co-localized almost exclusively in infiltrated macrophages and SMCs that have accumulated in or adjacent to early and advanced atherosclerotic plaques, while the anti-atherosclerotic enzyme eNOS is reduced in ECs. These data therefore suggest that increased formation of Ang II and iNOS in infiltrated macrophages and medial SMCs might well play important roles in the development and progression of human coronary atherosclerosis.

Keywords: Atherosclerosis, angiotensin converting enzyme, angiotensin II, human coronary artery, nitric oxide synthase, immunohistochemistry

Introduction

The development and progression of human coronary artery atherosclerosis are dependent on multiple factors, including genetics, life style, and the imbalance of pro-atherosclerotic and anti-atherosclerotic humoral influences. Many vasoactive hormones, growth factors and cytokines promote while others counteract the development and progression of coronary atherosclerosis. Angiotensin II (Ang II) is now well recognized as one of most important vasoactive pro-atherosclerotic factors [1-5]. Ang II, which is converted from Ang I by angiotensin converting

enzyme (ACE), can induce vascular injury due to its potent vasoconstrictor action [6] and ability of generating reactive oxygen species, and cause vascular hypertrophy by increasing smooth muscle cell (SMC) proliferation via acting on the AT₁ receptor [7]. While ACE occurs predominantly in the endothelium and adventitia of healthy human blood vessels [8-10], increased ACE expression has been observed in cellular structures beyond the endothelium and adventitia, such as the neointima developed following balloon injury in rat aorta [11]. Moreover, early atherosclerotic lesions commonly involve infiltration and/or migration of macro-

phages into the vessel wall, where increased expression of ACE has been reported in macrophages and vascular smooth muscle cells in atherosclerotic plaques of human coronary arteries [12-15]. These studies therefore suggest that local formation of Ang II is increased at the injured sites and therefore may play an important role in the development of human coronary atherosclerosis. However, it is not known whether local expression of AT₁ receptors is also increased in migrated macrophages and/or proliferative vascular SMCs in human atherosclerotic coronary arteries.

In contrast to Ang II, nitric oxide (NO) is generally a vasorelaxant and anti-atherosclerotic factor [16-18]. NO inhibits angiogenesis by preventing SMC proliferation [19]. Three isoforms of NO synthase (NOS) have been identified and cloned: neural NOS (nNOS), inducible NOS (iNOS) and endothelial NOS (eNOS) [20]. eNOS is constitutively present in endothelial cells of blood vessels and plays an important and beneficial role in maintaining normal endothelial function [21]. By contrast, iNOS is normally absent or expresses only at very low levels in different layers of blood vessels, but may become cytotoxic if excessively induced in infiltrated macrophages, proliferated SMCs, and endothelial cells by influences of vasoactive agents such as Ang II, cytokines, and/or under diseased states [18,22]. Although previous immunohistochemical studies have demonstrated that iNOS is expressed in macrophages of atherosclerotic coronary specimens taken from patients with unstable angina [23] and eNOS, though present, is reduced in endothelial cells of atherosclerotic lesions [24], co-localization and redistribution of different NOS isoforms during development and progression of human coronary atherosclerosis remain to be further investigated.

Similarly, while there is strong evidence that the renin-angiotensin system (RAS) and NO systems counteract each other in the regulation of vascular structures and tone physiologically as well as during the development and progression of human coronary atherosclerosis [25-29], to our knowledge there is still a lack of a comprehensive cellular mapping of the major components of the RAS (ACE and AT₁ receptor) and NO systems (eNOS and iNOS) in normal and early and advanced atherosclerotic human coronary arteries. In the present study, we hypothesized that

cellular expression of pro-atherosclerotic factors, ACE, AT₁ receptors and iNOS, is increased and co-localized in major cellular components of early atherosclerotic lesions and atheromatous plaques during the development and progression of human coronary disease. Increased co-expression or occurrence of pro-atherosclerotic factors ACE, which forms Ang II, the AT₁ receptor, which mediates actions of Ang II, and cytotoxic iNOS, while decreasing eNOS expression, in early and advanced atherosclerotic plaques may indeed promote human coronary atherosclerosis.

Methods

Tissue collection and American Heart Association (AHA) histological classification

The study was conducted on 88 coronary artery segments obtained at autopsy from 31 Caucasians patients who died from various causes including 24 males and 7 females with ages ranging from 21 to 83 years old (Table 1). 12 patients died from cardiovascular events including acute myocardial infarction, chronic heart failure and stroke, 11 from multiple organ injuries due to suicide or traffic accidents, and the remaining from lung or gastrointestinal diseases. The 88 coronary arterial samples consisted of 28 left main trunks, 22 left anterior descending arteries, 12 left circumflexes and 26 right coronary arteries according to American Heart Association classification. Collection of these autopsied coronary arteries for the current study was approved by the Human Ethics Committee of the Howard Florey Institute of Experimental Physiology and Medicine, The University of Melbourne.

All coronary arteries were obtained from autopsy within 24 hours after death, but most samples were collected between 8 to 12 hours. Samples were immediately frozen in liquid nitrogen and stored at -70°C before sectioning. Serial frozen sections, ~10 µm thick, were cut on a cryostat at -20°C for histological classification following haematoxylin and eosin staining, for immunohistochemical cellular localization of ACE, AT₁ receptors, iNOS or eNOS and cellular markers [12,13,30], and for autoradiographic quantification of ACE [8,9]. Classification of human coronary atherosclerosis was made according to American Heart Association Definitions with modifications [31-33]. Type I lesion

includes normal coronary artery with diffuse intimal thickening. Type II and/or type III are early atherosclerotic lesions, whereas type IV and/or type V shows advanced lesions. We further classified advanced lesions into atheromatous plaques and fibrosclerotic plaques. We had no type VI lesions designated complicated plaques.

