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### Research Article

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**Heparin cofactor II (HCII) specifically inhibits thrombin action at sites of injured arterial wall, and patients with HCII deficiency exhibit advanced atherosclerosis. However, the *in vivo* effects and the molecular mechanism underlying the action of HCII during vascular remodeling remain elusive. To clarify the role of HCII in vascular remodeling, we generated *HCII*-deficient mice by gene targeting. In contrast to a previous report, *HCII*<sup>-/-</sup> mice were embryonically lethal. In *HCII*<sup>+/-</sup> mice, prominent intimal hyperplasia with increased cellular proliferation was observed after tube cuff and wire vascular injury. The number of protease-activated receptor-1-positive (PAR-1-positive) cells was increased in the thickened vascular wall of *HCII*<sup>+/-</sup> mice, suggesting enhanced thrombin action in this region. Cuff injury also increased the expression levels of inflammatory cytokines and chemokines in the vascular wall of *HCII*<sup>+/-</sup> mice. The intimal hyperplasia in *HCII*<sup>+/-</sup> mice with vascular injury was abrogated by human HCII supplementation. Furthermore, HCII deficiency caused acceleration of aortic plaque formation with increased PAR-1 expression and oxidative stress in apoE-KO mice. These results demonstrate that HCII protects against thrombin-induced remodeling of an injured vascular wall by inhibiting thrombin action and suggest that HCII is potentially therapeutic against atherosclerosis without causing coagulatory disturbance.**

## Introduction

Thrombin not only acts as a key enzyme in the formation of fibrin clots during the blood coagulation process but also activates platelets, vascular endothelial cells, VSMCs, macrophages, and fibroblasts to enhance procoagulation (1), chemoattraction (2), mitogenesis (3), and proliferation (4) of these cells. Actions of thrombin in these cells are mediated through proteolytic processing of specific cell-surface receptors known as protease-activated receptors (PARs) (5). Among the 4 members of the family of PARs, PAR-1 is known as the most important thrombin receptor. Although approximately one-half of PAR-1-deficient mice are embryonically lethal (6), mice that survive manifest diminished neointimal hyperplasia after arterial injury compared with WT mice (7). In humans, PAR-1 is widely expressed in macrophages, VSMCs, and intimal cells of atherosclerotic lesions (8), and PAR-1 activation has been shown to play an important role in the development of cardiovascular diseases, including arterial thrombosis (9), stroke, and ischemic heart disease (10, 11). In addition, it has

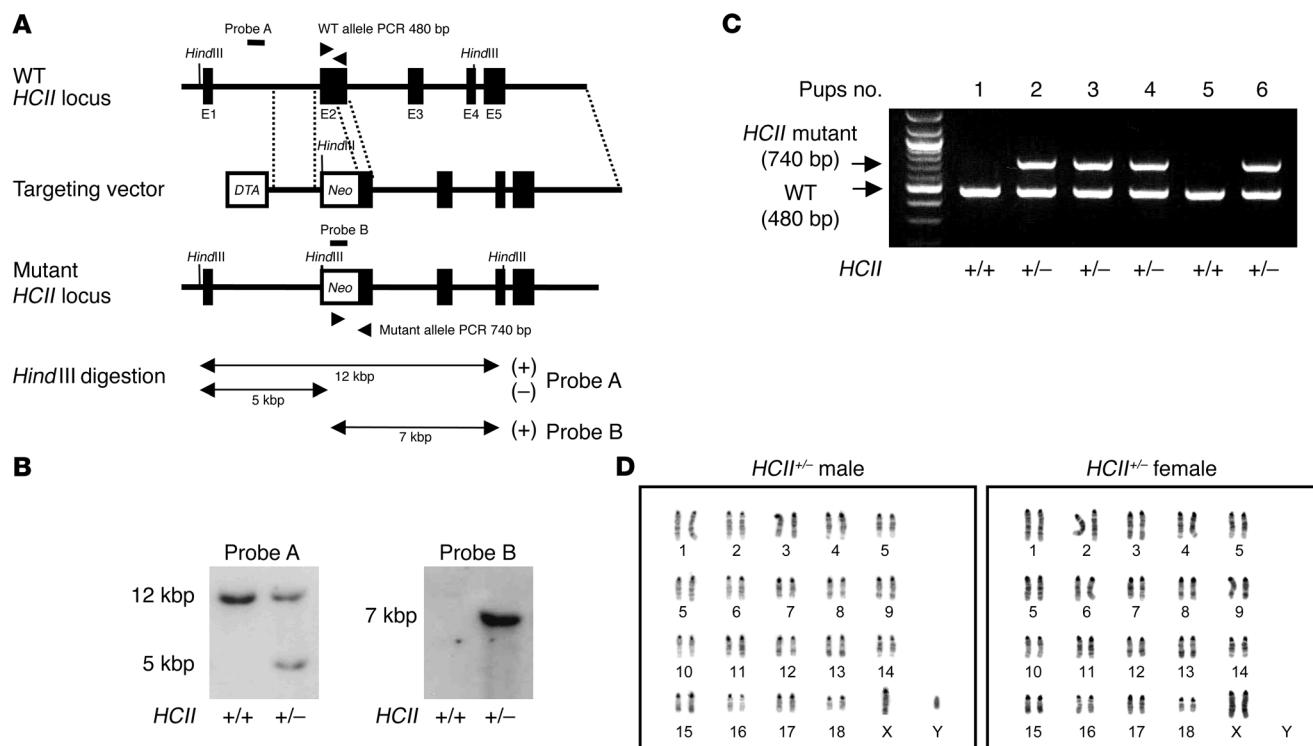
been reported that thrombogenicity-regulating factors such as tissue factor (TF), thrombomodulin (TM), hirudin, vitronectin, and plasminogen activator inhibitor-1 (PAI-1) regulate VSMC proliferation or migration leading to vascular remodeling (12–15). Those factors directly or indirectly contribute to modulation of thrombin action. Thrombin action-accelerating factors, including TF, vitronectin, and PAI-1, promote vascular remodeling, whereas thrombin action-attenuating factors such as TM and hirudin suppress vascular remodeling (12–15).

Enhanced thrombin action at sites of injured vascular endothelium is efficiently inhibited by antithrombin (AT) and heparin cofactor II (HCII). AT exhibits its activity by binding to heparan sulfate, and anticoagulatory active heparan sulfate proteoglycans are concentrated immediately beneath the aortic endothelium, with only a small amount present on the luminal surfaces of endothelial cells (16). HCII is also a serine protease inhibitor (serpin) with a molecular weight of 65.6 kDa. HCII is synthesized by hepatocytes and secreted into the blood stream at a concentration of about 1.0  $\mu\text{mol/l}$  (17) and is distributed in the intima and media of atherosclerotic lesions as well as walls of normal arteries. HCII inactivates thrombin without affecting other proteases involved in the blood coagulation cascade. Inactivation of thrombin action by HCII is enhanced by more than 1,000-fold after binding to dermatan sulfate (DS) (17, 18), which is synthesized and secreted by smooth muscle cells and fibroblasts and is deposited in the matrix

**Nonstandard abbreviations used:** AT, antithrombin; DHE, dihydroethidium; DS, dermatan sulfate; Egr-1, early growth response protein-1; HCII, heparin cofactor II; KLF5, Krüppel-like zinc finger transcription factor 5; MCP-1, monocyte chemoattractant protein-1; 8OHdG, 8-hydroxy-2'-deoxyguanosine; PCNA, proliferating cell nuclear antigen.

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**Figure 1**

Targeted disruption of murine *HCII* gene, genotyping, and karyotypes of *HCII*-mutant mice. **(A)** Genomic locus, targeting vector, and predicted targeting locus are illustrated. External (probe A) and internal (probe B) probes were used for Southern blot analysis. Two sets of PCR primers for detecting the 480-bp WT allele and 740-bp mutant allele were employed for genotyping. **(B)** Southern blot analysis of murine genomic DNA. A 5-kbp *HindIII* fragment denotes the homologous recombinant allele in probe A as an external probe, and a single 7-kbp *HindIII* fragment denotes the homologous recombinant allele without random integration in probe B as an internal probe. **(C)** Genotyping PCR. The mutant allele yielded a 740-bp band, and the WT allele yielded a 480-bp band. **(D)** Karyotype analysis of *HCII*<sup>+/-</sup> male and female mice. Chromosomes of splenic lymphocytes in both male and female *HCII* mutant mice showed normal karyotypes.

of vascular intima and media (19–21). Thus, AT inhibits thrombin action and elicits an anticoagulatory effect at the surface of the vascular wall, whereas HCII can most efficiently act in the intima and media of the vascular wall to protect them from the actions of thrombin on vascular remodeling.

We previously reported a 66-year-old woman with type I congenital HCII deficiency manifesting multiple atherosclerotic lesions, including multiple coronary vessel disease, bilateral carotid stenosis, right renal artery stenosis, and abdominal aortic aneurysm (22). The advanced atherosclerotic lesions in this patient as well as in another family member prompted us to speculate that HCII plays a protective role against the formation of atherosclerotic lesions in elderly individuals who have cardiovascular risk factors with injured vascular walls. We therefore conducted 2 clinical studies to determine the relationship between plasma HCII activity and development of atherosclerotic lesions. Studies on in-stent restenosis after percutaneous coronary intervention (23) and on carotid atherosclerosis (24) provided clinical evidence that HCII plays a protective role against the progression of atherosclerosis in the elderly with atherosclerotic risk factors. In 2002, He et al. generated *HCII* gene-KO mice and showed that the KO mice manifest normal growth and enhanced thrombogenicity after photochemical damage to the carotid endothelium (25). However, the mechanisms of HCII action as well as the importance of HCII in preventing the formation of atherosclerotic lesions are as yet

unclear. To address these issues, we generated *HCII*-deficient mice by gene targeting and analyzed vascular remodeling in WT and *HCII*-mutant mice using a vascular injury model and a hyperlipidemia-induced atherosclerosis model.

