Deficiency of cathepsin S reduces atherosclerosis in LDL receptor–deficient mice

Citation

Published Version
doi:10.1172/JCI14915

Permanent link
http://nrs.harvard.edu/urn-3:HUL.InstRepos:13506491

Terms of Use
This article was downloaded from Harvard University’s DASH repository, and is made available under the terms and conditions applicable to Other Posted Material, as set forth at http://nrs.harvard.edu/urn-3:HUL.InstRepos:dash.current.terms-of-use#LAA

Share Your Story
The Harvard community has made this article openly available. Please share how this access benefits you. Submit a story.

Accessibility
Deficiency of cathepsin S reduces atherosclerosis in LDL receptor–deficient mice

Galina K. Sukhova,1 Yaou Zhang,2 Jie-Hong Pan,2 Youichiro Wada,3 Takashi Yamamoto,3 Makoto Naito,3 Tatsuhiko Kodama,3 Sotirios Tsimikas,4 Joseph L. Witztum,4 Michael L. Lu,1 Yasuhiko Sakara,1 Michael T. Chin,1 Peter Libby,1 and Guo-Ping Shi2

1The Leducq Center for Cardiovascular Research, Department of Medicine, Brigham and Women’s Hospital and Harvard Medical School, Boston, Massachusetts, USA
2Department of Medicine, University of California, San Francisco, San Francisco, California, USA
3Department of Molecular Biology and Medicine, University of Tokyo, Tokyo, Japan
4Department of Medicine, University of California at San Diego, La Jolla, California, USA

Human atherosclerotic lesions overexpress the lysosomal cysteine protease cathepsin S (Cat S), one of the most potent mammalian elastases known. In contrast, atheromata have low levels of the endogenous Cat S inhibitor cystatin C compared with normal arteries, suggesting involvement of this protease in atherogenesis. The present study tested this hypothesis directly by crossing Cat S–deficient (CatS–/–) mice with LDL receptor–deficient (LDLR–/–) mice that develop atherosclerosis on a high-cholesterol diet. Compared with LDLR–/– mice, double-knockout mice (CatS–/– LDLR–/–) developed significantly less atherosclerosis, as indicated by plaque size (plaque area and intimal thickening) and stage of development. These mice also had markedly reduced content of intimal macrophages, lipids, smooth muscle cells, collagen, CD4+ T lymphocytes, and levels of IFN-γ. CatS–/– LDLR–/– monocytes showed impaired subendothelial basement membrane transmigration, and aortas from CatS–/– LDLR–/– mice had preserved elastic laminae. These findings establish a pivotal role for Cat S in atherogenesis.


Introduction

 ECM remodeling contributes to all stages of atherosclerosis. Proteolysis of ECM molecules probably participates in the initial mononuclear leukocyte translocation from the vascular lumen through the basement membrane into subendothelium, traversal of medial smooth muscle cells (SMCs) through the elastic lamina into the intima, compensatory arterial enlargement, plaque angiogenesis, and plaque disruption (1–3).

 Previous studies have suggested involvement of both serine proteases and MMPs in arterial diseases. Deficiency of plasminogen reduced the development of lesions in transplanted mouse arteries (4) and in arteries of electrically injured mice (5). Hyperlipidemic mice deficient in urokinase showed reduced aortic ectasia (6), an effect attributed to reduced plasmin-dependent activation of pro-MMPs (6).

 Human atheromata overexpress the interstitial collagenases MMP-1 (7), MMP-13 (8), and MMP-8 (9) at sites of interstitial collagen cleavage. MMPs 2, 3, 7, 9, 12, 14, and others also localize in human atherosclerotic plaques (7, 10–14). Overexpression of the MMP inhibitor TIMP-1 reduced formation of aneurysms (15) and atherosclerotic lesions (16) in animals, affirming a role for MMPs in arterial remodeling and atherogenesis.

 Beyond serine proteases and MMPs, we recently proposed the participation of cathepsin S (Cat S) and Cat K in atherogenesis. These cysteine proteases have potent elastolytic (17, 18) and collagenolytic activities (19). Diseased human arteries overexpress these enzymes and exhibit a reciprocal deficiency of cystatin C, the most abundant endogenous inhibitor of cysteine proteases (1, 2). Biochemical studies suggest that Cat S and Cat K account for most of the elastolytic activity extracted from human carotid plaques. Multiple cell types found in human atheroma can express Cat S. We have previously demonstrated that intimal macrophages as well as SMCs in the tunica media express Cat S in human atherosclerotic arteries (1). We also demonstrated colocalization of Cat S with luminal and microvascular endothelial cells (ECs) in human atheroma and with microvascular ECs in wounded mouse skin (G.-P. Shi et al., unpublished data). In vitro experiments showed secretion of active Cat S by these atheroma-associated cell types following exposure to inflammatory cytokines found in atheroma. Our findings implicate Cat S in atherogenesis along with serine proteases and MMPs. Furthermore, Cat S, an essential protease acquired for invariant chain processing, plays an important role in antigen presentation. Cat S deficiency leads to impaired
immune responses (20, 21), which might also affect atherogenesis (22, 23). Despite this plethora of observational and biochemical data, no direct evidence establishes a role of Cat S in atherogenesis. This study used Cat S–deficient (CatS–/–) mice crossbred with atherosclerosis-prone (LDLR–/–) mice to address this question directly.