Immunohistochemical localization

Antibodies: For cellular localization of ACE, a polyclonal antibody generated against a 25-amino-acid peptide located near the COOH terminus of human ACE was used (a gift from Professor Kunio Hiwada, Ehime University School of Medicine, Ehime, Japan). Its potency and specificity for human kidney ACE has been described previously [12,13,34]. The cellular localization of the AT₁ receptors was determined using a polyclonal antibody generated against the amino acid sequence corresponding to residues 15-24 of the human AT₁ receptor, Ac-QDDCPKAGRHC-NH₂ a hydrophilic portion from the amino-terminal extracellular domain coupled to an additional COOH-terminal cysteine (a gift from Professor Toshio Ogihara, Osaka University School of Medicine, Suita, Japan). Its specificity for the human AT₁ receptor has been reported previously [30,35,36].

To co-localize two isoforms of NO synthase with ACE and the AT₁ receptor in the same coronary section, we used a polyclonal anti-iNOS antibody (Transduction Laboratory Ltd.) and an anti-eNOS antibody (Transduction Laboratory Ltd.), respectively. We also identified cellular structures in each coronary artery using cell-specific antibodies with a monoclonal anti-smooth muscle antibody (1A4, DAKO Co Ltd.) for smooth muscle cells, a monoclonal anti-macrophage antibody (HAM56, DAKO Co Ltd.) for macrophages, and a monoclonal anti-von Willebrand factor antibody (DAKO Co Ltd.) for endothelial cells.

Immunohistochemistry: Immunohistochemical localization of ACE, AT₁ receptor, eNOS and iNOS was performed using an avidin-biotin complex method as described previously [12,13,36,37]. 10 µm-thick frozen sections were first fixed with acetone, endogenous peroxidase was blocked by 3% hydrogen peroxidase, and non-specific binding was blocked by incubation of sections in 10% normal goat serum, respectively. Sections were then incubated

with the primary antibodies for overnight at 4°C. After washes with phosphate buffered saline, sections were sequentially incubated with biotinylated secondary antibody and avidin-biotin complex (Vector Laboratories Inc.), and developed with 3-amino-9-ethylcarbazole (DAKO). Sections were then counter-stained with haematoxylin for histological identification. For negative controls, non-immune rabbit serum (DAKO) instead of the primary antibodies was used on adjacent sections.

In vitro autoradiographic localization and quantification of ACE: To further examine whether enhanced ACE protein expression in early and advanced atherosclerotic plaques was correlated with ACE activity in human coronary arteries, we determined ACE binding using quantitative *in vitro* autoradiography as described previously [8,9]. Briefly, ¹²⁵I-351A, a tyrosyl derivative of the ACE inhibitor, lisinopril (Merck Institute of Therapeutic Research) was used to label ACE, which binds to the active site of ACE and serves as an index of ACE activity. Four slide-mounted frozen sections of each coronary sample were pre-incubated in 10 mmol/L sodium phosphate buffer, pH 7.4, for 15 min, and then incubated in the same fresh buffer containing 0.2% bovine serum albumin (BSA) and ~0.3 mCi/mL of the radioligand, ¹²⁵I-351A, for 1 hour at 22°C. Non-specific binding was determined in the presence of 1 mmol/L EDTA, which inactivates ACE. After incubation, sections were washed with fresh buffer by 4 x 1 min, air-dried, and directly exposed to Agfa-Scopix CR3 X-ray film together with a set of radioactivity standards (Agfa-Gevaert) for 48 hours. ACE binding in the adventitia, early atherosclerotic lesions, atheromatous plaques, and fibrosclerotic plaques was quantitated by a computerized densitometry (MCID, Imaging Research Institute) compared with haematoxylin eosin-stained adjacent sections. Statistical analysis on ACE binding was performed using unpaired t test between normal and early or advanced atherosclerotic plaques. A value of p<0.05 was considered significant.

Results

Patient characteristics

Patients were classified into two broad groups, one died from known cardiovascular events and the other from non-cardiovascular causes. The former group of patients were generally older

(59.6 ± 4 vs. 42.6 ± 4 years old, $p < 0.05$), had higher body weight (71.6 ± 3.5 vs. 64 ± 3.5 kg, $p < 0.05$) and heart weight (464.5 ± 25 vs. 368.5 ± 11.3 g, $p < 0.01$) than the latter. However, body mass index (BMI) was similar between two groups.

Histological AHA classification

Of all coronary arterial samples, 33 were classified as normal coronary arteries with diffuse intimal thickening (see **Figure 1A**), 18 showed early atherosclerotic lesions (**Figure 2A**) and 37 had advanced lesions (**Figure 3A**). In advanced lesions, we identified 17 samples as atheromatous plaques and 10 as fibrous plaques. Normal coronary arteries with diffuse intimal thickening were identified primarily from patients died from non-cardiac events such as multiple organ injuries due to traffic accidents, suicide or other non-cardiovascular diseases, whereas advanced coronary lesions were identified predominantly in patients died from cardiovascular disorders ranging from myocardial infarction, hypertension and chronic heart failure.

Co-localization of ACE, AT₁ receptor, eNOS and iNOS by immunohistochemistry

Normal coronary arteries: **Figure 1** shows co-localization of ACE and eNOS, AT₁ receptor and iNOS in a representative normal coronary artery with diffuse intimal thickening. Intense ACE (**Figure 1B**) and eNOS immunostaining (**Figure 1C**) was co-localized predominantly in endothelial cells (**Figure 1D**), where only weak signals for AT₁ receptors (**Figure 1E**) and iNOS expression (**Figure 1F**) were present. By contrast, AT₁ receptors and iNOS were strongly expressed in vascular SMCs of thickened intima and medial SMCs. ACE and eNOS expression were absent in intimal and medial SMCs (**Figure 1B** and **Figure 1C**).

Atherosclerotic lesions

Early atherosclerotic lesions: In most early atherosclerotic lesions examined, ACE, iNOS and AT₁ receptors were co-localized in accumulated macrophages and SMCs in the intima, where eNOS expression is weak (data not shown).

