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Mast cells modulate the pathogenesis of elastase-induced abdominal aortic aneurysms in mice

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Abdominal aortic aneurysm (AAA), an inflammatory disease, involves leukocyte recruitment, immune responses, inflammatory cytokine production, vascular remodeling, neovascularization, and vascular cell apoptosis, all of which contribute to aortic dilatation. This study demonstrates that mast cells, key participants in human allergic immunity, participate in AAA pathogenesis in mice. Mast cells were found to accumulate in murine AAA lesions. Mast cell–deficient KitW−/−KitW−/− mice failed to develop AAA elicited by elastase perfusion or peri-aortic chemical injury. KitW−/−KitW−/− mice had reduced aortic expansion and internal elastic lamina degradation; decreased numbers of macrophages, CD3+ T lymphocytes, SMCs, apoptotic cells, and CD31+ microvessels; and decreased levels of aortic tissue IL-6 and IFN-γ. Activation of mast cells in WT mice via C48/80 injection resulted in enhanced AAA growth while mast cell stabilization with disodium cromoglycate diminished AAA formation. Mechanistic studies demonstrated that mast cells participated in angiogenesis, aortic SMC apoptosis, and matrix-degrading protease expression. Reconstitution of KitW−/−/KitW−/− mice with bone marrow–derived mast cells from WT or TNF-α−/− mice, but not from IL-6−/− or IFN-γ−/− mice, caused susceptibility to AAA formation to be regained. These results demonstrate that mast cells participate in AAA pathogenesis in mice by releasing pro-inflammatory cytokines IL-6 and IFN-γ, which may induce aortic SMC apoptosis, matrix-degrading protease expression, and vascular wall remodeling, important hallmarks of arterial aneurysms.

Introduction
Abdominal aortic aneurysm (AAA), a common aortic disease in the elderly, with an unknown pathogenesis, is characterized by extensive tissue remodeling, neovascularization, inflammatory cell infiltration, and vascular cell apoptosis, all of which contribute to aortic dilatation, the sine qua non of the disease (1). Leukocyte recruitment is a key feature of human AAA. After infiltration, the neutrophils, macrophages, and lymphocytes in human AAA lesions produce a spectrum of proinflammatory cytokines and matrix-degrading proteases that promote formation of AAA (1). Selective depletion of neutrophils (2), lymphocytes (3), matrix-degrading proteases (4, 5), or proinflammatory cytokines (6) impairs experimental AAA formation.

Mast cells are another inflammatory cell type recently identified in human aortic lesions, including those of atherosclerosis and AAA (7, 8). Classically, mast cells have been described as essential effector cells of immediate hypersensitivity and chronic allergic reactions that contribute to asthma, atopic dermatitis, and other allergic diseases (9). Despite the lack of direct evidence of mast cell participation in either atherosclerosis or AAA, enhanced accumulation of mast cells during human atherogenesis and AAA formation suggests that mast cells play a role in the pathogenesis of these vascular diseases. For example, active chymase, a unique mast cell protease important in the formation of angiotensin II and activation of MMP (10, 11), has been found at increased levels in extracts of human AAA lesions. Chymase-positive mast cells were detected in the media and adventitia in human AAA (8). Pharmacological inhibition of chymase activity reduces AAA progression in hamsters (12). In addition to a unique set of proteases, mast cells also release proinflammatory cytokines (IFN-γ, TNF-α, and IL-4, -5, -6, and -13) and chemokines (monocyte chemoattractant protein-1 [MCP-1], IL-8, RANTES), all of which may directly or indirectly participate in AAA formation. Stabilization of mast cells with disodium cromoglycate (DSCG) blocked the increase of plasma IL-6 in ApoE−/− mice (13). Many in vitro and in vivo studies have also demonstrated that IL-6 plays a role in lymphocyte activation and that IFN-γ stimulates lymphocytes, macrophages, ECs, and fibroblasts. In addition to activating T cells and B cells, IL-6 is required for fibroblast and EC MCP-1 expression, macrophage and leukocyte recruitment, MMP expression, angiogenesis, angiotensin II–mediated macrophage oxidized LDL uptake, and angiotensin II–mediated reactive oxygen species production and endothelial dysfunction, all of which are important to AAA pathogenesis (7). Mice lacking IFN-γ are resistant to AAA (3). Injection of IFN-γ into CD4+ T cell–depleted mice, which are resistant to aneurysm induction, can partially reconstitute aneurysms in CD4+ mice (3).