## Results

**Generation and characterization of *HCII*-KO mice.** In order to disrupt the *HCII* gene function, we deleted a 120-bp fragment of exon 2 including the ATG site in the *HCII* gene and inserted a *phosphoglycerate kinase–neomycin* cassette with polyA termination. The targeting vector (Figure 1A) was introduced into TT2 ES cells by electroporation. Among 354 ES clones, we found 3 clones with mutated *HCII* exon 2 by homologous recombination, which was confirmed by Southern blot analysis with external and internal probes (Figure 1B). Two of the 3 clones did not have any random integration (Figure 1B). Blastocyst injections of 2 independent ES clones yielded chimeric mice. The chimeric mice derived from 1 ES clone transmitted the mutant allele into the germ line. *HCII*<sup>+/-</sup> mice were produced by mating the chimeric mice with female C57BL/6J mice. Thereafter, *HCII*<sup>+/-</sup> mice were backcrossed for 10 generations with the C57BL/6J strain. By mating with male and female *HCII*<sup>+/-</sup> mice, we could not obtain *HCII*<sup>-/-</sup> mice due to early embryonic lethality at E6.5 to E8.5 (Table 1 and Figure 1C); however, *HCII*<sup>+/-</sup> mice were fertile and exhibited normal survival and growth (data not shown). G band staining demonstrated that male *HCII*<sup>+/-</sup> mice and female

**Table 1**

Genotypes of embryos and newborns derived through timed mating of *HCII*-mutant heterozygote

Days of gestation	Genotype of <i>HCII</i> gene			Total
	+/+	+/-	-/-	
Newborn	88	177	0	265
16.5	23	49	0	72
14.5	12	20	0	32
12.5	10	22	0	32
10.5	9	21	0	30
8.5	6	10	0	16

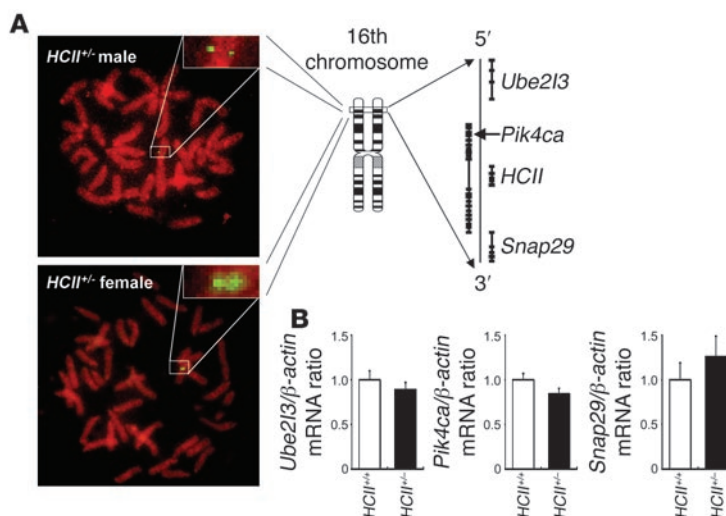
*HCII*<sup>-/-</sup> mice have 38,XY karyotype and 38,XX karyotype, respectively, with no obvious abnormal chromosomes (Figure 1D). FISH analysis using a *phosphoglycerate kinase*–*neomycin* cassette probe showed the expected position of the *phosphoglycerate kinase*–*neomycin* cassette at the sixteenth murine chromosome, and no random integration of the cassette was observed in male and female *HCII*<sup>-/-</sup> mice (Figure 2A). In addition, the expression levels of neighboring genes around the *HCII* genome, such as *Ube2l3*, *Pik4ca*, and *Snap29*, were the same in *HCII*<sup>+/+</sup> and *HCII*<sup>-/-</sup> mice (Figure 2B). Conventional RT-PCR analysis of the full-length *HCII* gene demonstrated no abnormal bands in either *HCII*<sup>+/+</sup> or *HCII*<sup>-/-</sup> mice (Figure 3A), and direct sequence analysis of each PCR product revealed no mutations in either *HCII*<sup>+/+</sup> or *HCII*<sup>-/-</sup> mice (data not shown). These results confirmed the absence of truncated *HCII* transcripts and noninterference for mRNA levels of neighboring genes around the *HCII* genome after *HCII* targeting vector introduction. *HCII*<sup>-/-</sup> mice showed half of the level of *HCII* protein content and mRNA in the liver and about 60% of the plasma *HCII* activity of *HCII*<sup>+/+</sup> mice (Figure 3, B–D). To clarify the difference between embryonic lethality of *HCII* mutant mice in the study by He et al. (25) (their heterozygous *HCII* mutant mice are designated “*HCII*<sup>-/-</sup> St. Louis” herein) and that of our *HCII* mutant mice (our heterozygous *HCII* mutant mice are designated “*HCII*<sup>-/-</sup> Tokushima” herein), we interbred *HCII*<sup>-/-</sup> Tokushima with *HCII*<sup>-/-</sup> St. Louis mice. *HCII* homozygous mutant pups having different *HCII* mutant alleles (Tokushima/St. Louis) were born close to the expected mendelian frequency (Table 2). Although the *HCII* (Tokushima/St. Louis) homozygous mutant mice showed undetectable plasma *HCII* antigen levels, the mutant mice exhibited 14.2% ± 2% of plasma *HCII* activities (Table 2). We conducted all the following examinations using *HCII*<sup>-/-</sup> Tokushima mice.

**Enhanced platelet aggregation in *HCII*<sup>-/-</sup> mice.** Hemostatic examinations, including examinations of plasma fibrinogen level, prothrombin time, and plasma AT activity, revealed no significant differences between *HCII*<sup>+/+</sup> and *HCII*<sup>-/-</sup> mice (Figure 4A). Next, we evaluated ADP-induced whole blood aggregation using a screen filtration pressure method in these mice. Platelet aggregatory threshold index values in *HCII*<sup>-/-</sup> mice were decreased markedly compared with those in *HCII*<sup>+/+</sup> mice (Figure 4B). These results indicate that enhanced platelet aggregation occurred in *HCII*<sup>-/-</sup> mice and suggest that *HCII* has a suppressive effect on platelet aggregability in vivo.

**Arterial cuff injury causes enhanced intimal and adventitial hyperplasia in *HCII*<sup>-/-</sup> mice.** No morphometrical differences were observed between uninjured femoral arteries in WT and *HCII*<sup>-/-</sup> mice (Figure 5, A and B). In an attempt to determine whether *HCII*<sup>-/-</sup> mice exhibit a difference in structure of the vascular wall after vascular injury, we compared structural changes in the walls of arteries after polyethylene tube cuff placement for 4 weeks. Cuff-injured femoral arteries in *HCII*<sup>-/-</sup> mice revealed a prominent increase in intimal and adventitial areas but not in the medial area compared with those in *HCII*<sup>+/+</sup> mice (Figure 5, A and B). As a consequence, the intima/media ratio and adventitia/media ratio were also significantly increased in *HCII*<sup>-/-</sup> compared with *HCII*<sup>+/+</sup> mice (Figure 5, A and B).

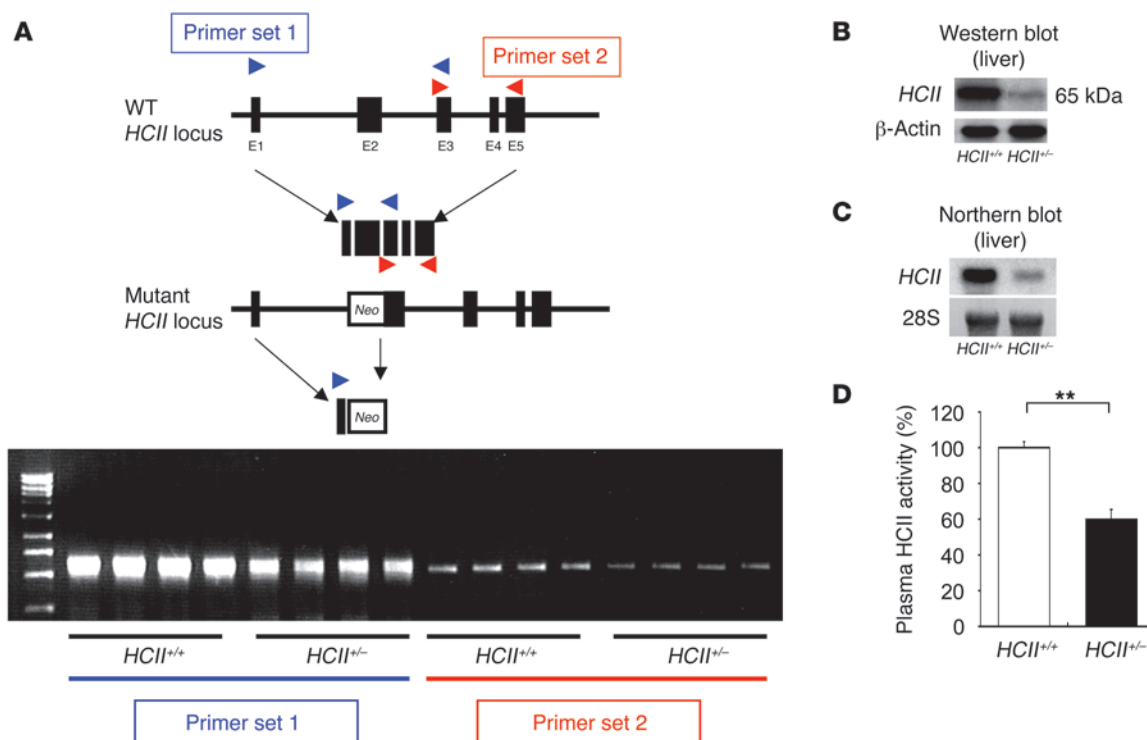
**Augmentation of cell proliferation in vascular walls of cuff-injured *HCII*<sup>-/-</sup> mice.** To identify and quantify proliferating vascular mesenchymal cells in these lesions, we performed proliferating cell nuclear antigen (PCNA) and BrdU staining. There were only a few positive nuclei for PCNA and BrdU staining at intima and media in both uninjured *HCII*<sup>+/+</sup> and *HCII*<sup>-/-</sup> mice (Figure 6, A–D). Although there was a marked increase in the number of PCNA- and BrdU-positive cells in both *HCII*<sup>+/+</sup> and *HCII*<sup>-/-</sup> mice after cuff injury, a much greater increase in PCNA- and BrdU-stained cells was observed in cuff-injured *HCII*<sup>-/-</sup> mice than in *HCII*<sup>+/+</sup> mice (Figure 6, A–D). These results demonstrate that the pronounced intimal hyperplasia in cuff-injured *HCII*<sup>-/-</sup> mice is due to enhanced proliferation of vascular mesenchymal cells.