Atheromatous plaques: **Figure 2** shows a typical representative atheromatous plaque, which

consists of a lipid core surrounded by a thickened intima and accumulated macrophages (**Figure 2D**). Macrophages adjacent to the lipid core were positive for ACE (**Figure 2B**), AT₁ receptors (**Figure 2E**) and iNOS (**Figure 2F**), but not for eNOS (**Figure 2C**). eNOS staining was only seen in endothelial cells and its expression level was much weaker than in normal coronary arteries (**Figure 1**). In addition to accumulated macrophages, AT₁ receptors were localized in SMCs around accumulated macrophages or adjacent to lipid cores in the intima, as well as in the media (**Figure 2E**).

Fibrosclerotic plaques: Infiltration of macrophages was still evident when atheromatous plaques advanced to fibrosclerotic plaques (**Figure 3D**). In addition, neovascularization occurs within the fibrous plaques. Co-localization of intensive ACE (**Figure 3B**), AT₁ receptor (**Figure 3E**) and iNOS immunostaining (**Figure 3F**) still occurred in infiltrated macrophages of fibrous plaques. ACE and eNOS immunostaining were weak in endothelial cells of coronary arteries, but appeared in endothelial cells of neovascularization within the plaque (**Figure 3B** and **Figure 3C**). Strong AT₁ receptor and iNOS expression was observed in medial SMCs (**Figure 3E** and **Figure 3F**).

Adventitia of advanced plaques: In the adventitia, ACE (**Figure 4B**) and eNOS (**Figure 4C**) were strongly expressed in, or co-localized with, endothelial cells (**Figure 4A**) of large and small vasa vasorum, whereas strong AT₁ receptor (**Figure 4E**) and iNOS expression (**Figure 4F**) was observed in the medial SMCs (**Figure 4D**) of vasa vasorum.

Quantification of ACE activity

Although ACE binding was not significantly different in the endothelia of various stages of coronary atherosclerosis (**Figure 5A**), ACE binding in early atherosclerotic lesions (44.3 ± 6.0 dpm/mm²) was significantly higher than that of normal coronary arteries (25.9 ± 2.1 dpm/mm²) ($p < 0.001$) or that in non-plaque samples ($p < 0.01$) (**Figure 5B**). ACE binding in atheromatous plaques (68.8 ± 5.5 dpm/mm²) was also significantly higher than that of early atherosclerotic lesion ($p < 0.001$), or that of fibrous plaque (33.3 ± 3.5 dpm/mm²) (**Figure 5B**) ($p < 0.001$). Examination of hematoxylin and eosin-stained adjacent sections confirmed that increased ACE