**Methods**

**CatS–/– LDLR–/– mouse generation and animal protocol.** CatS–/– mice were backcrossed more than ten generations onto the C57/BL6 background. These mice were crossbred with LDLR–/– mice (also on the C57/BL6 background; The Jackson Laboratory, Bar Harbor, Maine, USA). The resulting double knockout mouse genotypes were confirmed using mouse tail genomic DNA–derived PCR as described (20, 24). This study used both CatS–/– LDLR–/– and control LDLR–/– mice. Mice consumed an atherogenic diet (D12108; Research Diets Inc., New Brunswick, New Jersey, USA) containing 20.1% saturated fat, 1.37% cholesterol, and 0% sodium cholate (25). After 8, 12, and 26 weeks (n = 12/timepoint/group), mice were used as described below.

**Oil red O staining for lipids.** Deposition of lipids in en face preparations of abdominal aortas (fixed with 10% formalin) was determined by oil red O staining. Subsequently, the aortas were opened longitudinally to the iliac bifurcation and pinned out on the surface of black formalin (26). Formalin-fixed frozen sections were incubated for 4 hours in a freshly prepared 0.1% solution of Sirius Red F3BA (Polysciences Inc., Warington, Pennsylvania, USA) in saturated aqueous picric acid. After rinsing twice in 0.01 N HCl and distilled water, sections were briefly dehydrated and mounted in Permount (Vector Laboratories). Sirius red staining was analyzed by polarization microscopy.

**Mouse aortic SMC isolation and elastase assay.** Mouse aortic SMCs were isolated as described with modifications (26). Briefly, mouse aortas were removed and separated from fat and adventitia followed by digestion with 1x collagenase type II (Worthington Biochemical Corp., Lakewood, New Jersey, USA) at 37°C for 10–15 minutes. The adventitia was completely stripped under microscopic visualization, the aortas were opened, and ECs were removed by abrasion. The aortas were then chopped and digested in 4 ml of type I collagenase (1 mg/ml; Worthington Biochemical Corp.) and elastase III (0.125 mg/ml; Sigma-Aldrich) for 40–90 minutes at 37°C. Digestion was stopped by addition of 10 ml of DMEM containing 10% FCS. Cells were then plated on a fibronectin-coated 60-mm culture dish. After the first two passages, cells were cultured on gelatin-coated culture dishes. Cell identity was verified by immunostaining for α-smooth muscle actin and calponin. To assay SMC elastase activity, SMCs were passaged onto 24-well plates (1 × 10^5/well). After the cells reached confluence, 300 μg of [3H]elastin was added to each well followed by 24–48 hours of incubation in the presence or absence of IFN-γ (500 U/ml). Culture medium was collected by centrifugation (14,000 g for 15 minutes) and soluble radioactive elastin was quantified as described (1).

**Statistical analysis.** To measure plaque size as well as the percent-positive area for staining, longitudinal sections of the aortic arch were analyzed microscopically for all mice. As described previously (27), a 3-mm segment of the lesser curvature of the aortic arch was defined proximally by a perpendicular axis dropped from the right side of the innominate artery. The percentage of the total area of the aortic arch stained for SMCs, macrophages, collagen, and lipids was determined using computer-assisted image analysis (Image-Pro Plus; Media Cybernetics, Silver Springers, Maryland, USA).

We estimated atherosclerosis as well as immunohistochemical staining for MCP-1, VCAM-1, and ICAM-1 by semiquantitative grading using the following criteria: 0, no staining; 1, weak staining; 2, moderate staining; 3, strong staining. Atherosclerotic lesions were graded as: 0, no lesion; 1, few macrophages underlying endothelium; 2, fatty streak–like lesion; and three grades (3 through 5) of atheroma, with the lower grades corresponding to less-developed lesions. Scoring was performed independently by two blinded observers. Data for CatS–/– LDLR–/– and control LDLR–/– groups