**Enhanced vascular expression of PAR-1 in cuff-injured *HCII*<sup>-/-</sup> mice.** It has been well documented that enhanced PAR-1 expression results in development of restenosis after balloon angioplasty and atherosclerosis (7, 8, 26–28). Therefore, we hypothesized that one of the


**Figure 2**

Noninterference of *HCII* targeting vector introduction into murine genome. (A) FISH analysis using a *phosphoglycerate kinase*–*neomycin* cassette probe and genomic map around the *HCII* genome. Single green spots in male and female *HCII*<sup>+/+</sup> murine chromosomes indicate no random integration and adequate homologous recombination of the *HCII* targeting vector. Original magnification, ×1,000; ×4,000 (insets). (B) Real-time PCR analysis for mRNA quantification of genes located around the *HCII* genome in *HCII*<sup>+/+</sup> (white bars) and *HCII*<sup>-/-</sup> mice (black bars). Values were normalized by arbitrarily setting the measurement for *HCII*<sup>+/+</sup> mice to 1.0. *Ube2l3*, ubiquitin-conjugating enzyme E2L3; *Pik4ca*, phosphatidylinositol 4-kinase, catalytic, alpha polypeptide; *Snap29*, synaptosomal-associated protein.



**Figure 3**

Analysis of *HCII* gene transcripts, protein, and plasma activity. (A) Conventional RT-PCR analysis of full-length *HCII* gene. (B) Western blot analysis of hepatic *HCII* protein in *HCII*<sup>+/+</sup> and *HCII*<sup>-/-</sup> mice. (C) Northern blot analysis of hepatic *HCII* mRNA in *HCII*<sup>+/+</sup> and *HCII*<sup>-/-</sup> mice. (D) Plasma *HCII* activity in *HCII*<sup>+/+</sup> (white bar) and *HCII*<sup>-/-</sup> mice (black bar). Values are expressed as mean  $\pm$  SEM. \*\**P* < 0.01. *n* = 12 in each group for plasma *HCII* activity.

causes of accelerated intimal hyperplasia in cuff-injured *HCII*<sup>-/-</sup> mice is enhancement of thrombin receptor expression in the vascular wall. To test this hypothesis, we examined vascular expression of PAR-1 in the vascular walls of these mice. In uninjured vessels, PAR-1 expression was restricted to the thin endothelial layer in both *HCII*<sup>+/+</sup> and *HCII*<sup>-/-</sup> mice (Figure 7A). After cuff injury, PAR-1-expressing cells were distributed more widely, with a marked increase in the thickness of the intimal layer in *HCII*<sup>-/-</sup> mice compared with *HCII*<sup>+/+</sup> mice (Figure 7A).

*Enhanced expression of inflammatory cytokines, chemokines, and transcription factors involved in VSMC proliferation in cuff-injured HCII<sup>-/-</sup> mice.* Intimal and adventitial hyperplasia caused by 4 weeks of polyethylene tube cuff placement around femoral arteries has been shown to be due to enhanced inflammatory reactions in response to vascular injury. To quantify gene expression levels of vascular

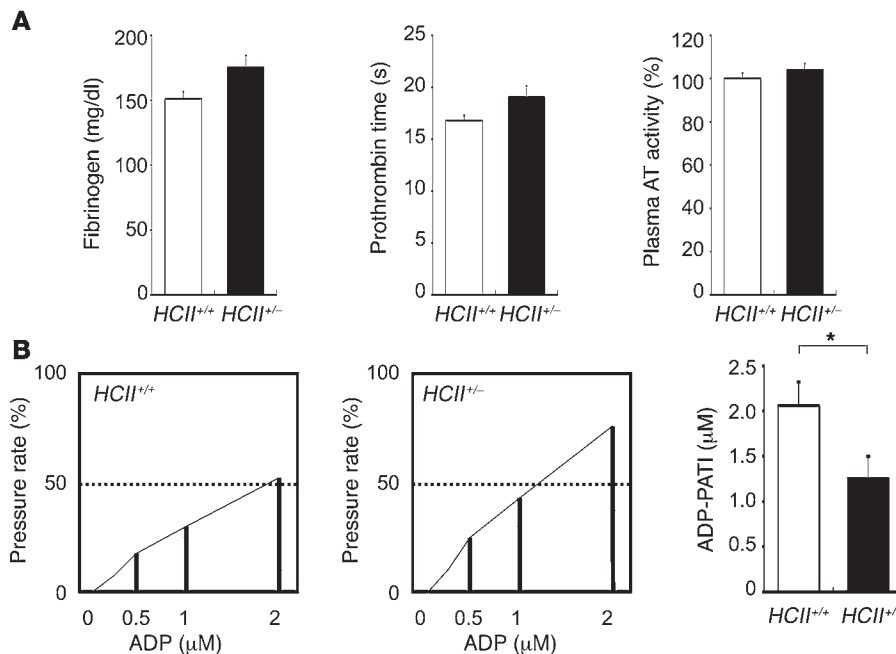
remodeling factors, including inflammatory cytokines, chemokines, and transcription factors, that enhance VSMC proliferation, we performed real-time PCR analysis using removed arteries containing surrounding tissues of these mice. Without cuff injury, there was no significant difference between gene expression levels of *Il1b*, *Il6*, monocyte chemoattractant protein-1 (*Mcp1*), early growth response protein-1 (*Egr1*) and Krüppel-like zinc finger transcription factor 5 (*Klf5*) in *HCII*<sup>+/+</sup> and *HCII*<sup>-/-</sup> mice (Figure 7B). Cuff injury caused notable enhancement of the expression of all of these genes in the vascular wall of *HCII*<sup>-/-</sup> mice, whereas only slight increases in the expression levels of these genes were observed in *HCII*<sup>+/+</sup> mice (Figure 7B). These results are consistent with the assumption that endogenous *HCII* attenuates inflammatory responses as well as transformation and migration of VSMCs in femoral arteries after cuff injury.

**Table 2**

Genotypes, plasma *HCII* antigen levels, and activity levels of second-generation mice derived from the mating of *HCII*<sup>+/-</sup> St. Louis and *HCII*<sup>+/-</sup> Tokushima mice

	Genotype of <i>HCII</i> gene			
	+/+	+/- (St. Louis)	+/- (Tokushima)	-/- (both mutants)
No. of pups	24	21	22	26
Plasma <i>HCII</i> antigen levels (%) <sup>A</sup>	102.0 $\pm$ 2.6	ND	43.4 $\pm$ 2.5	0.0 $\pm$ 0.1
Plasma <i>HCII</i> activity levels (%) <sup>A</sup>	101.3 $\pm$ 3.1	ND	59.4 $\pm$ 3.9	14.1 $\pm$ 2.0

Values are expressed as mean  $\pm$  SEM. <sup>A</sup>*n* = six 20- to 22-week-old mice in each group. ND, not determined; however, these values were estimated in ref. 25.

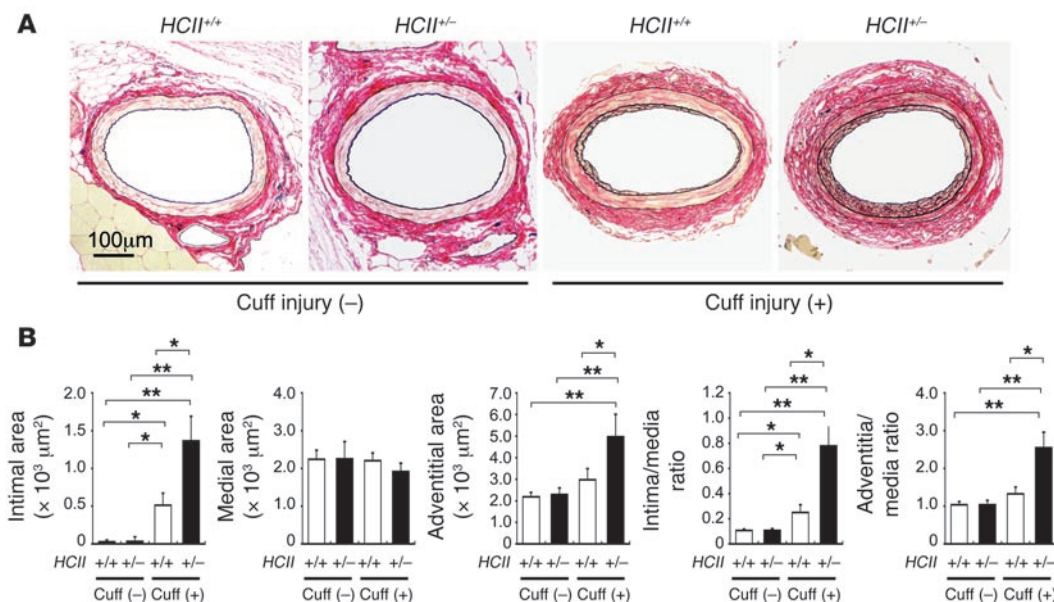
**Figure 4**

Effect of HCII deficiency on hemostatic examinations. **(A)** Plasma fibrinogen concentration, prothrombin time, and AT activity in *HCII*<sup>+/+</sup> (white bars) and *HCII*<sup>-/-</sup> mice (black bars). Values are expressed as mean ± SEM. *n* = 6–12 in each group. **(B)** The left and middle panels show representative results of platelet aggregation analyzed by the screen filtration pressure method in *HCII*<sup>+/+</sup> and *HCII*<sup>-/-</sup> mice. The platelet aggregation pressure of each reaction tube was determined as the pressure rate (percentage). For screen filtration pressure aggregometer study, the pressure rate was standardized using a grading curve produced by plotting 3 concentrations of ADP on the x axis and pressure rate (percentage) on the y axis. The concentration of ADP causing a 50% increase in pressure rate was calculated and applied as the platelet aggregatory threshold index (PATI) in *HCII*<sup>+/+</sup> (white bar) and *HCII*<sup>-/-</sup> mice (black bar). *n* = 6 in each group. Values are expressed as mean ± SEM. \**P* < 0.05.