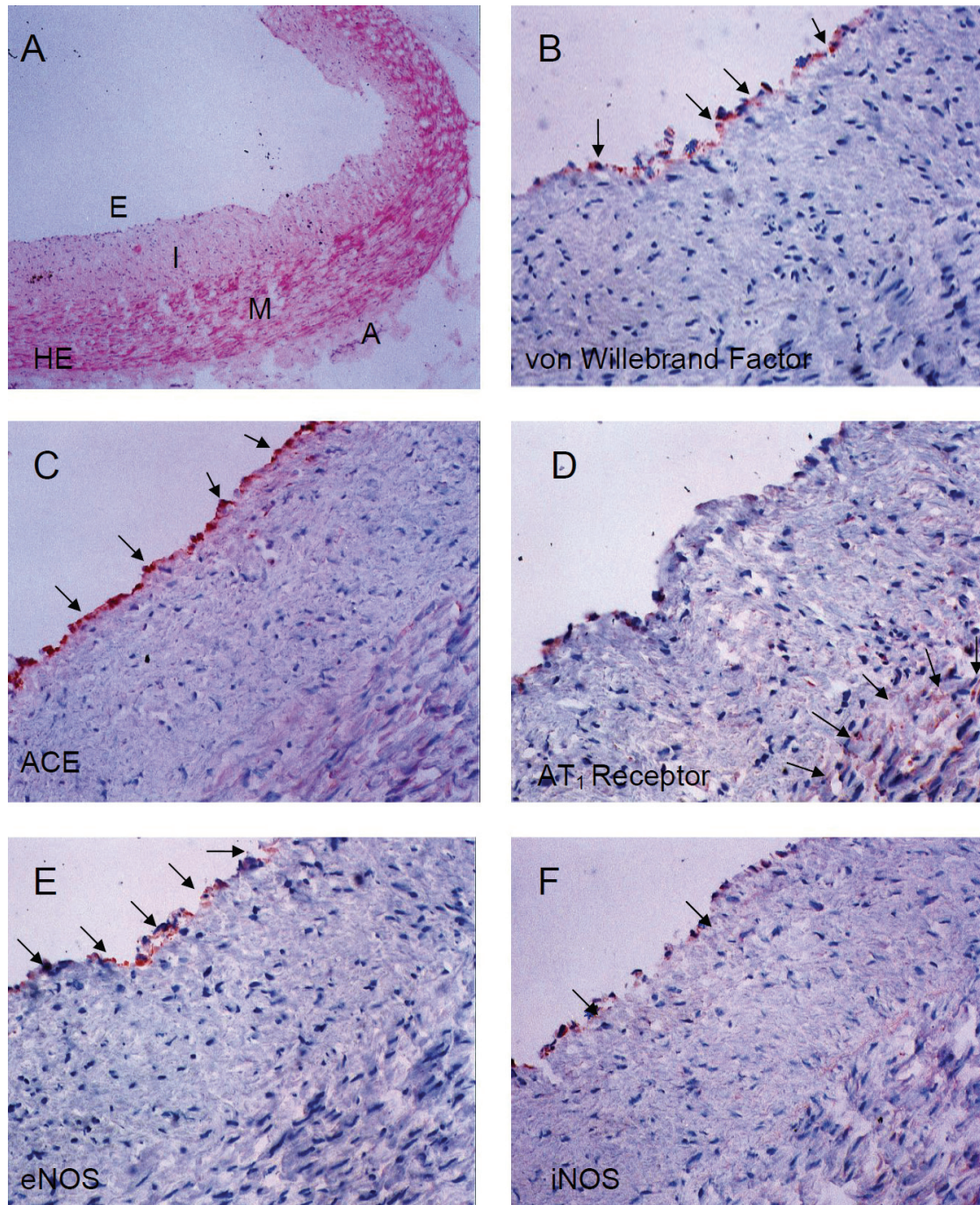


Figure 1. Cellular co-localization of ACE, AT₁ receptor, eNOS and iNOS expression in a representative normal human coronary artery with diffuse intimal thickening from a patient who died from a non-cardiovascular event. Panel A: a section stained with hematoxylin and eosin showing a mono-layer of endothelial cells (E), the thickened intima (I), medial smooth muscle cells (M), and adventitia (A). Panel B: a consecutive section of A stained with anti-von Willebrand factor antibody showing endothelial cells. Panel C: a consecutive section of B stained with anti-human ACE antibody showing ACE immunostaining co-localized with endothelial cells. Panel D: a consecutive section of C staining with anti-human AT₁ receptor antibody showing strong AT₁ receptor immunostaining mainly in vascular smooth muscle cells. Panel E: a consecutive section of D staining with anti-eNOS antibody showing eNOS immunostaining co-localized with endothelial cell marker and ACE. Panel F: a consecutive section of E stained with anti-iNOS antibody showing very weak iNOS immunostaining in endothelial cells and smooth muscle cells. Positive immunostaining is indicated by arrows. Magnification: Panel A, x 50; Panel B-F, x 200.

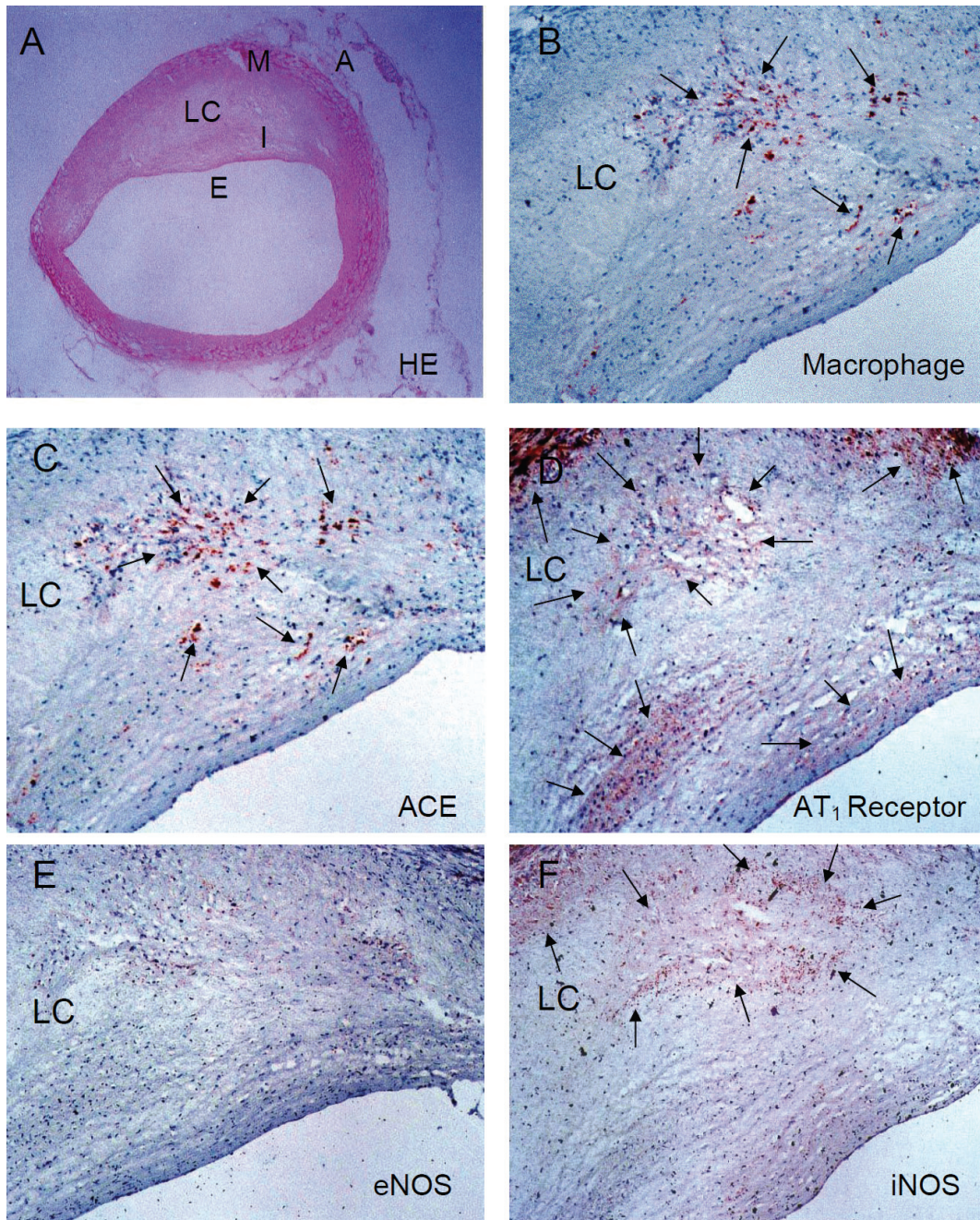


Figure 2. Cellular co-localization of ACE, AT₁ receptor, eNOS and iNOS expression in a representative atheromatous plaque of human coronary artery from a patient who died from a myocardial infarction. Panel A: a section stained with hematoxylin eosin showing an atheromatous plaque which has a lipid core (LC) surrounded by accumulated or infiltrated macrophages. Panel B: a consecutive section of A stained with anti-macrophage antibody (HAM56) showing intense immunostaining in the shoulder region of the plaque adjacent to the lipid core. Panel C: a consecutive section of B staining with anti-human ACE antibody showing ACE immunostaining co-localized with macrophage marker adjacent to the lipid core. Panel D: a consecutive section of C stained with anti-human AT₁ receptor showing strong AT₁ receptor immunostaining co-localized with thickening intima (I), infiltrated macrophages and in smooth muscle cells (M). Panel E: a consecutive section of D stained with anti-eNOS antibody showing reduced eNOS immunostaining in endothelial cells (E) and absent from atheromatous plaque. Panel F: a consecutive section of E stained with anti-iNOS antibody showing strong iNOS immunostaining co-localized with macrophage marker adjacent to the lipid core, and in smooth muscle cells. Positive staining is indicated by arrows. Magnification: Panel A, x 50; B-F, x 200.