However, other inflammatory cell types in AAA lesions, such as macrophages, neutrophils, and lymphocytes, also produce potent extracellular matrix proteases as well as the aforementioned mast
cell mediators. For example, macrophages are a major source of the vascular wall destructive cysteine proteases (14), MMP (4, 5), and the proinflammatory cytokines TNF-α, IL-1β, and IFN-γ (3, 15), implicated in the development of human and murine AAA. Lymphocytes produce Th1 and Th2 cytokines in human arterial diseases (3, 15). Our recent studies demonstrated that CD4+ T cells, the major lymphocytes in human AAA lesions (3), produce MMP-9 and the cysteine proteases cathepsins S, K, and L (16), all of which are essential to vasculature remodeling (4, 14).

In this study, we used an aorta elastase perfusion–initiated mouse AAA model to determine whether mast cells modulate AAA formation. Reconstitution of mast cells from different cytokine-deficient mice into mast cell–deficient mice identified mast cell mediators essential to AAA. Finally, we determined whether we can regulate mouse AAA formation by activating or stabilizing mast cell activities in vivo.

Results

Mast cell accumulation and AAA formation. Biochemical observations revealed high numbers of chymase- and tryptase-positive mast cells in human AAA lesions and suggested an association of mast cell accumulation and AAA progression (8). To test this hypothesis, we used an established elastase perfusion model to produce AAA in mice (2, 4). In this model, the infrarenal aorta is locally exposed to pancreatic elastase, and aortic expansion forms over a period of 2 weeks as defined by an increase of aortic diameter by 100% greater than that before perfusion (4). Consistently, all mice developed AAA 14 days after elastase perfusion (Figure 1A). While normal aortas contained few if any mast cells (Figure 1, B and E), a significant increase of mast cell content appeared in the adventitia 7 days after perfusion (Figure 1, C and E) with further increase at 14 days after perfusion (Figure 1, D and E). Murine mast cells appear predominantly in the adventitia (Figure 1, C and D). These observations indicate that mast cells populate AAA induced by elastase perfusion and provide an opportunity to determine experimentally whether AAA formation requires mast cells.

Reduced AAA in mast cell–deficient mice. Accumulation of mast cells in AAA lesions suggests a participation of these cells in this disease. To test this hypothesis, we studied mast cell–deficient KitW-sh/KitW-sh mice in the elastase perfusion–induced AAA model. KitW-sh/KitW-sh mice contain a de novo inversion mutation consisting of a break point at the upstream regulatory region of the c-kit receptor gene that results in reduced Kit receptor expression and defective mast cell maturation and survival (17). However, this mutation does not affect the number or function of blood leukocytes, including lymphocytes (18). Elastase activities of bone marrow–derived neutrophils (6,185 ± 94 versus 6,195 ± 26, OD 515nm; P = 0.92) and blood monocytes (5,980 ± 167 versus 6,274 ± 62, OD 515nm; P = 0.18) showed no significant difference between WT mice and KitW-sh/KitW-sh mice. At baseline, histological analysis did not reveal any aortic wall elastin degradation, loss of medial SMCs, medial SMC apoptosis, or infiltration of macrophages and T cells in aortas from either WT or KitW-sh/KitW-sh mice before the elastase perfusion (data not shown). However, KitW-sh/KitW-sh mice were protected from AAA formation during 56 days following elastase perfusion. After perfusion expansion (day 0), the abdominal aortas of KitW-sh/KitW-sh mice showed no further expansion on days 7, 14, and 56, whereas all WT control mice had an average of more than 110% and approximately 170% aortic expansion 14 and 56 days after perfusion, respectively (Figure 2A). Importantly, while the lumen diameters of both WT and KitW-sh/KitW-sh mice increased at day 0 by nearly 30% due to elastase perfusion, the lumen diameters of KitW-sh/KitW-sh mice decreased continuously to those of preperfusion at day 56 after perfusion. In contrast, although there was a decrease on day 7, lumen diameters of WT mice increased consistently over time and reached 30% at day 56 after perfusion (Figure 2B).
Therefore, the AAA protection in KitW-sh/KitW-sh mice at day 14 was largely due to attenuation of adventitial thickening, whereas AAA protection in KitW-sh/KitW-sh mice at 56 days was the result of a combination of attenuation of both luminal expansion and adventitial thickening. Histological analysis of KitW-sh/KitW-sh mice showed better preservation of the internal elastic laminae (IELs) (Figure 2C), consistent with higher medial SMC content on day 14, as determined by semiquantitative grading (Figure 2D). Therefore, reduced levels of medial SMCs in WT mice may result from enhanced IEL fragmentation and an increase in medial SMC death. Although most mast cells reside in the adventitia, clusters of mast cells exist in the media where elastin degradation and SMC apoptosis occur. Immunostaining of serial sections with mouse MCP-4 (mMCP-4) (mast cells) (19) and \( \alpha \)-actin (SMCs) demonstrated numerous mast cells but fewer SMCs in the media (Figure 2E). Further, activated mast cells in the adventitia may also release granular contents capable of acting on cells at a distance, such as those in the media. Compared with KitW-sh/KitW-sh mice, WT mice showed overall more apoptotic cells on days 7 and 14 (\( \sim 4 \)-fold) (Figure 2F). By staining adjacent sections for both SMCs (\( \alpha \)-actin) (Figure 2G) and apoptosis (TUNEL) (Figure 2H), we explored the contribution...
of cell death to the loss of medial SMCs. Media poor in α-actin had abundant TUNEL staining. Significantly reduced levels of macrophages (Figure 2I) and CD3+ T cells (Figure 2J) in elastase-perfused KitW-sh/KitW-sh mice confirmed decreased inflammation. Neovascularization plays an essential role in AAA formation, and its presence associates with AAA lesion rupture (20). To determine whether mast cells modulate neovascularization, we stained tissue sections with an anti-CD31 antibody and found a reduced content of adventitial CD31+ microvessels in KitW-sh/KitW-sh mice both 7 and 14 days after elastase infusion (Figure 2K). Immunostaining of AAA serial sections from WT mice with mMCP-4 and CD31 antibodies demonstrated the presence of both microvessels and mast cells in the adventitia (Figure 2L), indicating a possible interaction between mast cells and ECs.