---

**Immunohistochemistry.** Serial cryostat sections (6 μm) of mouse aortic arches were fixed in acetone, air dried, and stained by the avidin-biotin-peroxidase method. After limiting endogenous peroxidase activity with 0.3% H2O2 and nonspecific binding of primary antibody with 5% species-appropriate normal serum (Vector Laboratories, Burlingame, California, USA), we incubated sections with primary antibodies diluted in PBS supplemented with 5% species-appropriate normal serum for 90 minutes at room temperature. Incubation with secondary antibodies for 45 minutes was followed by avidin–biotin complex (ABC; Vector Laboratories, Burlingame, California, USA), we incubated sections with primary antibodies diluted in PBS supplemented with 5% species-appropriate normal serum (Vector Laboratories, Burlingame, California, USA) followed by counterstaining with Gill’s hematoxylin solution (Sigma-Aldrich, St. Louis, Missouri, USA). Antibodies used were: rabbit anti–mouse Cat S (1:90; ref. 20), goat anti-smooth muscle α-actin (1:75; Santa Cruz Biotechnology Inc., Santa Cruz, California, USA), rat anti–mouse macrophage (Mac-3, 1:1,000; Pharmingen, San Diego, California, USA), and rat anti–mouse T cell (CD4, 1:100; Pharmingen), goat anti–rat IFN-γ (1:75; R&D Systems Inc., Minneapolis, Minnesota, USA), hamster anti–mouse monocyte chemoattractant protein-1 (MCP-1) (1:50; Pharmingen), biotinylated anti–mouse VCAM-1 (1:60), and biotinylated anti–mouse ICAM-1 (1:50; Pharmingen).
were presented as mean ± SD and compared by the nonparametric Mann-Whitney test. A value of \( P \leq 0.05 \) was considered significant.

**Lipoprotein measurement.** Blood samples were collected by retro-orbital venous plexus puncture. Serum total cholesterol and triglyceride were assayed by enzymatic methods using Sigma Diagnostics kit nos. 401 and 343 (Sigma-Aldrich), respectively, adapted for microtiter plate assay. HDL was separated from LDL and VLDL cholesterol with kit no. 352-4, also adapted for microtiter plate assay. HDL cholesterol levels were then measured using kit no. 401. LDL cholesterol was calculated as follows: serum LDL cholesterol concentration (mg/dl) = total cholesterol – HDL cholesterol – (triglycerides/5).

**Mouse monocyte isolation, transmigration measurement, and invasion.** Monocytes from CatS\(^{-/-}\) LDLR\(^{-/-}\) and LDLR\(^{-/-}\) mice were isolated by Percoll gradient centrifugation. Briefly, mouse blood was collected into heparin (1:200; Sigma-Aldrich) followed by 1:1 dilution with 1× PBS. Diluted blood was laid over a Percoll gradient (1.04 g/ml; Amersham Biosciences Corp., Piscataway, New Jersey, USA) and centrifuged for 30 minutes at 900 g. The interface layer of monocytes/lymphocytes was collected and cultured on a Petri dish for 2 hours. Unattached lymphocytes were removed by washing with RPMI and monocytes were further cultured in RPMI.

Rabbit aortic SMCs and ECs were cocultured as described (28). Briefly, 1 × 10\(^6\) rabbit aortic SMCs were cultivated with EGM-2 media (containing 2% rabbit plasma) on the filter for more than 3 weeks. The mixture of type I and type IV collagen was then laid on the filter of a Chemotaxicell chamber (Kurabo Inc., Osaka, Japan) and incubated at 37°C for 1 hour to complete gel formation. Then 2 × 10\(^5\) rabbit aortic EC’s were seeded on top of the gel; these became confluent in a week. Monocytes (1 × 10\(^6\)) isolated from peripheral blood of CatS\(^{-/-}\) LDLR\(^{-/-}\) or LDLR\(^{-/-}\) mice were applied to the EC layer (day 0). After 7 days of culture, samples were fixed with 10% formalin, embedded in paraffin, and sectioned (10 µm) for immunohistochemical study. Sections were incubated with primary antibodies: F4/80 (anti–mouse monocyte, ref. 29) or anti–macrophage scavenger receptor class A, type I (anti-MSR) (30) followed by secondary antibody conjugated with HRP (Amersham Biosciences Corp.). After visualization with 3,3’-diaminobenzidine (Dojin Chemical Co., Kumamoto, Japan), nuclei were counterstained with hematoxylin. The numbers of monocytes/macrophages attached to ECs and those that had transmigrated into the matrix layer were counted by two blinded observers using confocal microscopy (TCS SP; Leica Microsystems GmbH, Heidelberg, Germany) as described previously (28). Results are presented as mean ± SD. Statistical comparisons were made using the two-tailed Student paired t test; a value of \( P \leq 0.01 \) was considered significant. The experiment was repeated four times.

For the mouse monocyte invasion/transmigration assay, cells were incubated overnight at room temperature in PBS in Boyden transwell chambers coated with type I collagen (110 µg/ml; Celtrix Pharmaceuticals Inc., Santa Clara, California, USA) and type IV collagen (30 µg/ml; BD Biosciences, Bedford, Massachusetts, USA). Briefly, monocytes were resuspended in serum-free media at 1 × 10\(^5\)/ml, and 0.5 ml of suspension was added to each well. The lower chamber was supplied with 0.8 ml of complete media with 0.5 µg of mouse recombinant MCP-1 (Pharmingen). The invasion assay was carried out at 37°C for 16 hours followed by crystal violet staining. Six microscopic fields were randomly chosen to count transmigrated cells.