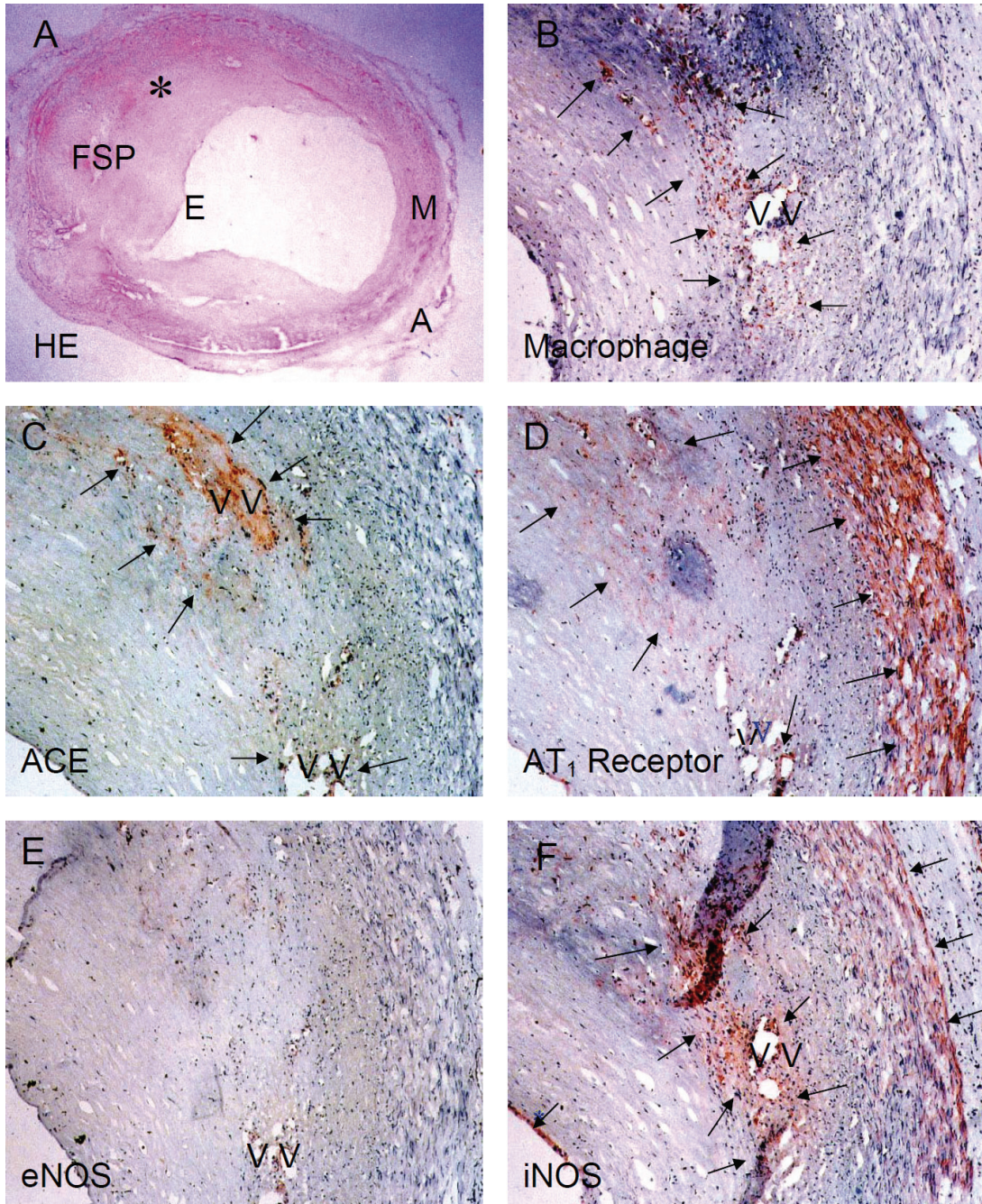


Figure 3. Cellular co-localization of ACE, AT₁ receptor, eNOS and iNOS expression in a representative fibrosclerotic plaque of human coronary artery from a patient who died from myocardial infarction. Panel A: a section stained with hematoxylin and eosin showing a fibrosclerotic plaque (FSP), which consists of dense connective tissues in atherosclerotic lesions and neovascularization. Vasa vasorum (VV) is occasionally found within the plaque. Panel B: a consecutive section of A showing macrophage marker immunostaining surrounding a vasa vasorum within the plaque. Panel C: a consecutive section of B showing strong ACE immunostaining co-localized with macrophages and neovascularization, but reduced in endothelial cells. Panel D: a consecutive section of C showing strong AT₁ receptor immunostaining localized in macrophages, thickening intima and medial smooth muscle cells. Panel E: a consecutive section of D showing reduced eNOS immunostaining in endothelial cells as well as neovascularization. Panel F: a consecutive section of E showing strong iNOS immunostaining co-localized with accumulated macrophages surrounding vasa vasorum and medial smooth muscle cells. Positive staining is indicated by arrows in macrophage-rich lesions around vasa vasorum and medial smooth muscle cells. Magnification: Panel A, x 50; Panel B-F, x 200.