Observations from the KitW-sh/KitW-sh mice show that mast cells regulate AAA formation and suggest that they do so by modulating directly or indirectly the level of cellular inflammation, angiogenesis, and SMC apoptosis in aortic wall. However, the present data do not exclude the possibility that the reduced inflammation observed resulted from decreased AAA formation rather than the converse.

Mast cells promote periaortic CaCl2-induced AAA formation. To further test the hypothesis of mast cell participation in AAA pathogenesis, we induced AAA formation in WT and KitW-sh/KitW-sh mice using a second model of experimental AAA (5). Six weeks after periaortic CaCl2 injury, aortic diameter in WT mice increased more than 70%. In contrast, KitW-sh/KitW-sh mice showed a less than 25% increase in

Figure 4
BMMC reconstitution and mast cell activation and stabilization. (A) Aortic expansion in WT, KitW-sh/KitW-sh, and KitW-sh/KitW-sh mice reconstituted with BMMCs from WT, IL-6−/−, IFN-γ−/−, and TNF-α−/− mice. (B) Percentage of macrophage-positive areas in AAA lesions from different groups of mice. (C) IFN-γ levels in AAA lesions from different groups of mice. (D) IFN-γ immunostaining in aortic sections from KitW-sh/KitW-sh or WT mice treated with or without C48/80 or DSCG. (E) IL-6 levels in tissue extracts from different groups of mice. (F) Aortic expansion in WT mice treated with or without C48/80 or DSCG. (G) Elastic laminae degradation in WT mice treated with or without C48/80 or DSCG. (H) Mast cell contents in aortas are not affected by C48/80 but decrease in DSCG-treated mice. Different groups of mice are designated with different colored bars as indicated in the legend. All data are from the 14-day time point experiments. Data are presented as mean ± SE, and P < 0.05 was considered significant, Mann-Whitney test. Number of animals is indicated in the bars. *Compared with WT mice.
aortic diameter (Figure 3A) and preservation of the IEL (Figure 3B). Under these experimental conditions, no lumenal changes were noticed in WT mice 6 weeks after injury and absence of mast cells had no effect on lumen diameters. Our data presented in Figure 3, A and B, indicate that the prevention of AAA formation in Kit<sup>W-sh/</sup>Kit<sup>W-sh</sup> mice is independent of the animal models used.