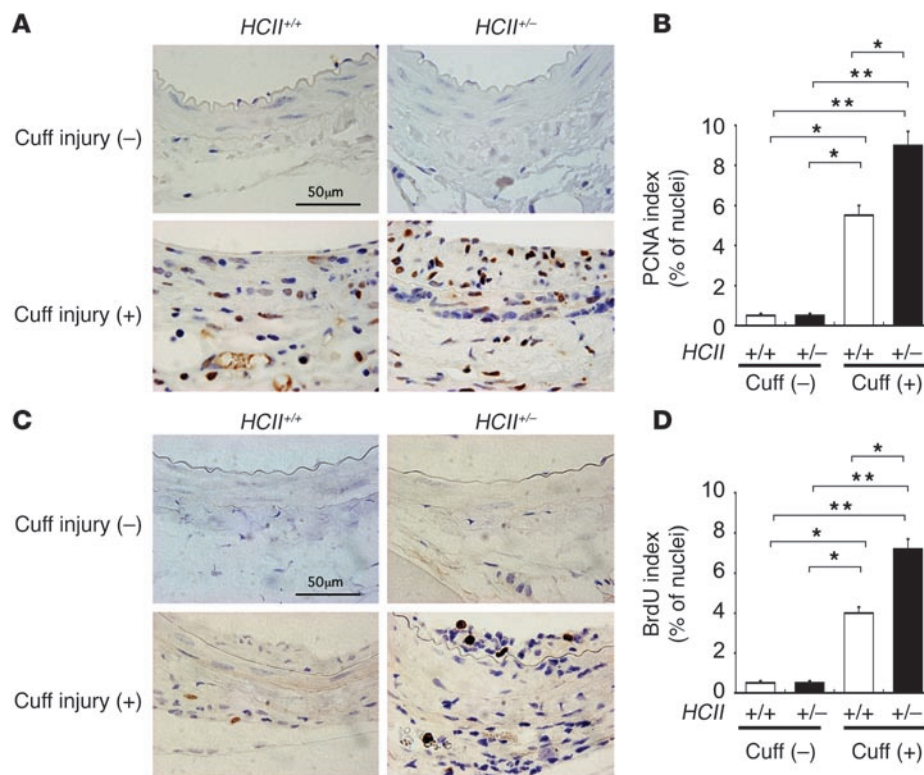
Human purified HCII protein ameliorates aberrant vascular remodeling in cuff-injured *HCII*<sup>-/-</sup> mice. In order to determine whether supplementation with a large amount of HCII in vivo can prevent accelerated vascular remodeling in cuff-injured *HCII*<sup>-/-</sup> mice, we administered human purified HCII protein for 4 weeks after surgery in *HCII*<sup>-/-</sup> mice. Since intraperitoneal injection of 45 mg/kg human purified HCII protein was able to maintain plasma HCII activity of greater than 100% for 48 hours in *HCII*<sup>-/-</sup> mice (Figure 8A), we administered this dose of HCII protein 3 times a week for 4 weeks. HCII protein

administration markedly attenuated intimal hyperplasia and adventitial hyperplasia in cuff-injured *HCII*<sup>-/-</sup> mice to levels similar to those observed in cuff-injured WT mice treated with vehicle (Figure 8, B and C). Thus, supplementation with HCII can ameliorate injury-induced vascular remodeling in *HCII*-deficient mice in vivo.

Enhanced neointimal hyperplasia, increased incidence of arterial thrombosis, and normalization of aberrant vascular remodeling by human purified HCII protein supplementation in wire-injured *HCII*<sup>-/-</sup> mice. In addition to the cuff injury model, we estimated vascular remodeling in

**Figure 5**

Histopathology of femoral arteries with and without cuff injury in *HCII*<sup>+/+</sup> and *HCII*<sup>-/-</sup> mice. **(A)** Histological features of uninjured and cuff-injured femoral arteries. Cross sections of uninjured and cuff-injured arteries were obtained on day 28 of cuff injury and stained with Elastica-van Gieson. **(B)** Quantitative analysis of arterial intimal area, medial area, adventitial area, intima/media ratio, and adventitia/media ratio with and without cuff injury in *HCII*<sup>+/+</sup> (white bars) and *HCII*<sup>-/-</sup> mice (black bars). Values are expressed as mean ± SEM. \**P* < 0.05; \*\**P* < 0.01. *n* = 12 in each group.



**Figure 6**

Immunostaining of proliferative vascular mesenchymal cells with PCNA and BrdU at the vascular wall with and without cuff injury in *HCII*<sup>+/+</sup> and *HCII*<sup>+/-</sup> mice. (A) PCNA staining of the vascular wall with and without cuff injury in *HCII*<sup>+/+</sup> and *HCII*<sup>+/-</sup> mice. (B) Quantitative analysis of PCNA-positive cells in intima and media with and without cuff injury in *HCII*<sup>+/+</sup> (white bars) and *HCII*<sup>+/-</sup> mice (black bars). Values are expressed as mean  $\pm$  SEM. \**P* < 0.05; \*\**P* < 0.01. *n* = 8 in each group. (C) BrdU staining of the vascular wall with and without cuff injury in *HCII*<sup>+/+</sup> and *HCII*<sup>+/-</sup> mice. (D) Quantitative analysis of BrdU-positive cells in intima and media with and without cuff injury in *HCII*<sup>+/+</sup> (white bars) and *HCII*<sup>+/-</sup> mice (black bars). Values are expressed as mean  $\pm$  SEM. \**P* < 0.05; \*\**P* < 0.01. *n* = 8 in each group.

*HCII*<sup>+/+</sup> and *HCII*<sup>+/-</sup> mice using a wire insertion injury model (29). Wire-injured femoral arteries in *HCII*<sup>+/-</sup> mice revealed prominent neointimal hyperplasia and a high intima/media ratio compared with those in *HCII*<sup>+/+</sup> mice (Figure 9, A and B). Moreover, wire injury caused a higher incidence (20%) of arterial occlusion due to thrombosis in *HCII*<sup>+/-</sup> mice than in *HCII*<sup>+/+</sup> mice (10%) (Figure 9A). Those abnormalities were ameliorated by human purified HCII protein supplementation, as was found in the cuff injury model in *HCII*<sup>+/-</sup> mice (Figure 9, A and B).

*HCII* deficiency causes accelerated atherosclerotic plaque formation and PAR-1 expression in the aortic root of *apoE*<sup>-/-</sup> mice. In order to examine the role of HCII action against lipid-induced atherosclerosis, we generated *HCII*<sup>+/-</sup>*apoE*<sup>-/-</sup> mice and compared the double-mutant mice with *HCII*<sup>+/+</sup>*apoE*<sup>-/-</sup> mice. There were no significant differences in fasting plasma levels of total cholesterol, triglyceride, high-density lipoprotein cholesterol, free fatty acid, and glucose (*HCII*<sup>+/+</sup>*apoE*<sup>-/-</sup> versus *HCII*<sup>+/-</sup>*apoE*<sup>-/-</sup> mice: 300  $\pm$  15.4 versus 280.1  $\pm$  17.2 mg/dl; 41.3  $\pm$  2.6 versus 44.5  $\pm$  4.9 mg/dl; 10.3  $\pm$  0.8 versus 11.1  $\pm$  1.3 mg/dl; 554.2  $\pm$  58.4 versus 506.4  $\pm$  54.5 mEq/l; and 81.1  $\pm$  5.0 versus 81.7  $\pm$  6.4 mg/dl; respectively; *n* = 14 in each group). Atherosclerotic plaque area was significantly increased in the aortic root of *HCII*<sup>+/-</sup>*apoE*<sup>-/-</sup> mice compared with that in *HCII*<sup>+/+</sup>*apoE*<sup>-/-</sup> mice (Figure 10A). In addition, lipid deposition and PAR-1-positive cells in the plaques were more prominently observed in the aortic root of *HCII*<sup>+/-</sup>*apoE*<sup>-/-</sup> mice than *HCII*<sup>+/+</sup>*apoE*<sup>-/-</sup> mice (Figure 10B).

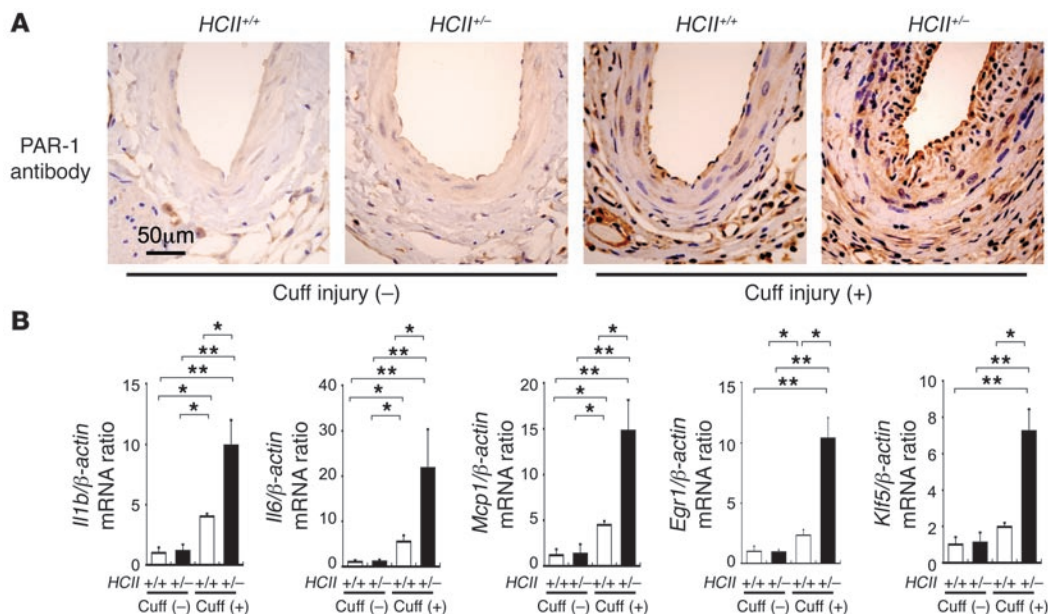
*HCII* deficiency increases oxidative stress in *apoE*<sup>-/-</sup> mice. Because it has been recognized that increased oxidative stress promotes cardiovascular diseases such as atherosclerosis and hypertension (30), we investigated superoxide production in *HCII*<sup>+/+</sup>*apoE*<sup>-/-</sup> and *HCII*<sup>+/-</sup>*apoE*<sup>-/-</sup> mice. The number of dihydroethidium-stained (DHE-stained) spots in the aortic root was greater in *HCII*<sup>+/-</sup>

*apoE*<sup>-/-</sup> mice than in *HCII*<sup>+/+</sup>*apoE*<sup>-/-</sup> mice (Figure 11A). Moreover, daily urinary excretion levels of 8-hydroxy-2'-deoxyguanosine (8OHdG), as a marker of oxidative stress-induced DNA damage, were higher in *HCII*<sup>+/-</sup>*apoE*<sup>-/-</sup> mice than in *HCII*<sup>+/+</sup>*apoE*<sup>-/-</sup> mice (Figure 11B). These results suggest that HCII attenuates oxidative stress and cellular damage. The mechanisms by which HCII exerts such an effect remain to be clarified.