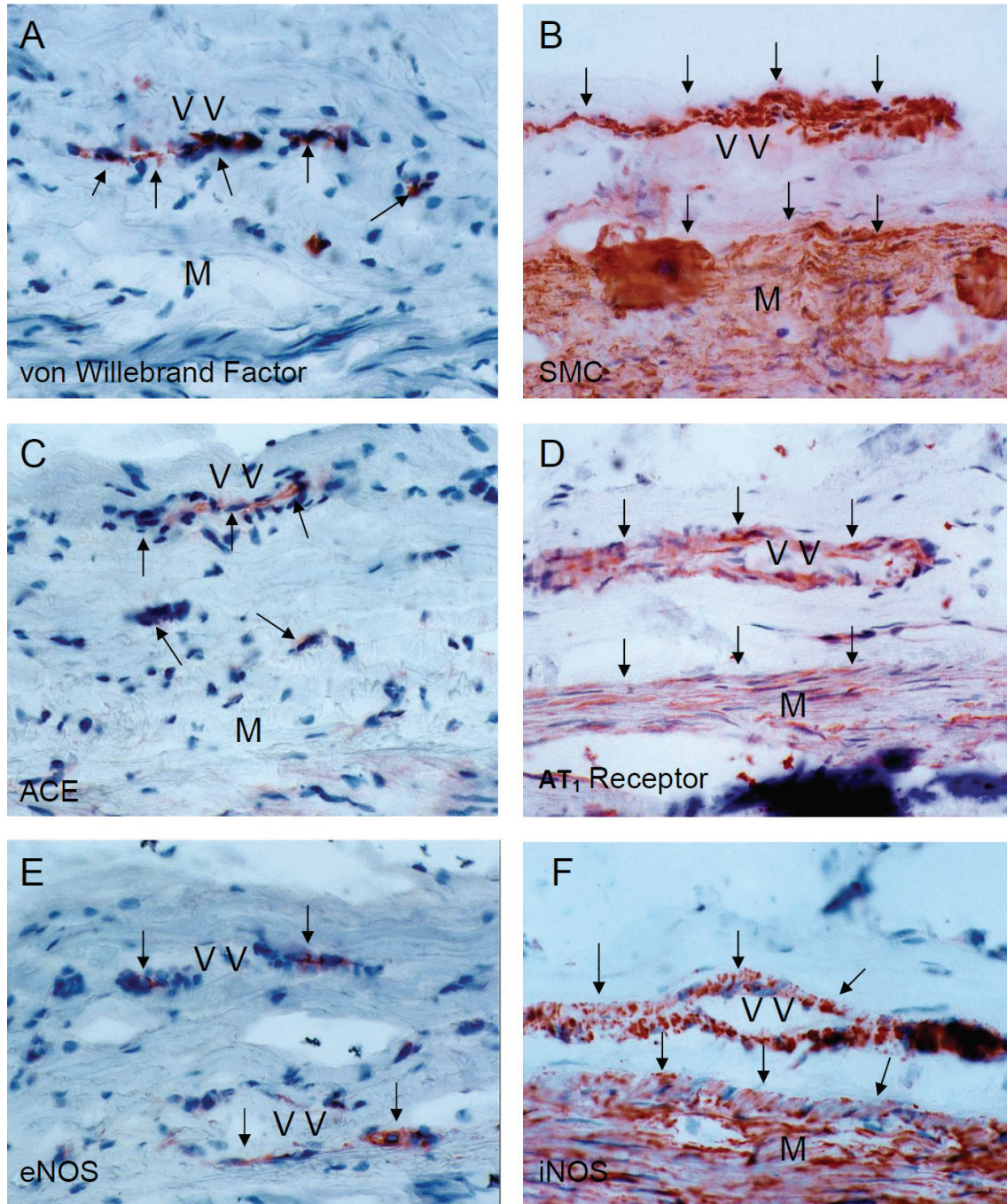


Figure 4. Cellular co-localization of ACE, AT₁ receptor, eNOS and iNOS expression in the adventitia of a representative advanced atherosclerotic plaque of human coronary artery from a patient who died from myocardial infarction. Panel A: a section stained with anti-von Willebrand factor antibody showing that the endothelium of vasa vasorum (VV) in the adventitia (A) of this advanced plaque were positive for the endothelial marker. Panel B: a consecutive section of A stained with the smooth muscle cell marker, HAM56. Panel C: a consecutive section of B showing strong ACE immunostaining co-localized with the endothelial marker in vasa vasorum. Panel D: a consecutive section of C showing strong AT₁ receptor immunostaining in smooth muscle cells of large vasa vasorum and medial smooth muscle cells. Panel E: a consecutive section of D showing eNOS immunostaining localized in endothelial cells of vasa vasorum. Panel F: a consecutive section of E showing strong iNOS immunostaining co-localized with the AT₁ receptor in smooth muscle cells of large vasa vasorum and medial smooth muscle cells. Positive staining is indicated by arrows in endothelial cells of vasa vasorum and medial smooth muscle cells. Magnification: Panels A-F, x 200.