### Table 1

**Plasma lipid contents in mice fed with chow or high cholesterol diet**

<table>
<thead>
<tr>
<th>Genotype</th>
<th>Diet</th>
<th>n</th>
<th>Plasma total cholesterol (mg/dl)</th>
<th>Plasma total triglyceride (mg/dl)</th>
<th>Plasma total HDL (mg/dl)</th>
<th>Plasma total LDL (mg/dl)</th>
</tr>
</thead>
<tbody>
<tr>
<td>WT</td>
<td>Chow</td>
<td>12</td>
<td>171.5 ± 50.0</td>
<td>113.0 ± 42.3</td>
<td>55.6 ± 5.6</td>
<td>93.9 ± 24.0</td>
</tr>
<tr>
<td>CatS(^{-/-})</td>
<td>Chow</td>
<td>6</td>
<td>291.1 ± 44.7</td>
<td>112.8 ± 26.8</td>
<td>46.3 ± 18.7</td>
<td>122.3 ± 52.4</td>
</tr>
<tr>
<td>LDLR(^{-/-})</td>
<td>Chow</td>
<td>6</td>
<td>237.2 ± 28.2(^{a})</td>
<td>116.3 ± 28.0</td>
<td>49.6 ± 18.8</td>
<td>164.4 ± 23.4(^{a})</td>
</tr>
<tr>
<td>WT</td>
<td>Chow</td>
<td>8</td>
<td>200.1 ± 43.3</td>
<td>115.2 ± 28.7</td>
<td>52.4 ± 8.9</td>
<td>124.6 ± 33.3</td>
</tr>
<tr>
<td>CatS(^{-/-})</td>
<td>Chow</td>
<td>8</td>
<td>215.2 ± 31.3</td>
<td>120.1 ± 25.3</td>
<td>62.3 ± 15.9</td>
<td>129.0 ± 25.7</td>
</tr>
<tr>
<td>LDLR(^{-/-})</td>
<td>Chow</td>
<td>6</td>
<td>1,519.7 ± 124.0</td>
<td>207.2 ± 38.1</td>
<td>49.1 ± 27.9</td>
<td>1,429.1 ± 149.8</td>
</tr>
<tr>
<td>CatS(^{-/-})/LDLR(^{-/-})</td>
<td>Chow (8 weeks)</td>
<td>8</td>
<td>1,348.8 ± 236.9</td>
<td>162.8 ± 82.6</td>
<td>50.4 ± 22.7</td>
<td>1,265.8 ± 233.9</td>
</tr>
<tr>
<td>WT</td>
<td>Chow</td>
<td>8</td>
<td>221.6 ± 52.7</td>
<td>118.0 ± 23.7</td>
<td>55.5 ± 8.7</td>
<td>142.5 ± 30.0</td>
</tr>
<tr>
<td>CatS(^{-/-})</td>
<td>Chow</td>
<td>8</td>
<td>235.1 ± 30.5</td>
<td>112.7 ± 24.2</td>
<td>57.6 ± 28.3</td>
<td>155.0 ± 33.1</td>
</tr>
<tr>
<td>LDLR(^{-/-})</td>
<td>Chow (12 weeks)</td>
<td>12</td>
<td>1,355.2 ± 340.1</td>
<td>204.8 ± 60.3</td>
<td>70.2 ± 15.0</td>
<td>1,103.2 ± 396.2</td>
</tr>
<tr>
<td>WT</td>
<td>Chow</td>
<td>7</td>
<td>195.6 ± 35.7</td>
<td>121.5 ± 28.8</td>
<td>58.7 ± 13.2</td>
<td>112.5 ± 32.4</td>
</tr>
<tr>
<td>CatS(^{-/-})</td>
<td>Chow</td>
<td>7</td>
<td>180.5 ± 43.2</td>
<td>119.1 ± 31.5</td>
<td>62.0 ± 22.7</td>
<td>94.7 ± 24.2</td>
</tr>
<tr>
<td>LDLR(^{-/-})</td>
<td>Chow (26 weeks)</td>
<td>11</td>
<td>851.7 ± 162.1</td>
<td>NA</td>
<td>54.1 ± 5.0</td>
<td>770.4 ± 198.8</td>
</tr>
<tr>
<td>CatS(^{-/-})/LDLR(^{-/-})</td>
<td>Chow (26 weeks)</td>
<td>11</td>
<td>949.1 ± 210.9</td>
<td>NA</td>
<td>57.8 ± 8.6</td>
<td>862.8 ± 170.5</td>
</tr>
</tbody>
</table>

\(^{a}\)P < 0.04, CatS\(^{-/-}\)/LDLR\(^{-/-}\) mice fed with chow diet vs. LDLR\(^{-/-}\) mice fed with chow diet. NA, not assayed.
Results

Generation of CatS−/− LDLR−/− mice. LDLR−/− mice develop atherosclerotic plaques when consuming a high-cholesterol diet and provide a well-established model of atherosclerosis (31).