## Discussion

In the present study, *HCII*-deficient mice were created by targeted disruption of the *HCII* gene and were used to examine the role as well as the mechanism of protective action of HCII against thrombin-induced vascular remodeling. Unexpectedly, mice with homozygous deletion of *HCII* exhibited embryonic lethality after careful backcrossing for 10 generations with C57BL/6J strain (Table 1 and Figure 1C). *HCII*<sup>-/-</sup> mice generated previously by He et al. survived throughout the fetal period and after birth with normal growth and life span (25). He et al. deleted exon 1 in their targeting vector, which is the same as exon 2 in our description of the *HCII* gene. Based on the results of a study by Kamp and Ragg showing that the genomic structure of murine and human *HCII* consists of 5 exons that are interrupted by 4 introns (31), we called exon 1 in their report exon 2 in this paper (Figure 1A). He et al. inserted a *phosphoglycerate kinase-neomycin* cassette to replace exon 2 (exon 1 in their description), whereas we inserted a *neomycin* cassette with polyA termination to silence transcription. In addition, they screened a bacterial artificial chromosome (BAC) library made from 129/SvJ mouse genomic DNA to construct the *HCII* targeting vector, and the targeting vector was introduced into 129/SvJ ES cells. Then their *HCII* heterozygous mutant mice (*HCII*<sup>+/-</sup> St. Louis) were backcrossed with WT C57BL/6 mice to produce a nearly congenic





**Figure 7**

Expression of PAR-1, inflammatory cytokines and chemokines, and vascular remodeling-related transcription factors with and without cuff injury in *HCII*<sup>+/+</sup> and *HCII*<sup>+/-</sup> mice. **(A)** Immunohistochemical staining of PAR-1 with and without cuff injury in *HCII*<sup>+/+</sup> and *HCII*<sup>+/-</sup> mice. **(B)** Gene expression of vascular remodeling factors at the vascular wall with and without cuff injury in *HCII*<sup>+/+</sup> (white bars) and *HCII*<sup>+/-</sup> mice (black bars). Real-time PCR analyses of *Il1b*, *Il6*, *Mcp1*, *Egr1*, and *Klf5* were performed in *HCII*<sup>+/+</sup> and *HCII*<sup>+/-</sup> mice. Values were normalized by arbitrarily setting the measurement in *HCII*<sup>+/+</sup> mice without cuff injury to 1.0. Values are expressed as mean  $\pm$  SEM. \**P* < 0.05; \*\**P* < 0.01. *n* = 10 in each group.

strain of mice containing the *HCII*-null allele (25). On the other hand, we screened a phage library made from C57BL/6 mouse genomic DNA, and we introduced the targeting vector into TT2 (hybrid with C57BL/6J and CBA) ES cells. Our *HCII* heterozygous mutant mice (*HCII*<sup>+/-</sup> Tokushima) were mated with WT C57BL/6J mice and backcrossed for 10 generations. From these results, we have reached the conclusion that complete KO of *HCII* gene results in embryonic lethality. We do not have plausible explanation for the discrepancy in fetal development between the studies by He et al. and ours. We speculate that a minor strain-dependent difference in genetic background may have caused the loss of embryonic lethality of *HCII*<sup>-/-</sup> St. Louis mice.

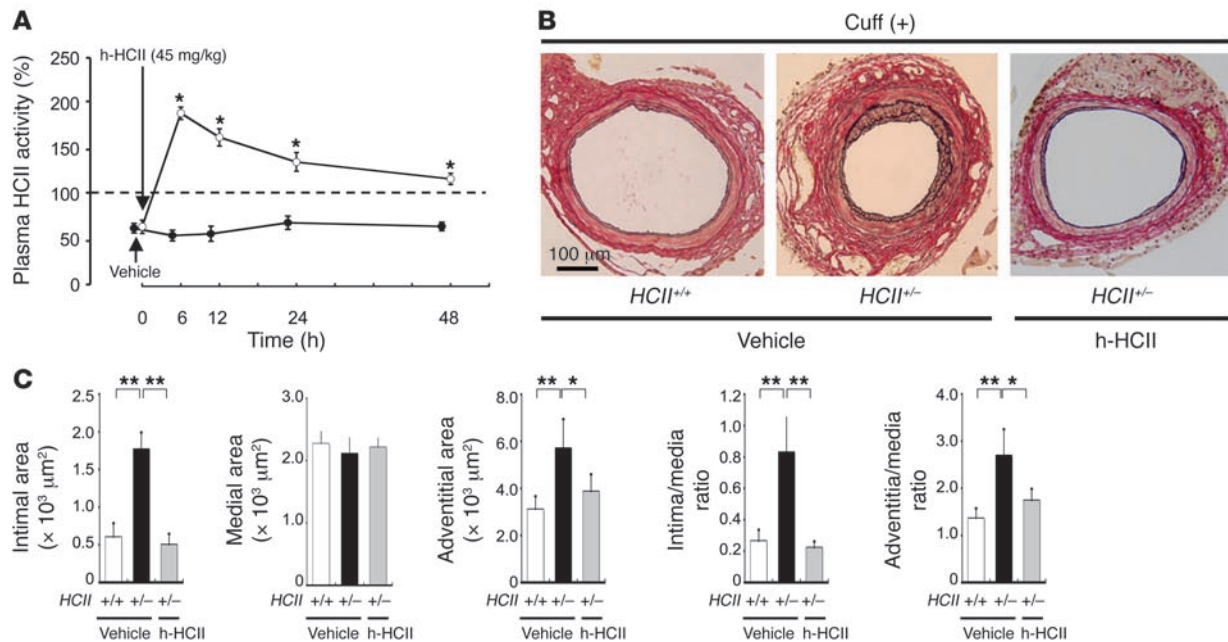
Although a family with congenital homozygous *HCII* deficiency was reported (32) with a Glu428Lys missense mutation in both alleles (33), the homozygous *HCII*-deficient patients showed 5%–13% of normal *HCII* activity, and there has been no report of humans with complete *HCII* deficiency. Delorme et al. reported that thrombin generation is increased during normal pregnancy and that a greater proportion of thrombin molecules form a complex with *HCII* in plasma in pregnant woman than in nonpregnant women (34). It has also been reported that *HCII* levels are significantly decreased in women with severe preeclampsia (35). Giri and Tollefsen have suggested that DS activates *HCII* in the fetal blood vessels or in the stroma of placental villi after injury to the syncytiotrophoblast layer and thereby inhibit fibrin generation in the placenta (36). Thus, a certain level of *HCII* activity may be required for the survival of human fetuses. In the present study, *HCII* homozygous mutant mice generated by mating *HCII*<sup>+/-</sup> Tokushima mice with *HCII*<sup>+/-</sup> St. Louis mice survived. Although

plasma *HCII* antigen was not detected in *HCII*<sup>-/-</sup> Tokushima/St. Louis mice, very low levels of plasma *HCII* activities remained in those mice. Because those murine plasma *HCII* activities were measured using the STACHROM *HCII* colorimetric assay (Diagnostica Stago) designed for measurement of human plasma, there is a possibility that the assay may have overestimated *HCII* activity in mice. Nevertheless, the possibility remains that at least a certain level of plasma *HCII* activity may be necessary for survival of fetuses during pregnancy in mice as well.

Humans with heterozygous *HCII* deficiency develop only occasional deep vein thrombosis and have no atherosclerotic lesions or vascular damage while they are

young. Although larger clinical studies are needed, heterozygous *HCII*-deficient subjects appear to suffer from advanced atherosclerotic lesions after the development of vascular injuries with aging (37). Indeed, a female member of a family with type I congenital *HCII* deficiency with about half of the normal plasma *HCII* activity level was found to have multiple atherosclerotic lesions, including lesions in coronary, carotid, renal, and abdominal aortic arteries, when she was 66 years old (22). Her elder brother, who had the same heterozygous mutation and reduced level of plasma *HCII* activity, also had multiple cerebral infarctions when he was 63 years of age. In addition, our 2 clinical studies on the protective role of *HCII* against the development of atherosclerosis demonstrated a protective role of *HCII* against progression of carotid atherosclerosis and in-stent restenosis after percutaneous coronary intervention (23, 24, 38). Schillinger et al. also demonstrated that a high level of plasma *HCII* activity is associated with reduced incidence of restenosis after femoropopliteal stenting (39). In the present study, *HCII*<sup>+/-</sup> mice exhibited no obvious differences in vascular structure compared with *HCII*<sup>+/+</sup> mice under the condition of no vascular stress. However, arterial cuff injury and wire injury caused prominent hyperplasia of the intimal area in *HCII*<sup>+/-</sup> mice compared with *HCII*<sup>+/+</sup> mice (Figures 5 and 9). Since *HCII* exerts its antithrombin action via a complex formation with DS and since DS is produced by VSMCs or fibroblasts and is deposited within the matrix of the vascular wall, the antithrombin action of *HCII* is exerted mostly at the injured vascular wall, where *HCII* is activated by its binding to DS. In the present study, cuff injury of the femoral artery of *HCII*<sup>+/-</sup> mice resulted in broad expression of PAR-1 in the vascular wall (Figure 7A). The enhanced expression of



**Figure 8**

Effects of human purified HCII supplementation on cuff-injured vascular remodeling in *HCII*<sup>+/-</sup> mice. **(A)** Time course of plasma HCII activity in *HCII*<sup>+/-</sup> mice after an intraperitoneal injection of human purified HCII protein (h-HCII) (open circles) or a vehicle alone (filled circles). \**P* < 0.05. *n* = 4 in each group. **(B)** Representative histological findings of cuff-injured femoral arteries. Micrographs of cross sections of cuff-injured arteries stained with Elastica-van Gieson on day 28 in *HCII*<sup>+/+</sup> mice treated with vehicle and in *HCII*<sup>+/-</sup> mice treated with vehicle alone or human purified HCII protein are shown. **(C)** Quantitative comparison of arterial intimal area, medial area, adventitial area, intima/media ratio, and adventitia/media ratio in cuff-injured *HCII*<sup>+/+</sup> mice treated with vehicle (white bars) and *HCII*<sup>+/-</sup> mice treated with vehicle (black bars) or human purified HCII protein (gray bars). Values are expressed as mean  $\pm$  SEM. \**P* < 0.05; \*\**P* < 0.01. *n* = 10 in each group.