Mast cell IL-6 and IFN-γ contribute to AAA formation. In Kit<sup>W-sh/</sup>Kit<sup>W-sh</sup> mice, protection from AAA formation may result either from altered c-kit signaling caused by the inversion mutation of the c-kit receptor gene or the absence of mast cells. To differentiate these possibilities, we introduced bone marrow–derived mast cells (BMMCs) from WT mice and mice deficient in mast cell cytokines IL-6, IFN-γ, and TNF-α into the tail veins of Kit<sup>W-sh/</sup>Kit<sup>W-sh</sup> mice using an established protocol (18, 19, 21, 22), followed by elastase perfusion to induce AAA formation. Reconstitution with WT BMMCs largely restored AAA formation and inflammatory cell (macrophages) infiltration in Kit<sup>W-sh/</sup>Kit<sup>W-sh</sup> mice 14 days after elastase perfusion (Figure 4, A and B), confirming that the absence of mast cells, rather than altered c-kit signaling, protects Kit<sup>W-sh/</sup>Kit<sup>W-sh</sup> mice from AAA formation. Despite TNF-α’s role in mediating some forms of mast cell–associated inflammation (23), Kit<sup>W-sh/</sup>Kit<sup>W-sh</sup> mice reconstituted with TNF-α–deficient BMMCs developed AAA and enhanced macrophage accumulation similar to those achieved with WT BMMCs. In contrast, BMMCs from IL-6– or IFN-γ–deficient mice did not have these effects (Figure 4, A and B). Failed restoration of AAA in Kit<sup>W-sh/</sup>Kit<sup>W-sh</sup> mice reconstituted with IL-6– and IFN-γ–deficient BMMCs did not result from altered maturation of IL-6– or IFN-γ–deficient BMMCs. Granular morphology, tryptase production, and cell-surface expression of FcεR of IL-6– and IFN-γ–deficient BMMCs did not differ from...
those of WT BMMCs (19). Further, immunostaining with antibodies against mouse chymase mMCP-4 demonstrated that the accumulation of IL-6– or IFN-γ–deficient mast cells that appeared in reconstituted KitW−/KitW− mice abdominal aortas 14 days after perfusion (19.3 ± 7.4 and 24.5 ± 3.7 mast cells/mm², respectively) did not differ significantly from that in abdominal aortas of WT or TNF-α–deficient BMMC-reconstituted KitW−/KitW− mice (29.7 ± 6.8 and 20.2 ± 4.9 mast cells/mm², respectively), indicating similar recruitment of the IL-6– and IFN-γ–deficient mast cells to the abdominal aortas following elastase perfusion. Therefore, mast cell–derived IL-6 and IFN-γ may play key roles in aortic remodeling and expansion. In agreement with this hypothesis, aortas from KitW−/KitW− mice and those reconstituted with IFN-γ– and IL-6–deficient BMMCs had significantly fewer IFN-γ–positive areas than those in aortic lesions from KitW−/KitW− mice reconstituted with WT or TNF-α–deficient BMMCs (Figure 4, C and D). The AAA tissue extract ELISA detected significant reduction of IL-6 in KitW−/KitW− mice relative to WT mice (Figure 4E). Therefore, from these experiments, we conclude that mast cell–derived IL-6 and IFN-γ play key roles in aortic remodeling and expansion.