To study the role of Cat S in atherogenesis, we generated double-knockout (CatS−/−LDLR−/−) mice by crossing CatS−/− mice with LDLR−/− mice, both on the C57/BL6 background. PCR analysis of tail genomic DNA confirmed the presence of mutant alleles for both Cat S (170 bp) (20) and LDLR (800 bp) (24), as well as the absence of their corresponding wild-type alleles (data not shown). Both CatS−/− and CatS−/−LDLR−/− mice were apparently healthy and fertile, and exhibited mendelian transmission of the mutant alleles. We noticed no obvious abnormality in vasculature, organ development, gestation time, or food/water consumption.

Cat S deficiency does not affect serum lipids. In agreement with earlier findings, LDLR deficiency significantly increased plasma total cholesterol and LDL levels but did not change either HDL or triglyceride levels (24). Disruption of Cat s gene function alone did not affect plasma total cholesterol, triglyceride, HDL, or LDL levels in mice consuming an atherogenic diet (Table 1). CatS−/−LDLR−/− mice that consumed either a chow or atherogenic diet showed no significant differences in plasma total cholesterol and total LDL levels compared with similarly fed LDLR−/− mice (Table 1).

Cat S deficiency reduces atherosclerosis in LDLR−/− mice. LDLR−/− mice fed an atherogenic diet developed atherosclerotic lesions with morphological characteristics similar in many aspects to those seen in humans (31). To evaluate the severity of atherosclerosis in mice, we characterized lesions by (a) lesion area in the abdominal aorta delineated by lipid staining with oil red O (Figure 1), (b) quantitative morphometric measurement of intimal thickening (area between internal elastic lamina and lumen) in longitudinal sections of aortic arches (intimal area; Figure 2a), and (c) histological grading of lesions in aortic arches (see details in Methods) (Figure 2b). Analysis of en-face preparations of abdominal aorta stained with oil red O showed a statistically significant reduction of plaque area in CatS−/−LDLR−/− mice after 12 weeks (60% reduction, P < 0.0001) and 26 weeks (30% reduction, P < 0.005) on an atherogenic diet (Figure 1). Consistent with the observations on the abdominal aorta, the aortic arches from CatS−/−LDLR−/− mice also showed a statistically significant reduction in plaque size (intimal area; Figure 2a) as well as stage of plaque development (Figure 2b) compared with LDLR−/− mice at all three timepoints tested. In comparison, Cat S single-deficient mice consuming the high-cholesterol diet
Preserved elastin in CatS<sup>−/−</sup> LDLR<sup>−/−</sup> mice. (a) Verhoeff–van Gieson staining for elastin shows aortic arch elastic laminae. The graph shows the number of discontinuities in the elastic lamina enumerated in the 3-mm aortic arch sections. Significant differences in elastin preservation between CatS<sup>−/−</sup> LDLR<sup>−/−</sup> and LDLR<sup>−/−</sup> mice occurred at both the 12-week and 26-week timepoints. (b) Cat S–positive medial SMCs colocalized with sites of elastin degradation in LDLR<sup>−/−</sup> mice. Serial sections of aortic arches were stained for α-actin (top left) and Cat S (bottom left). Verhoeff–van Gieson staining for elastin performed after Cat S staining showed Cat S expression to be colocalized with elastin fragmentation; arrows indicate elastin breaks (right). (c) SMC elastase assay. IFN-γ-stimulated SMCs from CatS<sup>−/−</sup> mouse aortas showed significantly weaker elastase activity than was found in wild-type cells (P < 0.0001). Three independent experiments yielded similar observations. (d and e) Reduction of intimal SMCs, collagen content, and fibrous cap thickness in CatS<sup>−/−</sup> LDLR<sup>−/−</sup> mice. Longitudinal sections of aortic arch were immunostained for α-actin (SMC content) and for collagen content with picrosirius red. CatS<sup>−/−</sup> LDLR<sup>−/−</sup> mice had significantly less SMC and collagen in intima than did LDLR<sup>−/−</sup> mice. Percentages of positively stained areas were calculated using computer-assisted image quantification as described (27). Values represent mean ± SD. P values ≤ 0.05 were considered significant. (f) The thickness of the fibrous cap of an aortic atheroma of each mouse was measured using sections stained with oil red O as described in Methods. Few if any fibrous caps formed at earlier timepoints.

Reduced SMC and interstitial collagen content of intimal lesions in CatS<sup>−/−</sup> LDLR<sup>−/−</sup> mice. SMC migration from the tunica media into the intima requires proteolytic degradation of elastin. The early lesions in aortas of CatS<sup>−/−</sup> LDLR<sup>−/−</sup> mice contained significantly fewer SMCs than those in LDLR<sup>−/−</sup> mice, measured as the percent of α-actin–positive area at 12 weeks (P < 0.05) and 26 weeks (P < 0.003) (Figure 3, d and e). Picrosirius red staining of the aortic arches revealed a significant reduction in collagen content in CatS<sup>−/−</sup> LDLR<sup>−/−</sup> mice compared with LDLR<sup>−/−</sup> control mice at the 12-week timepoint (P < 0.01) but not at the 26-week timepoint (Figure 3e). Atheromata in CatS<sup>−/−</sup> LDLR<sup>−/−</sup> mice had significantly thinner fibrous caps at 26 weeks (Figure 3, d and f, P < 0.001). The fibrous caps were insufficiently developed at earlier timepoints to permit accurate measurement. Reduction of both the collagen content and the fibrous cap thickness may reflect the decreased number of SMCs (the major source of collagen in arteries) due to impaired SMC migration across the internal elastic membrane that separates the tunica media from the intima.