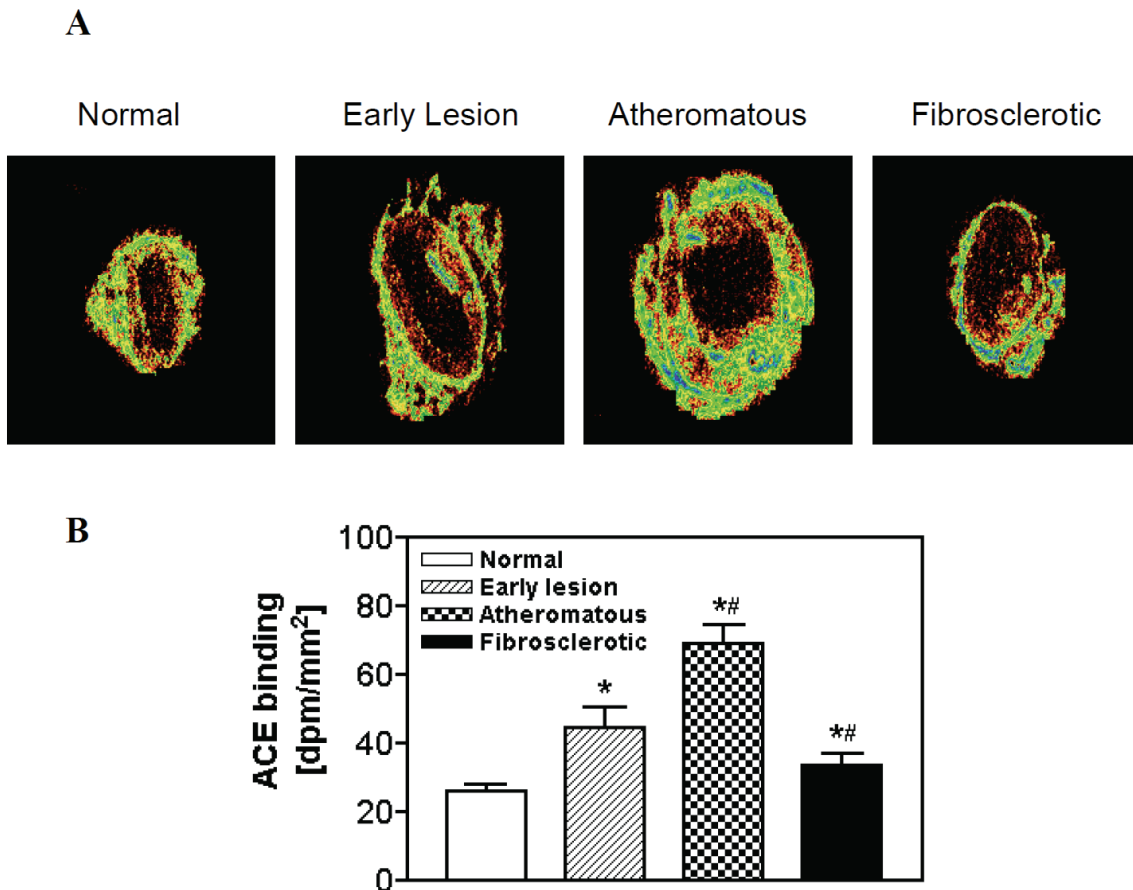


Figure 5. *In vitro* autoradiography showing increased ACE binding in early atherosclerotic lesions and advanced plaques of human coronary arteries (A) with the quantitated data shown in Panel B. Yellow shows the highest level of ACE binding, while green and black the moderate and background levels, respectively. *, $p < 0.05$ vs. normal; #, $p < 0.05$, atheromatous vs. early lesions; or atheromatous vs. fibrosclerotic.

binding at these coronary arteries was closely associated with accumulated macrophages, adventitial vasa vasorum, and endothelial cells of neovascularization in advanced plaques.

Discussion

The present study demonstrates that during the development and progression of human coronary atherosclerosis, cellular expression of pro-atherosclerotic factors, ACE, AT₁ receptor and iNOS, is markedly increased at cellular sites with early and advanced atherosclerotic lesions or plaques. This enhanced expression of ACE, AT₁ receptor and iNOS at the protein level occurs almost exclusively in accumulated or infiltrated macrophages and vascular smooth muscle cells within or adjacent to the shoulder of

atherosclerotic lesions or lipid cores. By contrast, eNOS expression is markedly reduced in endothelial cells and no eNOS expression occurs in accumulated macrophages and smooth muscle cells within and around the plaques. To our knowledge our study represents the first comprehensive immunohistochemical investigation showing increased expression and co-localization of ACE, AT₁ receptor and iNOS in the same early and advanced atherosclerotic lesions during the development and progression of human coronary artery atherosclerosis.

It should be acknowledged that the current study has strengths as well as limitations. Because all normal and atherosclerotic coronary artery segments were obtained from autopsied patients several hours after death, only immu-

nohistochemistry was performed to co-localize these vasoactive factors with the exception for ACE activity, which can be determined by quantitative *in vitro* autoradiography. For the same reason, we were also unable to measure Ang II or NO production in these autopsied coronary arteries *ex vivo*. However, collection of normal and atherosclerotic coronary arteries from patients died from known cardiovascular events and those from other non-cardiac causes allowed us to classify all coronary tissue sections from these patients into four different groups according to the AHA classification before immunohistochemistry was performed. This approach made it possible to co-localize ACE, AT₁ receptor and two isoforms of NOS, eNOS and iNOS, together with cellular markers including antibodies against von Willenbrand factor (endothelial cells), 1A4 (smooth muscle cells) and HAM56 (macrophages) in consecutive sections of the same coronary artery.

The results of the present study on ACE and AT₁ receptors are consistent with previous studies in experimentally injured vessels of animals or in early and advanced atherosclerotic plaques of human coronary arteries [11-13,38,39]. Rakugi et al have demonstrated that increased expression of ACE occurs in the developing neointima two weeks following induction of aorta balloon injury in rats [11]. In monkeys with experimental atherosclerosis induced by high cholesterol diet, both ACE and AT₁ receptor binding significantly increase in atherosclerotic vessels [38]. Increased AT₁ receptor expression has also been reported in a rabbit model of hypercholesterolemic atherosclerosis [39]. In human studies, we [12,13,40] and other investigators [14,15] further showed increased ACE expression in accumulated macrophages of early atherosclerotic lesions and atheromatous plaque, as well as at sites of injury following percutaneous transluminal coronary angioplasty in humans [13]. However, AT₁ receptor was not co-localized with ACE in accumulated macrophages and smooth muscle cells within and around early and advanced atherosclerotic plaques in afore-mentioned studies. Because ACE is commonly expressed in the endothelium and adventitia but not in medial smooth muscle cells of normal coronary artery, increased ACE expression in accumulated macrophages and smooth muscle cells adjacent to atherosclerotic plaques strongly suggests an increase in local formation of Ang II at these cellular sites. In-

deed, in a study in patients with acute coronary syndrome ACE activity increased 4-fold in coronary artery specimens [41]. Using quantitative *in vitro* autoradiography the present study was able to demonstrate increased ACE binding as a marker of ACE activity in early and advanced atherosclerotic plaques, where ACE co-localized with macrophages and smooth muscle cells (**Figure 5**). These data are consistent with our previous studies in which Ang II immunostaining was identified in macrophages and smooth muscle cells of human coronary hypercellular lesions and atheromatous plaques [40], thus providing further support to the concept that Ang II is produced locally in infiltrated macrophages and smooth muscle cells during the development and progression of human coronary atherosclerosis.