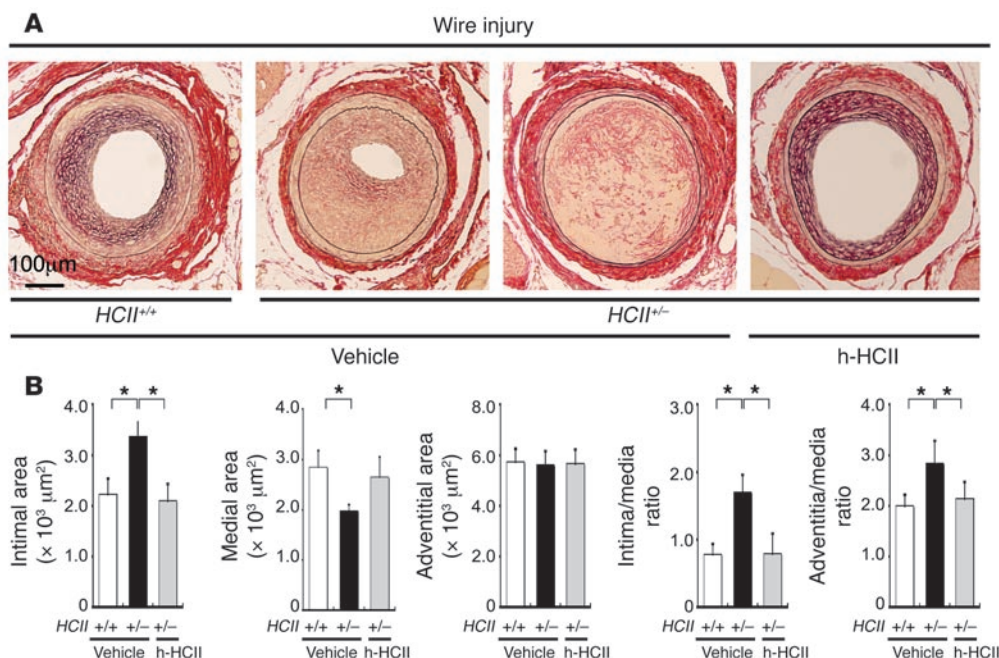
PAR-1 in turn is expected to further promote vascular remodeling in *HCII*<sup>+/-</sup> mice. Taken together, these results are consistent with the assumption that the vascular protective role of HCII is exerted most efficiently at sites with vascular injury where thrombin is activated to stimulate vascular remodeling.

Thrombin stimulates migration and proliferation of mesenchymal cells, including VSMCs, via binding to its cell-surface receptors, PARs (40–42). Receptor-mediated actions of thrombin enhance the expression of inflammatory cytokines and chemokines and are associated with pathological conditions such as atherosclerosis and restenosis after vascular interventions (43, 44). Activation of PAR-1 by thrombin also enhances NF- $\kappa$ B activation in endothelial cells (45) and VSMCs (46). Because upregulation of these cytokines and chemokines that contribute to the promotion of vascular remodeling (47–50) is known to enhance NF- $\kappa$ B activation, there is a possibility that the action of thrombin on migration and proliferation of cells in the vascular wall is mediated by its stimulatory effect on the expression of inflammatory cytokines and chemokines. This possibility is supported by an observation by Isoda et al. that deficiency of an IL-1 receptor antagonist promotes neointimal formation in mice after cuff injury (51). The present study demonstrated that gene expression of inflammatory cytokines (*Il1b* and *Il6*), a chemokine (*Mcp1*), and transcriptional factors involved in VSMC proliferation (*Egr1* and *Klf5*) was markedly upregulated in cuff-injured *HCII*<sup>+/-</sup> mice (Figure 7B). Based on these observations, we suggest that HCII inhibits vascular remodeling via its inhibitory effect on the expression of these cytokines and chemokines by blocking receptor-mediated thrombin action.

In addition, wire injury caused not only prominent neointimal hyperplasia but also a high incidence of arterial occlusion due to thrombosis. This result is consistent with the previous finding that *HCII*-deficient mice manifested acceleration of arterial thrombosis formation after photochemical endothelium injury (25). In the present study, we found that HCII deficiency enhanced platelet aggregation. Since PARs exists in platelets, HCII may have a suppressive effect on platelet aggregation.

Since it is well known that dyslipidemia promotes arterial plaque formation leading to cardiovascular events, we investigated the role of HCII action against high cholesterol-induced atherosclerosis using *HCII*<sup>+/-</sup>*apoE*<sup>-/-</sup> mice. As expected, the double mutant mice manifested exaggeration of atherosclerotic lesion formation with increased lipid deposition and enhanced PAR-1 expression. Interestingly, the double mutant mice showed higher levels of oxidative stress than *apoE* single mutant mice. These results suggest that HCII has antioxidant potency at the injured vascular wall. Based upon these observations, inactivation of PAR-mediated thrombin actions and attenuation of oxidative stress at sites of injury of the arterial wall is an attractive approach for protection against atherosclerosis and restenosis after percutaneous coronary intervention.

Our observations were further corroborated by a study using *HCII*<sup>-/-</sup> St. Louis mice on an *apoE*<sup>+/+</sup> or *apoE*<sup>-/-</sup> background (Douglas M. Tollefsen, unpublished observations). In that study, *HCII*<sup>-/-</sup> St. Louis mice exhibited greater neointimal hyperplasia of carotid arteries after wire injury than *HCII*<sup>+/+</sup> mice. In addition, it was revealed that *HCII*<sup>-/-</sup>*apoE*<sup>-/-</sup> mice developed larger aortic



**Figure 9**

Effects of HCII deficiency and of human purified HCII supplementation on wire-injured vascular remodeling. (A) Representative histological findings for wire-injured femoral arteries. Micrographs of cross sections of wire-injured arteries stained with Elastica–van Gieson on day 28 in *HCII*<sup>+/+</sup> mice treated with vehicle and in *HCII*<sup>-/-</sup> mice treated with vehicle alone or human purified HCII protein are shown. (B) Quantitative comparison of arterial intimal area, medial area, adventitial area, intima/media ratio, and adventitia/media ratio in wire-injured *HCII*<sup>+/+</sup> mice treated with vehicle (white bars) and *HCII*<sup>-/-</sup> mice treated with vehicle (black bars) or human purified HCII protein (gray bars). Values are expressed as mean ± SEM. \**P* < 0.05. *n* = 8–10 in each group.

plaque formation than *HCII*<sup>+/+</sup>*apoE*<sup>-/-</sup> mice. Taken together, the results of the present study and those of the study by Tollefsen and colleagues demonstrate that HCII protects against thrombin-induced remodeling of an injured vascular wall by inhibiting thrombin action. Since HCII is activated mainly at the vascular wall by complex formation with DS, HCII or agents that mimic these characteristics of HCII can act mostly at the vascular wall. Such agents can protect against vascular remodeling without causing coagulatory disturbance or bleeding tendency in patients with atherosclerosis or cardiovascular disease.

## Methods

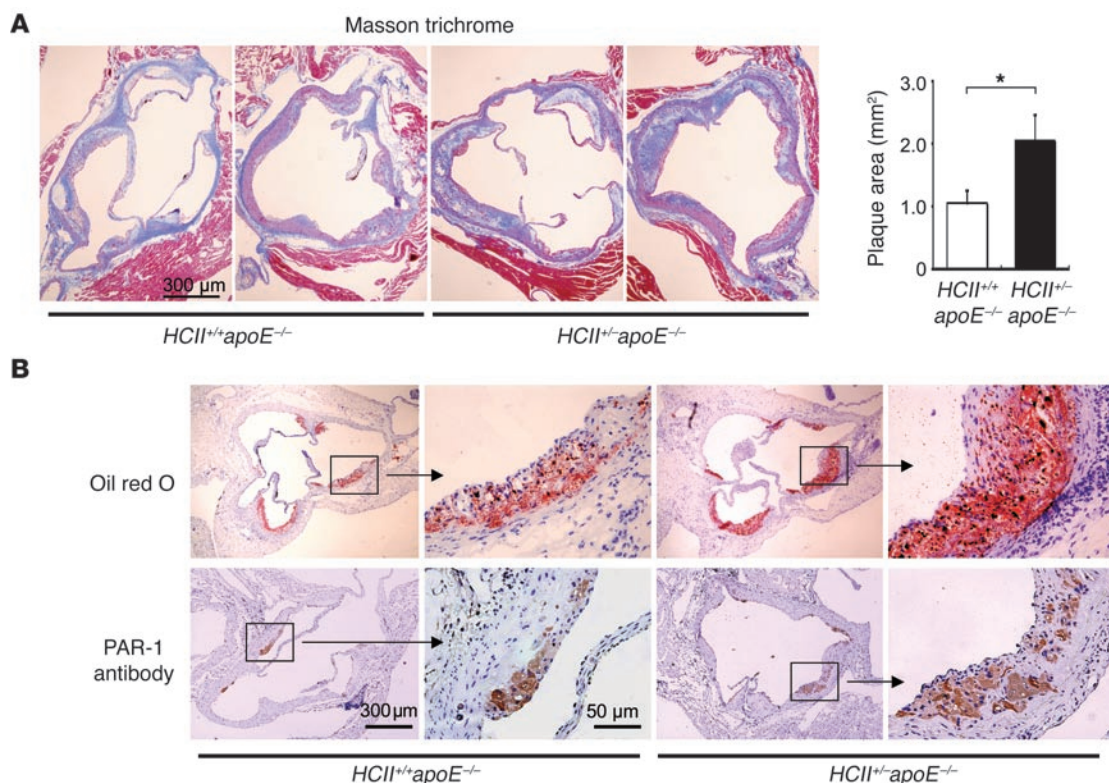
**Gene targeting and genotyping for HCII mutant mice.** A mouse genomic C57BL/6 library (female, 6–8 weeks old, vector: Lambda DASH II, insert size: 9–23 kb; Stratagene) was screened with a human *HCII* cDNA probe. A 17-kb fragment of the mouse *HCII* genome containing coding exons 2–5 was used to construct a targeting vector. A 120-bp fragment containing the ATG sequence of exon 2 was deleted, and a *phosphoglycerate kinase–neomycin* cassette including polyA termination was inserted. Diphtheria toxin fragment A gene (DTA) was bound to the 5' end of the targeting vector for negative selection. TT2 (hybrid with C57BL/6J and CBA) ES cells were transfected with a linearized targeting vector (25 mg per 1.0 × 10<sup>7</sup> cells) using a Bio-Rad Gene Pulser II at 250 V and 500 F and grown under G418 selection as described previously (52–55). Targeted ES cell clones were identified by Southern blot analysis with probe A as an external probe (WT allele denotes a 12-kbp band; mutant allele denotes a 5-kbp band) and probe B as an internal probe (mutant allele without random integration denotes a single 7-kbp band; Figure 1A) after *HindIII* diges-