Cat S deficiency decreases macrophage and leukocyte accumulation in aortic lesions. Macrophages sustain inflammation within atheroma and probably influence plaque stability. Human atheroma that have ruptured and caused fatal thrombosis typically have abundant macrophages (32). The percentage of plaque area...
immunoreactive Cat S (data not shown), suggesting that Cat S might also influence T lymphocyte trafficking during atherogenesis.

Cat S deficiency impairs leukocyte transmigration through subendothelial basement membrane. We first tested the possibility that reduced leukocyte accumulation in CatS–/– LDLR–/– mice resulted from decreased expression of adhesion and chemotactic molecules. However, we detected no difference in the expression of VCAM-1, ICAM-1, or MCP-1 at the early timepoint (8 weeks) most important for white blood cell recruitment (Table 2) (35). After 12 weeks and 26 weeks on an atherogenic diet when atheromatous lesions were rather well developed, lesions did show a statistically significant decrease in immunoreactive VCAM-1 (P < 0.02 for both the 12-week and 26-week timepoints) or MCP-1 (P < 0.004 and P < 0.006 for 12-week and 26-week timepoints, respectively). In addition to reduced local inflammatory activation, these changes may reflect decreased atheroma formation (36). ICAM-1 protein content did not differ among groups at any time studied.

Alternatively, Cat S might participate in leukocyte migration from peripheral blood by facilitating proteolytic degradation of subendothelial vascular basement membrane. The nascent atheroma in LDLR–/– mice (Figure 2, grade 1 and grade 2) had Cat S–positive macrophages underlying the endothelium (Figure 5a). We previously showed a similar pattern of Cat S expression in human fatty streaks (1). We tested this hypothesis in vitro in a monocyte migration and differentiation assay described previously (28). This assay measures migration of peripheral blood monocytes isolated from CatS–/– LDLR–/– and control LDLR–/– mice through a monolayer of ECs cultured atop a mixture of type I and type IV collagens, prominent components of the subendothelial basement membrane (see details in Methods).

The formation of an EC monolayer as well as the deposition of a collagen I and collagen IV–containing basement membrane has been previously demonstrated (28). After 7 days of incubation, 91% of the F4/80-positive monocytes/macrophages (29) from CatS–/– LDLR–/– mice remained in the upper chamber, and only 9% of these cells transmigrated through the endothelial monolayer and the subjacent collagen-containing substrate (Figure 5, d and f). In contrast, over 50% of LDLR–/– monocytes/macrophages crossed into the SMC layer (Figure 5,
26 weeks
LDLR–/–
NS, no significant difference.

Grading of adhesion molecule expression in atherosclerotic lesions
Table 2

<table>
<thead>
<tr>
<th>Western diet</th>
<th>Mouse genotype</th>
<th>MCP-1</th>
<th>VCAM-1</th>
<th>ICAM-1</th>
</tr>
</thead>
<tbody>
<tr>
<td>8 weeks</td>
<td>LDLR–/–</td>
<td>0.79 ± 0.88</td>
<td>1.37 ± 0.38</td>
<td>1.75 ± 0.69</td>
</tr>
<tr>
<td></td>
<td>CatS–/– LDLR–/–</td>
<td>1.00 ± 0.71</td>
<td>1.00 ± 0.77</td>
<td>1.33 ± 0.26</td>
</tr>
<tr>
<td></td>
<td></td>
<td>NS</td>
<td>NS</td>
<td>NS</td>
</tr>
<tr>
<td>12 weeks</td>
<td>LDLR–/–</td>
<td>2.33 ± 0.61</td>
<td>2.75 ± 0.42</td>
<td>1.50 ± 0.71</td>
</tr>
<tr>
<td></td>
<td>CatS–/– LDLR–/–</td>
<td>0.29 ± 0.25</td>
<td>1.96 ± 0.56</td>
<td>1.17 ± 0.41</td>
</tr>
<tr>
<td></td>
<td></td>
<td><em>P</em> &lt; 0.004</td>
<td><em>P</em> &lt; 0.02</td>
<td>NS</td>
</tr>
<tr>
<td>26 weeks</td>
<td>LDLR–/–</td>
<td>2.25 ± 0.69</td>
<td>3.13 ± 0.44</td>
<td>0.37 ± 0.34</td>
</tr>
<tr>
<td></td>
<td>CatS–/– LDLR–/–</td>
<td>0.50 ± 0.52</td>
<td>2.37 ± 0.38</td>
<td>0.83 ± 0.74</td>
</tr>
<tr>
<td></td>
<td></td>
<td><em>P</em> &lt; 0.006</td>
<td><em>P</em> &lt; 0.02</td>
<td>NS</td>
</tr>
</tbody>
</table>

NS, no significant difference.