While AT₁ receptor expression occurs in medial smooth muscle cells as expected, it is highly significant for the present study that AT₁ receptor expression is increased and co-localized with ACE and iNOS (see below) in accumulated macrophages and smooth muscle cells within and around atherosclerotic lesions. This enhanced expression was observed in most coronary sections classified as early atherosclerotic lesions, atheromatous plaques and fibrosclerotic plaques examined. It is also of particular interest that AT₁ receptor immunostaining occurs in accumulated macrophages and smooth muscle cells adjacent to lipid cores in addition to medial smooth muscle cells (**Figure 2E**). These findings have not been reported previously in human coronary atherosclerotic lesions, though AT₁ receptors mRNA has been found in human macrophages [30].

The significance of co-localization of ACE, Ang II and AT₁ receptors at these atherosclerotic lesions is still poorly understood, but may imply important implications, which warrant further investigations. It has been suggested that AT₁ receptors may mediate vasoconstriction in the contractile phenotype [6] and cell proliferation in the synthetic phenotype of vascular smooth muscle cells [7]. Although we were not able to differentiate whether smooth muscle cells adjacent to the lipid core or to the shoulder of atherosclerotic lesions are of the synthetic phenotype, AT₁ receptors may play dual roles in mediating both cell contraction and proliferation at these sites. Thus, it is conceivable that Ang II-induced vascular smooth muscle cell contrac-

tion and proliferation, together with Ang II-induced iNOS expression (see discussion below), may be one of most important contributors to the development of human coronary atherosclerosis. Additionally, expression of ACE and AT₁ receptors in inflammatory macrophages accumulated at early and advanced atherosclerotic lesions also implies an unspecified but important role for interactions between Ang II and macrophages. Ang II has been shown to increase macrophage-induced oxidation of low-density lipoprotein or peroxide production via the AT₁ receptor [42] or via a lipoxygenase-dependent pathway [43]. Furthermore, stimulation of peroxide production or induction of iNOS expression (see discussion below) in accumulated macrophages by Ang II via the AT₁ receptor has also been reported. In keeping with this context, it is not surprising to note that Ang II promotes atherosclerotic lesions and aneurysms in apolipoprotein E-deficiency mice [1,2]. Taken together, Ang II and macrophages may interact to promote atherosclerosis via multiple mechanisms related to cell contraction and proliferation, and synthesis and/or release of cytokines, elastase, collagenase and free radicals.

Other equally important findings in the present study are that while eNOS expression was reduced (**Figure 2C** and **Figure 3C**) in endothelial cells comparing with normal coronary artery (**Figure 1C**) and was absent at cellular sites with early atherosclerotic lesions and advanced atherosclerotic plaques, enhanced iNOS expression was localized in accumulated macrophages and smooth muscle cells of the shoulder lesions adjacent to the lipid core (**Figure 2F** and **Figure 3F**). This cellular distribution of iNOS expression coincides with those of enhanced expression of ACE (**Figure 2B** and **Figure 3B**) and AT₁ receptors (**Figure 2E** and **Figure 3E**) in early and advanced atherosclerotic lesions. Reduced eNOS expression and production has been reported previously in atherosclerotic human carotid arteries compared with normal internal mammary arteries [24]. However, it is not known whether this applies to human coronary atherosclerotic arteries. In this context, our results provide morphological evidence that eNOS expression and production may be reduced during the progression of human coronary atherosclerosis. Indeed, accelerated atherosclerosis and ischemic heart disease has been reported in apolipoprotein E/eNOS double knockout mice [44].

As for iNOS expression, our results are consistent with previous studies on iNOS expression during vascular injury and human coronary atherosclerosis. For example, induction of vascular iNOS expression has been reported in animal models of vascular injury, such as those induced by periarterial collar [45] or by endothelial denudation [46]. In human coronary disease, iNOS expression was increased in macrophages as well as smooth muscle cells of transplanted human coronary arteries [17], and in human coronary atherosclerotic plaques of patients [23,47,48]. However, it is not clear whether induction of iNOS is associated with, or accompanied by, enhanced ACE and AT₁ receptor expression and reduced eNOS expression in these vascular lesions. The roles of NO in the development of human coronary atherosclerosis are currently under intensive investigations. It has been suggested that NO, produced by eNOS and iNOS, may serve dual roles in the development and progression of human coronary atherosclerosis [16]. Endothelial NOS is located predominantly in endothelial cells and produces small amounts of NO in response to physiological shear stress from raised arterial pressure or to stimulation by kinins via the B₂ receptor [18]. NO produced by eNOS is beneficial, anti-atherosclerotic, and important in maintaining normal endothelial function as it induces vasorelaxation, inhibits leukocyte adhesion and platelet aggregation, decreases expression of adhesion molecules and chemotactic factors, and inhibits smooth muscle cell proliferation [16,18]. By contrast, iNOS induced locally in macrophages and smooth muscle cells by cytokines or during the inflammatory processes including Ang II can produce large quantity of NO at sites of its expression; this NO can be cytotoxic and pro-atherosclerotic [18,49]. The pro-atherosclerotic properties of iNOS as a result of producing large amount of NO include: a) interaction with the free radical superoxide to form peroxynitrate, which is known to damage cellular proteins; b) induction of vascular hyperreactivity; and c) promotion of leukocyte adhesion and platelet aggregation [16-18]. Thus, increased expression of pro-atherosclerotic iNOS and reduced expression of anti-atherosclerotic eNOS in early and advanced atherosclerotic lesions or plaques as shown in the present study may play an important role in human coronary atherosclerosis.

In summary, the present study demonstrates for

the first time that expression of ACE, AT₁ receptor and iNOS at the protein level are co-localized almost exclusively in infiltrated macrophages and vascular smooth muscle cells adjacent to the lipid core of early and advanced atherosclerotic lesions or plaques, while eNOS expression is reduced in endothelial cells of atherosclerotic human coronary artery. These data therefore suggest that over-expression or formation of Ang II and iNOS in infiltrated macrophages and migrated smooth muscle cells plays an important role in the development and progression of human coronary atherosclerosis.

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