Control of AAA by regulating mast cell activation. Activation of mast cells and the subsequent release of their granular constituents is a major mechanism whereby mast cells participate in pathobiological processes (24). Mast cells likely affect AAA pathogenesis by releasing IL-6, IFN-γ, and MMPs (25), K (27), and L (16) reduce atherogenesis in mice. Mice deficient in cathepsins C (28) or S (unpublished data) and MMP-9 or MMP-2 (4, 5) also develop significantly less AAA in experimental models. Mast cells release a unique set of proteases, i.e., tryptase and chymase, that may participate in AAA directly or indirectly by activating MMP (11). Our recent in vitro study demonstrates that mast cells induce elastinolytic cathepsin expression in both aortic SMCs and ECs (19). Labeling the cysteine protease active site with [125I]JPM (29) (Figure 5, A and B) and determining MMP activity by gelatin zymography (Figure 5, A and C) (30) revealed much lower cathepsin and MMP activities in aortic extracts from KitW−/KitW− mice than in WT mice 14 days after elastase perfusion. Importantly, the activities of cathepsin and MMP in...
the aortas of KitW–/KitW– mice increased following reconstitution with BMMCs from WT or TNF-α–deficient mice but not those from IL-6– or IFN-γ–deficient mice (Figure 5, B and C), suggesting that mast cell–derived IL-6 and IFN-γ stimulate matrix-degrading protease IEL degradation (Figure 4, F and G). These observations agree with prior in vitro studies (19). BMMCs from IL-6– or IFN-γ–deficient mice produce less than 5% IL-6 or IFN-γ of those from WT mice. In contrast, BMMCs from TNF-α–deficient mice release 50% IL-6 and 30% IFN-γ of those from WT mice and thus may suffice to induce vascular cell protease expression in vitro (19) and in vivo (Figure 5, B and C). Activities of both cathepsin (e.g., cathepsin B) and MMP (e.g., MMP-2) increased in aortic extracts from C48/80-treated mice but decreased to the levels found in KitW–/KitW– mice following the stabilization of mast cells with DSCG (Figure 5, A–C) when both assays contained an equal amount of proteins (Figure 5A). In situ elastin zymography optimized for cysteine proteases (pH 5.5 with both EDTA and diithiothreitol) supported these observations. While negligible elastinolytic activities (green fluorescence in the adventitia) were detected in normal aorta, saline-treated aorta, and heat-inactivated elastase–treated aorta, AAA lesions in WT mice showed much higher elastinolytic activity than did those from KitW–/KitW– mice, as indicated by the differences in percentage area of adventitia fluorescence in Figure 5D. Media green fluorescence may come from both increased elastase activity in AAA lesions and intact elastin filament autofluorescence, which also appeared in normal adventitia in Figure 5D. Nonselective cathepsin inhibitor E64d (20 μM) greatly inhibited such activity in parallel sections. Increased AAA expansion (Figure 4F) and IEL fragmentation (Figure 4G) in mice treated with the mast cell activator C48/80 accompanied enhanced in situ activity of cathepsin elastase. Likewise, impaired AAA and reduced IEL degradation in DSCG-treated mice (Figure 4, F and G) may have resided, at least in part, from a reduction in elastinolytic cathepsin activity (Figure 5D). Omission of fluorescent elastin substrate was utilized for the control (Figure 5D).

Role of mast cells in neovascularization and aortic SMC apoptosis. Angiogenesis and apoptosis, like elastinolysis, are common in human AAA (20). Inhibition of angiogenesis or apoptosis reduces AAA formation in rodents (31, 32). In KitW–/KitW– mice, decreases in the number of microvessels (Figure 2K) and total apoptotic cells (Figure 2F) suggest that mast cells participate in these pathologic events. After incubating mouse aortic rings in Matrigel with degranulated mast cell supernatant from WT mice (Figure 6, B and C) but not after incubation with culture medium alone (Figure 6, A and C), microvessel sprouting, similar to that seen in the aortic rings treated with IL-6–, IFN-γ–, and TNF-α–deficient mast cell supernatant (data not shown), occurred at substantial levels, suggesting that additional factors from mast cells contributed to microvessel growth in AAA. Immunostaining a cross section of the Matrigel with mouse CD31 antibody revealed well-formed microvessels derived from the aorta after stimulation with medium conditioned by mast cell degranulation (Figure 6D).

α-Actin staining of SMCs of AAA lesions in WT mice showed areas negative for α-actin (Figure 2G) but rich in apoptotic cells (Figure 2H) in contrast to reduced apoptotic cells in lesions of KitW–/KitW– mice (Figure 2F), consistent with a role of mast cells in aortic SMC apoptosis (33). To identify mast cell–derived mediators important for SMC apoptosis, we used pyrroline dithiocarbamate (PDTC) with or without supernatant from degranulated BMMCs to induce apoptosis of mouse aortic SMCs. TUNEL-positive cells appeared within hours. However, supernatant from degranulated BMMCs from WT and TNF-α–deficient mice but not that from IL-6– and IFN-γ–deficient mice enhanced SMC apoptosis (Figure 6, E and F), supporting a role of mast cell–derived IL-6 and IFN-γ in aortic SMC apoptosis.

Discussion

Although mast cells have been identified in human AAA lesions, evidence for a direct participation of mast cells in this vascular disease is lacking. The availability of mast cell–deficient mice and established mouse AAA models made it possible to determine whether these cells are essential mediators of AAA and not merely a hallmark.