tion and were aggregated with CD-1 8-cell embryos to generate chimeras as described previously (52–55). Chimeras were crossed with C57BL/6 female mice to produce germline transmission of the targeted allele. Offspring were genotyped either by Southern blotting with probes A and B or by PCR with 3 specific primers (WT allele forward, 5'-GTGTTA-AGAGAATCTGCTGTCTGC-3', and reverse, 5'-GTAGCCTG-GTCCTTCAGGACTCGG-3'; *HCII* mutant allele forward, 5'-CTTCCTCGTGTCTTACGGTA-3', and identical reverse primer). The PCR conditions were 35 cycles of denaturation for 30 seconds at 94°C, annealing for 30 seconds at 54°C, and extension for 45 seconds at 72°C. The WT allele gave a 480-bp band, and the mutant allele gave a 740-bp band. PCRs for the WT allele and mutant allele detections were performed separately. These offspring mice were backcrossed to C57BL/6J mice for 10 generations. After the backcrossing, male and female *HCII* heterozygous mutant mice

were intercrossed to obtain homozygous mutant mice. In addition, we obtained reproductive *HCII* heterozygote mutant mice established by Tollefsen's laboratory (25). Genotyping for detection of their *HCII* mutant allele was performed as described previously (25). Reproductive C57BL/6 background *apoE*<sup>-/-</sup> mice were purchased from Taconic. *HCII*<sup>+/+</sup>*apoE*<sup>-/-</sup> mice and *HCII*<sup>-/-</sup>*apoE*<sup>-/-</sup> mice were generated by crossbreedings of male and female *HCII*<sup>+/+</sup>*apoE*<sup>-/-</sup> mice. These animals were housed in a specific pathogen-free facility under climate-controlled conditions with a 12-hour light/12-hour dark cycle and were provided with standard diet chow (Oriental Yeast Co.) and water ad libitum. *HCII*<sup>+/+</sup> and *HCII*<sup>-/-</sup> mice were sacrificed and analyzed at the age of 20 weeks, whereas *HCII*<sup>+/+</sup>*apoE*<sup>-/-</sup> mice and *HCII*<sup>-/-</sup>*apoE*<sup>-/-</sup> mice were sacrificed and analyzed at the age of 30 weeks. All experimental procedures were approved of by the Animal Research Committee of the University of Tokushima Graduate School.

**Chromosome preparation and staining of splenic lymphocytes in *HCII* mutant mice.** Culture of murine lymphocytes was performed as described previously (56). Lymphocytes were isolated from spleens of adult *HCII*<sup>+/+</sup> male and female mice, washed twice with PBS solution, and transferred to 65-mm dishes containing 5 ml RPMI 1640 supplemented with 20% fetal calf serum, 3 mg/ml concanavalin A (Type IV-S; Sigma-Aldrich), 10 mg/ml lipopolysaccharide (Sigma-Aldrich), and 5 × 10<sup>-5</sup> M mercaptoethanol. Cultures with concentrations of 0.5 × 10<sup>6</sup> to 1.0 × 10<sup>6</sup> cells/ml were incubated at 37°C in a humidified mixture of 95% air and 5% CO<sub>2</sub>. Either thymidine (300 mg/ml) or 2 × 10<sup>-7</sup> M methotrexate and 200 mg/ml BrdU (Sigma-Aldrich) were added to the cultures, and then the lymphocytes were washed and transferred to culture medium. Finally, the cells were fixed with 3:1 methanol/glacial acetic acid and were air dried. The chromosome slides were stained with 3% Giemsa solution (pH 6.8) for analysis of G- and R-banding patterns.





**Figure 10**

Quantification and characterization of atherosclerotic lesions in *HCII*<sup>+/+</sup>*apoE*<sup>-/-</sup> and *HCII*<sup>+/-</sup>*apoE*<sup>-/-</sup> mice. **(A)** The left panel shows representative histological findings in Masson trichrome–stained serial sections of the aortic root in *HCII*<sup>+/+</sup>*apoE*<sup>-/-</sup> and *HCII*<sup>+/-</sup>*apoE*<sup>-/-</sup> mice. The right panel shows mean values of the atherosclerotic plaque area in *HCII*<sup>+/+</sup>*apoE*<sup>-/-</sup> (white bar) and *HCII*<sup>+/-</sup>*apoE*<sup>-/-</sup> mice (black bar). Values are expressed as mean ± SEM. \**P* < 0.05. *n* = 12 in each group. **(B)** Representative histological findings in oil red O– (upper panels) and PAR-1–stained (lower panels) serial sections of the aortic root in *HCII*<sup>+/+</sup>*apoE*<sup>-/-</sup> and *HCII*<sup>+/-</sup>*apoE*<sup>-/-</sup> mice.

**FISH analysis using a phosphoglycerate kinase–neomycin cassette probe.** Phosphoglycerate kinase–neomycin cassette cDNA of 1.6 kbp was labeled by nick translation (Roche) with biotin 16-dUTP (Roche). The labeled probe was ethanol precipitated with 625 µg/ml sonicated salmon sperm DNA and transfer RNA and then denatured for 10 minutes at 75°C in 100% formamide and kept at 4°C. The hybridization solution was mixed with an equal volume of denatured probe in formamide to make a final concentration of 50% formamide, 2× sodium chloride–sodium citrate, 10% dextran sulfate, and 2 mg/ml BSA. The biotin-labeled DNA was mixed at a final concentration of 5 µg/ml and used at 100 ng per slide. The mixture was put on denatured chromosome slides, covered with paraffin, and incubated overnight at 37°C. In situ hybridization of metaphase chromosomes was subsequently performed as described previously (56).

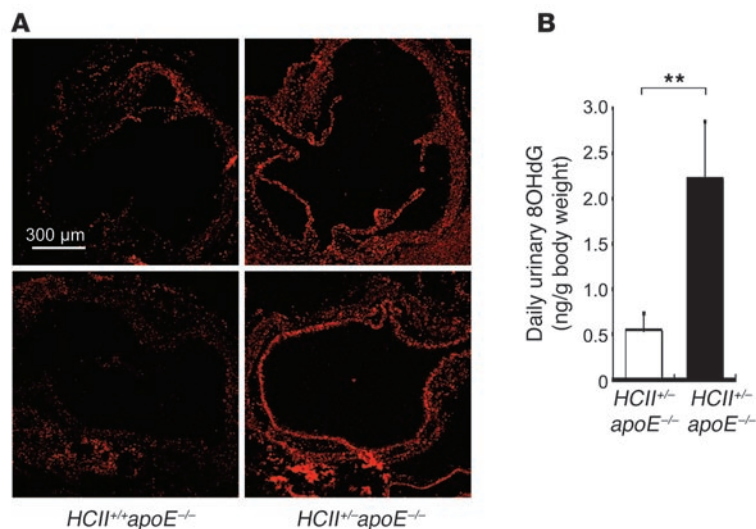
**RNA isolation.** Liver and femoral arteries were perfused and rinsed with physiological saline. Total RNA was isolated by acid guanidinium thiocyanate–phenol–chloroform extraction using TRIzol Reagent (Invitrogen). RNA concentrations were measured spectrophotometrically at 260 nm, and samples were stored in diethylpyrcarbonate-treated water at –80°C.

**Conventional RT-PCR analysis of full-length *HCII* gene.** Hepatic RNA was prepared with the use of TRIzol reagent (Invitrogen), and conventional RT-PCR to detect *HCII* gene transcripts was performed as described previously (57). PCR primers were designed for the front side of the *HCII* gene (top of exon 1 to tail of exon 3): forward, 5'-CTGCCAGACCACTGCTCAGAGGTA-3', and reverse, 5'-CTGTTTGTTCATGCTTTTTTGGCC-3'; and the tail side of the *HCII* gene (top of exon 3 to tail of exon 5): forward, GAACTTGGGT-

GAATAAATCCCAG-3', and reverse, 5'-CATGGTAAATATGGTTGTTTA-3'. The PCR conditions were 35 cycles of denaturation for 60 seconds at 94°C, annealing for 60 seconds at 54°C, and extension for 90 seconds at 72°C. The WT allele of the front side of the *HCII* gene gave a 1,264-bp band, and that of the tail side of the *HCII* gene gave a 1,139-bp band.

**Northern blot analysis.** Approximately 20 µg of total RNA of liver was fractionated on 1% formaldehyde–agarose gels and transferred to Hybond nylon membranes (GE Healthcare) by capillary action in high-salt solution (20× sodium chloride–sodium citrate). Blots were prehybridized in a hybridization solution for 1 hour at 42°C, followed by overnight hybridization with digoxigenin-labeled specific oligonucleotide probes (DIG Northern Starter Kit; Roche Diagnostics) as described previously (57). *HCII* mRNAs were estimated after correction for loading differences by measuring the amount of 28S rRNA. The intensity of the bands was also measured using ImageJ 1.29 (<http://rsb.info.nih.gov/ij/>). The forward and reverse sequences of the *HCII* oligonucleotide probe were 5'-GAACTTGGGTGAATAAATCCCAG-3' and 5'-CTGTTTGTTCATGCTTTTTTGGCC-3', respectively.