Discussion
Recent observations have implicated the elastolytic cathepsins S and K in atherosclerosis (1, 2). The present study sought direct evidence for Cat S involvement in atherogenesis using genetically modified mice. These experiments showed in vivo that lack of Cat S reduced atherogenesis in LDLR–/– mice by more than 50% after 8 or 12 weeks of consumption of an atherogenic diet, and mice still showed significantly diminished atheroma (∼30% reduction) at 26 weeks (Figure 1 and Figure 2). This attenuation of atherosclerosis in CatS–/– LDLR–/– mice likely results primarily from the lack of Cat S. We found no difference in other cysteine proteases (Cat B and Cat L, examined by cysteine protease active site [125I]-JPM labeling) nor in the expression of MMPs (MMP-2, MMP-3, or MMP-9 examined by Western blot analysis) in CatS–/– LDLR–/– compared with LDLR–/– macrophages (data not shown). Nor did deficiency of Cat S affect serum lipid levels or expression of other cathepsins or MMPs by ECs (G.-P. Shi et al., unpublished data). The striking magnitude and duration of this blunted lesion formation highlights the importance of understanding the mechanism(s) of Cat S involvement in atherogenesis.

Although not conclusive, our results suggest several possible mechanisms by which Cat S may contribute to lesion formation. First, Cat S, one of the most potent mammalian elastases (40), remains active at neutral pH (18) and may participate in proteolysis of elastic laminae in arteries (1, 2). We previously demonstrated in vitro that SMCs activated by IFN-γ as well as differentiated macrophages can express and secrete active Cat S and degrade extracellular elastin. Either natural (cystatin C) or synthetic inhibitors of Cat S can substantially block this elastolysis (1, 2, 41, 42). Within the artery wall, elastin fibers contribute to the vessel’s structural integrity and regulate SMC migration and proliferation (43). Mice with targeted disruption of the elastin gene die 2–3 days after birth due to malformed blood vessels (44). Our present observations indicate that degradation or remodeling of elastic laminae contributes to generation of atherosclerotic lesions.

Figure 5
Cat S deficiency impairs leukocyte transmigration. (a) Increased local expression of Cat S by ECs and macrophages abutting the arterial basement membrane. Paraffin sections of LDLR–/– mouse fatty streaks were stained with rabbit anti–mouse Cat S antibodies (red color) and non-immune rabbit IgG (b). (c–e) In vitro monocyte transmigration. Cat S deficiency inhibited transmigration of monocytes/macrophages through the EC monolayer and collagen-containing film prepared as described in Methods. Transmigrated macrophages were detected by the monocyte/macrophage marker F4/80 (arrows) (c and d) and macrophage scavenger receptor (MSR) antibodies (arrowheads) (e). Asterisks and Fs indicate collagen matrix and microporous filter, respectively. (f) The bar graph shows percentage of monocytes/macrophages remaining on the EC layer (white bars) and the percentage that transmigrated into the collagen-containing gel (black bars). Six independent experiments showed similar results.
and development of atheromatous lesions, indicating a critical role for elastin preservation in this postnatally acquired disease as well. Indeed, in this study, CatS had the greatest overall effect on lesion development at 12 weeks, when CatS–/– LDLR–/– mice still showed better preserved elastic laminae in aortic arches (Figure 3a) and abdominal aortas (data not shown) than in LDLR–/– mice. Similar to our earlier findings in human atheromata (1), medial SMCs in LDLR–/– mice also express abundant Cat S at the sites of elastic lamina degradation (Figure 3b), suggesting a role for Cat S in this process. Primary cultured CatS–/– SMCs stimulated with IFN-γ showed significantly lower elastolysis than did SMCs from control mice (Figure 3c). Reduced elastin breakdown may limit SMC entry into the intima across the elastin barriers in CatS–/– LDLR–/– mice. The thinner fibrous cap and reduced intimal collagen content may follow decreased SMC accumulation in lesions, as these cells produce most of the collagen in atherosclerotic lesions (43).