This study used a BMCC reconstitution technique to determine whether mast cell–derived IL-6, IFN-γ, and TNF-α plays a role in AAA formation (Figure 4A). TNF-α is expressed in macrophages and lymphocytes both in the atherosclerotic (34) and AAA (35) lesions. In ApoE−/− mice with diet-induced atherosclerosis, a lack of TNF-α reduces atherosclerosis (36). In this model, TNF-α plays a proatherogenic role by upregulating expression of the vascular adhesion molecules ICAM-1 and VCAM-1 and the chemokine MCP-1 in the vascular wall and by inducing expression of scavenger receptor class A and uptake of oxidized LDLs in macrophages. Although it remains to be determined whether TNF-α plays a similar role to that in atherosclerosis, both serum (37) and aneurysmal tissues (35) from AAA patients have increased TNF-α levels. However, our data with TNF-α−/− mast cells did not support a direct role of mast cell–derived TNF-α in AAA. This does not exclude the possibility that TNF-α plays a role in AAA formation; rather, it is possible that TNF-α produced from other cells may compensate for its absence from mast cells.

AAA formation requires mast cell IFN-γ and IL-6 but not TNF-α. Reconstitution of mast cell–deficient KitW–/KitW– mice with IFN-γ−/− or IL-6−/− BMCCs did not restore the AAA phenotype (Figure 4A). These observations agree with the findings in IFN-γ−/− mice (3), which are protected from CaCl2–induced AAA. Both this study (3) and others using a histocompatibility-mismatched aortic transplantation model in IFN-γ receptor–deficient recipient mice (6) demonstrated that Th2-slanted immune responses promote AAA formation. Indeed, mast cells from IL-6−/− and IFN-γ−/− mice produce negligible IFN-γ and IL-6, suggesting cooperative regulation of these 2 cytokines (19). HeLa cells and monocytes exhibit IFN-γ-regulated expression of IL-6 (38, 39). Enhanced IL-6 and IFN-γ, which may come in part from mast cells, were detected in cultured explants of human AAA lesions and in serum of AAA patients (15, 40). AAA sections or tissue extracts from WT mice contained much higher levels of both IFN-γ and IL-6 than those from KitW–/KitW– mice (Figure 4, C–E). In addition to their possible roles in activating macrophages and inducing MMP and cysteine protease expression (41), which are also critical to AAA formation (4, 5), IL-6 and IFN-γ appeared to regulate the formation of AAA by modulating lesional SMC apoptosis (Figure 6, E and F). Our observations from in vitro cell-based assay and the murine AAA model not only demonstrate that mast cell–derived IL-6 and IFN-γ are essential to AAA but also suggest that other sources of IL-6 and IFN-γ, e.g., inflammatory cells such as T cells (3) and macrophages (42), do not compensate for their absence in mast cells. Such cell type–dependent functional specificity is not unique to IL-6 and IFN-γ, as similar activity has been reported for other cytokines. For example, mast cell TNF-α is required for recruiting neutrophils to...
infected tissue compartments (43), macrophage and neutrophil TNF-α are required for resistance to intracellular *Listeria* infection, and T cell TNF-α is required for protection from high bacterial loads (44). Thus, while it may seem intuitive that, because cytokines are made by many cell types, the absence of their production by one cell type would be compensated for by their production by another cell type, when tested in experimental models, this compensation does not always prove to be true, and depending on the end point measured, a single cellular source of a cytokine can have profound influence over modulating that end point.

Our in vitro experiments demonstrate that mast cell–derived IL-6 and IFN-γ are required for apoptosis but not neovascularization. Insignificant differences in the microvessel growth from the aortic ring Matrigel assay between WT and IL-6–, IFN-γ–, and TNF-α–deficient BMMCs suggest that mast cell mediators other than IL-6, IFN-γ, and TNF-α modulate neovascularization in this assay even though IL-γ, IL-6, and TNF-α have been implicated in angiogenesis. In fact, mast cell–derived bFGF, angiopoietin-1, chymase/trypsin, and a number of other proinflammatory cytokines and chemokines (45) have been implicated in angiogenesis. These mast cell mediators may have obscured the effect of mast cell IL-6, IFN-γ, and TNF-α in our in vitro assays (Figure 6, A–C).