**Western blot analysis.** Hepatic tissue samples were homogenized in an ice-cold tissue protein extraction reagent (T-PER; Pierce) containing protease inhibitors. Equal amounts of protein (50 µg) were loaded onto 10% acrylamide gels and then transferred to nitrocellulose membranes using a wet blot apparatus. Membranes were blocked overnight at 4°C in blocking buffer (10 mM Tris-HCl, 10 mM NaCl, 0.1% polyoxyethyl- enesorbitan monolaurate) with 5% skim milk. The first antibody, goat



**Figure 11**  
Effect of HCII deficiency on oxidative stress in *apoE*<sup>-/-</sup> mice. (A) Representative findings in serial sections of the aortic root stained with DHE in *HCII*<sup>+/+</sup>*apoE*<sup>-/-</sup> and *HCII*<sup>-/-</sup>*apoE*<sup>-/-</sup> mice. (B) Mean values of urinary excretion of 8OHdG in *HCII*<sup>+/+</sup>*apoE*<sup>-/-</sup> (white bar) and *HCII*<sup>-/-</sup>*apoE*<sup>-/-</sup> mice (black bar). Values are expressed as mean ± SEM. \*\**P* < 0.01. *n* = 6 in each group.

anti-human HCII antibody (1:1,000 dilution; Affinity Biologicals Inc.), was applied for 1 hour at room temperature. After a 30-minute wash with the blocking buffer, membranes were probed with the second antibody [1:5,000 dilution peroxidase-conjugated AffiniPure F(ab')<sub>2</sub> fragment rabbit anti-goat IgG antibody; Jackson ImmunoResearch Laboratories Inc.] for 1 hour at room temperature. Membranes were washed again for 30 minutes. The membranes were then incubated with chemiluminescent reagents (Amersham ECL Plus Western Blotting Detection System; GE Healthcare) for 5 minutes and then exposed to x-ray film. The band intensities were quantified using ImageJ 1.29.

**Measurements of plasma HCII activities and hemostatic evaluations of murine plasma.** Blood was drawn through a heart puncture and collected into a tube containing a 1:10 volume of 3.8% sodium citrate and then centrifuged at 2,000 *g* for 20 minutes. Plasma was stored at -80°C until use. Plasma HCII activity was measured on the basis of antithrombin activity in the presence of DS, with the use of the STACHROM HCII colorimetric assay (Diagnostica Stago). HCII activity of pooled murine plasma of 16-week-old C57BL/6J mice (8 males) was set at 100%. The intraassay and interassay coefficients of variation of this kit were 3.9% and 4.3%, respectively (23, 24). Fibrinogen concentration was evaluated by Clauss's coagulometric technique, and prothrombin time and plasma AT activity were measured as described previously (57). Measurements of platelet aggregation with a screen filtration pressure aggregometer (WBA analyzer from SSR Engineering Co.) were performed according to a previously described method (57).

**Measurements of plasma HCII antigen levels.** Murine plasma HCII antigen levels were estimated by ELISA described previously as online supplemental data (<http://atvb.ahajournals.org/cgi/content/full/25/12/2689/DC1>; ref. 58).

**Vascular injury by cuff placement around femoral arteries.** Cuff placement surgery was carried out on 16-week-old male mice as described previously (48). In brief, the right femoral artery was isolated from surrounding tissues, and then a polyethylene tube (2-mm PE-50; inner diameter, 0.56 mm; outer diameter, 0.965 mm; BD) was cut longitudinally, loosely placed around the artery, and then closed with suture. At the same time, the left femoral artery was also isolated from surrounding tissues and

immediately sutured without cuff placement. Four weeks after the surgery, the mice were sacrificed with high-dose pentobarbital and perfused first with physiological saline and then with 10% neutral buffered formalin at 100 mmHg. Then the cuffed arteries and sham-operated arteries were removed with surrounding tissues.

**Vascular injury by wire insertion into femoral arteries.** Wire insertion surgery was performed on 16-week-old male mice as described previously (59). Either the left or right femoral artery was exposed by blunt dissection. A small branch between the rectus femoris and vastus medialis muscles was isolated and looped proximally and ligated distally with 6-0 silk sutures. Veins and connective tissues around the artery were removed. The exposed muscular branch artery was dilated by topical application of 1 drop of 1% lidocaine hydrochloride. Transverse arteriotomy was performed in the muscular branch with Vannas-style iris spring scissors (Fine Science Tools). A straight spring wire (0.38 mm in diameter; no. C-SF-15-15; COOK Inc.) was inserted into the femoral artery for more than 5 mm toward the iliac artery. The wire was left in place for 1 minute to denude and dilate the artery. Then the wire was removed, and the silk suture looped at the proximal portion of the muscular branch artery was secured. Four weeks after the surgery, the mice were sacrificed with high-dose pentobarbital and perfused first with physiological saline and then with 10% neutral buffered formalin at 100 mmHg. Then the cuffed arteries and sham-operated arteries were removed with surrounding tissues.

**Real-time PCR analysis.** Hepatic tissues and cuffed arteries after 4 weeks of cuff placement and control intact arteries were perfused with physiological saline and then removed. RNA was prepared with the use of TRIzol reagent (Invitrogen), and real-time PCR was performed as described previously (60). PCR primers were designed for ubiquitin-conjugating enzyme E2L3 (*Ube2l3*), forward, 5'-TCCGTAACATCCAGTTGATGAAG-3', and reverse, 5'-GGTTTGAATGGATACTCTGCTGGAA-3'; phosphatidylinositol 4-kinase, catalytic, alpha polypeptide (*Pik4ca*), forward, 5'-AACCAGCAGTGTGCCAGCATC-3', and reverse, 5'-CAGTCCCATCAGGTAGGCAAGAA-3'; synaptosomal-associated protein (*Snapt29*), forward, 5'-GACAGGCAGCATCTGAGACA-3', and reverse, 5'-GCTCATCTTCAAATCCTGATCA-3'; *Il1b*, forward, 5'-ATGGCAACTGTTCTCTGAA-3', and reverse, 5'-GTACAAAGCTCATGGAGA-3'; *Il6*, forward, 5'-TTCCCTACTTCA-CAAGTC-3', and reverse, 5'-GGTTTGCCGAGTAGATCT-3'; *Mcp1*, forward, 5'-CCAACTCTCACTGAAGCCAGTCT-3', and reverse, 5'-TTGGGATCATCTTGCTGGTGAA-3'; *Egr1*, forward, 5'-CTATGAGCACCTGAC-CACA-3', and reverse, 5'-TAGTCAGGGATCATGGGAAC-3'; *Klf5*, forward, 5'-CTCCGGAGACGATCTGAAACAC-3', and reverse, 5'-CCACTGAG-GCACTGTCTCGTCTA-3'; and  $\beta$ -actin, forward, 5'-TGGAATCCTGTGGCATCCATGAAAC-3', and reverse, 5'-TAAACGCAGCTCAGTAA-CAGTCCG-3'. Expression levels of *Ube2l3*, *Pik4ca*, *Snapt29*, *Il1b*, *Il6*, *Mcp1*, *Egr1*, and *Klf5* were expressed relative to  $\beta$ -actin signal.

**Histological and immunohistochemical studies.** The removed arterial tissues (aortic root, femoral arteries) were fixed in 10% neutral buffered formalin overnight. The aortic root and middle segment of the femoral arteries were cut into subserial 5- $\mu$ m-thick cross sections with intervals of 200  $\mu$ m. Then sections were stained with Elastica-van Gieson (femoral arteries), Masson trichrome, and PAR-1 [Thrombin R (H-111):sc-5605; Santa Cruz Biotechnology Inc.] (aortic root). Antibody distribution was visualized by the avidin-biotin complex technique and Vector Red substrate (Vector Laboratories) (61). The average from 5 sections for each animal tissue was used to determine lesion size.





**Detection of proliferative cells by BrdU and PCNA staining.** BrdU (100 mg/kg in saline) was injected intraperitoneally, and the mice were perfused 6 hours later with 10% neutral buffered formalin; then their femoral arteries were fixed in 10% neutral buffered formalin in PBS. Immunohistochemical staining was performed using a Cell Proliferation Kit (GE Healthcare) according to the instructions of the manufacturer. For immunohistochemistry, sections were fixed in 10% neutral buffered formalin for 10 minutes at room temperature, embedded in paraffin, sectioned at 2  $\mu$ m, and counterstained with hematoxylin. Labeled cells and total cells in the intima and media were counted in all sections per specimen, and labeling indices were calculated as the labeled cell/total cell ratio of intima and media.

**Human purified HCII protein administration in cuff-injured or wire-injured HCII mutant mice.** Human HCII protein was purified by a heparin affinity column from human pooled plasma as previously reported (62) and buffered with 10 mM sodium phosphate and 5% sorbitol. Because heparin was covalently bound to this column bed, no elution of heparin was observed. The purity of HCII protein measured by HPLC was more than 99%. We first measured the time course of plasma HCII activities in *HCII*<sup>-/-</sup> mice after intraperitoneal injection of human purified HCII protein (45 mg/kg) or vehicle alone. Then cuff-injured or wire-injured WT mice and 1 group of *HCII*<sup>-/-</sup> mice were injected with the vehicle intraperitoneally 3 times per week for 4 weeks, while another group of cuff-injured or wire-injured *HCII*<sup>-/-</sup> mice were injected with purified human HCII protein (45 mg/kg) intraperitoneally 3 times per week for 4 weeks. Cuff-injured or wire-injured arteries in these mice were removed and used for histological evaluation by Elastic-van Gieson staining.

**Superoxide detection and oil red O staining in the aortic root.** The aortic root and ascending aorta were excised from mice, rinsed in normal saline, and frozen in optimal cutting temperature compound (Tissue-Tek; Sakura Finetek) until use. Transverse sections (10  $\mu$ m thick) were cut with a cryostat and placed on silane-coated glass slides. Oil red O staining was performed as described previously (63). The slide glasses were incubated with DHE in PBS (10  $\mu$ M/l) in a dark, humidified container at room temperature for 30 minutes. DHE is oxidized on reaction with superoxide to ethidium bromide, which binds to DNA in the nucleus and fluoresces red. After placement of cover glass, aortic tissues were observed using a laser scanning confocal microscope (Nikon Digital Eclipse CFI-Ready). The excitation wavelength was 488 nm, and emission fluorescence was detected with a 568-nm long-pass filter.

**Measurement of urinary 8OHdG excretion.** To evaluate urinary excretion of 8OHdG, as an oxidative DNA damage marker, mice were individually

housed in metabolic cages that provided free access to tap water and food for 24 hours, and collected urine was analyzed using a commercially available ELISA assay kit (Nikken SEIL Corp.). The amount of urinary 8OHdG excretion was corrected to the volume of urine in a given day and body weight (g).

**Statistics.** Values for parameters within a group are expressed as mean  $\pm$  SEM. For comparisons among groups, statistical significance was assessed using an unpaired 2-tailed Student's *t* test or 1-way ANOVA, and the significance of each difference was determined by post-hoc testing using the Tukey-Kramer method. These analyses were performed on an Apple Macintosh computer with the use of Excel (Microsoft) and StatView statistical package (StatView 5.0; SAS Institute Inc.). A *P* value of less than 0.05 was considered significant.

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