A second mechanism of Cat S function may involve remodeling of the basal lamina or basement membrane of the artery’s intimal layer. Immunostaining for Cat S showed strong signals in the basement membrane regions of human fatty streaks (1). We observed a similar pattern of Cat S expression in fatty streaks of LDLR–/– mice (Figure 3a). To enter the intima during early atherogenesis, leukocytes must penetrate the arterial basement membrane, which contains type IV and type I collagen, laminin, and fibronectin (46). This study monitored monocyte transmigration in vitro through layers of SMCs, collagen matrix, and ECs. Scanning EM analysis demonstrated monocyte traversal through both the EC monolayer and the underlying collagen matrix (28). The present experiments suggest that Cat S may pave the way for leukocyte entry by breaking down the subendothelial basement membrane. Monocytes isolated from CatS–/– LDLR–/– mice did not transmigrate the endothelial monolayer overlying a layer of type I and type IV collagen (Figure 5, d and f), suggesting a role for Cat S in blood-borne leukocyte penetration through the arterial basement membrane after endothelial adhesion. Although we currently cannot define a clear mechanism for this process, Cat S may play multiple roles including ECM degradation, monocyte differentiation, and possibly even “reverse transmigration.” Primary cultured monocytes from CatS–/– and control mice did not show significantly different invasion/transmigration in a collagen-coated Boyden chamber, suggesting that Cat S–mediated collagenolysis may not be the only factor affecting monocyte basement membrane transmigration. Monocyte immunostaining with F4/80, a marker for both monocytes and macrophages, indicated that most of the F4/80-positive cells from CatS–/– LDLR–/– mice are much smaller in size (and are possibly monocytes) (Figure 5d) than those (possibly macrophages) from control mice (Figure 5, c and e). Therefore Cat S may also affect monocyte differentiation in this context, although we lack direct evidence on this point. During atherogenesis, monocytes also traverse endothelium in the basal-to-apical direction (reverse transmigration) (47). Randolph et al. have shown that endogenous signals can stimulate monocyte traversal across an endothelial monolayer atop collagen matrix and that a significant fraction of these cells becomes dendritic cells after reverse transmigration (48). Cat S may thus affect monocyte reverse transmigration, although this issue requires further investigation.

Our observations did not suggest a role for Cat S in leukocyte recruitment as none of the adhesion or chemotactic molecules measured in lesions showed differences between the two groups of mice (Table 2, 8-week timepoint). Impaired leukocyte transmigration at the early stage of lesion development and/or reduced levels of cell adhesion molecule (i.e., VCAM) and chemotactic agent (i.e., MCP-1) in lesions at later stages in CatS–/– LDLR–/– mice (Table 2) may account for the reduced number of lesional CD4+ T cells and macrophages and the reduced amount of IFN-γ (Figure 4). In particular, MCP-1, which was most affected by Cat S deficiency, plays a critical role in monocyte-macrophage recruitment and migration (49). However, the monocyte recruitment/transmigration defect observed in CatS–/– monocytes (Figure 5) may not be the major cause of the retarded atherosclerosis observed in this study. Indeed, MCP-1 levels declined strikingly at both the 12-week and 26-week timepoints (Table 2), in parallel with diminished lesion macrophage content (Figure 4, a and b). Deeper understanding of the importance of MCP-1 participation in this setting may require further experiments such as bone marrow transplantation.

Cat S may also contribute to atherogenesis by effects on formation and growth of neovessels. Microvessels may influence plaque growth and lesion stability (50). Administration of the angiogenic factor VEGF was shown to significantly augment the experimental atheroma (51). In contrast, atherosclerotic animals treated with inhibitors of angiogenesis showed not only reduced intimal neovascularization but also reduced plaque size (52). Our recent observations suggested that Cat S may contribute to the degradation of arterial ECM, which provides a path for neovessel growth (53). Cat S deficiency reduces microtubule formation in vitro and microvessel growth in vivo (G.-P. Shi et al., unpublished data). Therefore, the impaired microvessel growth observed in CatS–/– mice might limit atherosclerotic plaque growth.

In addition, we and others have previously reported that Cat S plays an important role in invariant chain processing in antigen-presenting cells, which leads to altered antibody class switching and production (20, 21). Increased serum autoantibodies against oxidized LDL correlate with atherosclerosis in humans and animals (54, 55). Cat S deficiency may cause impaired immune responses and T lymphocyte activation and expansion. Indeed, the number of T lymphocytes also decreased markedly in lesions of CatS–/– LDLR–/– mice. Further, CD1 molecules mediate presentation of lipid antigen to T cells and further T cell activation by lipid-laden macrophages in...
human atheroma. CD1\* foamy macrophages congregate with T cells in atherosclerotic plaques (56). CatS\(+/–\) mice have defective CD1-restricted antigen presentation (57), consistent with a role for this protease in antigen presentation by CD1\* macrophages. Therefore, altered immune responses found in CatS\(+/–\) mice may also contribute to the retarded atherogenesis in CatS\(+/–\)/LDLR\(+/–\) mice.

In conclusion, this study provides direct in vivo evidence for involvement of the cysteine protease Cat S in atherogenesis in mice. The data presented support several possible mechanisms that contribute to this atherogenic role of Cat S. Our previous observations in diseased human arterial tissue suggest a similar function for this protease in human lesions. The present findings firmly place this cysteine protease alongside MMPs and serine proteases as proteolytic enzymes involved in atherosclerosis. This work raises numerous questions for future investigation and underscores the need to consider Cat S as a therapeutic target in arterial diseases.

Acknowledgments

The authors thank Harold A. Chapman and Karen Williams for reading the manuscript, and Eugenia Shvartz, Kenji Ohyauchi, Susumu Momozaki, Mika Kobayashi, and Akashi Izumi for their excellent technical assistance. This work is supported by grants from the NIH National Heart, Lung, and Blood Institute to G.-P. Shi (HL-60942) and P. Libby (HL-34636 and SCOR grant HL-56985).