Pharmacologic regulation of mast cell activation and degranulation directly controlled AAA formation in mice (Figure 4, F and G). These data not only strengthen the hypothesis that mast cells are essential mediators of AAA pathogenesis but also suggest that treatments blocking mast cell degranulation may prevent or delay AAA progression in humans. Compound 48/80 has been widely used to degranulate mast cells in live animals (46). Although the detailed mechanisms of how C48/80 activates mast cells need to be investigated further, recent observations suggest that C48/80 acts on mast cells via activation of phospholipase D (47) and Gα13 G protein subunits (48) to trigger the release of histamine as well as other granule components including cathepsins and MMP-activating chymases or cytokines, which may influence neighboring vascular cells. This hypothesis is supported by the findings of increased cysteine protease and MMP activities, elevation of AAA tissue IFN-γ and IL-6, and enhanced medial elastin degradation in aortic tissues isolated from C48/80-treated mice (Figure 4). In contrast, DSCG is a drug widely used in the treatment of asthmatic patients. Observations from in vitro tests and animal models show that the effect of DSCG is related to mast cell stabilization. Importantly, this compound reduces mouse AAA by 40% (Figure 4). Impaired AAA in DSCG-treated mice is likely due to reduced release of mast cell granule contents. This hypothesis is consistent with reduced AAA lesion cysteine protease and MMP activities and IFN-γ and IL-6 levels found in DSCG-treated mice (Figures 4 and 5).

While this study provides evidence that mast cells regulate AAA formation, it also leaves several important questions unanswered. First, depletion of neutrophils in mice impairs elastase perfusion–induced AAA (2), and mice lacking T lymphocytes have protection from CaCl2–induced AAA (3). Determination of whether lack of mast cells in the current study affected the function of either neutrophils or T lymphocytes will require further investigation. Although we did not find significant differences of elastase activity of neutrophils from WT and Kitw/+ Kitw/+ mice, unrecognized interactions between these cell types may exist. Second, it remains unknown what mediated the increased number of mast cells in the injured aortas following elastase perfusion. Five weeks after reconstitution, we detected few if any mast cells in the aortas.

In contrast, elastase perfusion enhanced mast cell numbers in the aortas. One possible explanation is that elastase perfusion induces aortic wall SMC expression of chemokines MCP-1 and RANTES (49), which may attract mast cells (50). Further, elastase perfusion may generate elastin degradation products that signal fibroblast, monocyte, and macrophage chemotaxis (51). Delineation of the exact mechanism will require further investigation. Third, mast cell–mediated neovascularization and SMC apoptosis remain poorly understood. Although our data suggest that SMC apoptosis involves mast cell–derived IFN-γ and IL-6, whether these cytokines regulate SMC apoptosis directly or indirectly remains unknown. Our prior study demonstrated that mast cell IFN-γ and IL-6 augment SMC and EC expression of cysteine proteases (19), which participate in cell apoptosis (52). IFN-γ and IL-6 from mast cells may have induced cathepsin expression and further led to SMC apoptosis, a hypothesis that has not been tested in the current study. Fourth, it remains uncertain whether increased expression and activities of cathepsins and MMP in AAA lesions (Figure 5) come from mast cells or other vascular cells. Mast cells contain large quantities of cathepsins (53) and mast cell chymases that participate importantly in MMP activation (11). However, mast cell cytokines (e.g., IL-6 and IFN-γ) are known stimulators of cathepsin and MMP expression, and our previous study (19) and the current study demonstrate concurrently their importance in arterial wall remodeling. Therefore, it is possible that mast cells contribute to AAA by releasing both proteases and cytokines, a hypothesis that requires further investigation.

In summary, data from this study establish that mast cells play a role in the development of AAA by affecting matrix-degrading protease expression, medial SMC apoptosis, and microvessel growth. More importantly, our data suggest a novel and effective therapeutic strategy to control human AAA progression by preventing release of mast cell products using mast cell–stabilizing compounds.

**Methods**

**AAA production.** Eight- to ten-week-old male WT mice (from The Jackson Laboratory, C57BL/6 background) and male Kitw/+Kitw/+ mice (C57BL/6 background) (17, 21) were used for aortic elastase perfusion, as detailed elsewhere (4). Aortic diameters were measured before the surgery (day −1), immediately following elastase perfusion (day 0), and 7, 14, and 56 days after perfusion. Abdominal aortas were harvested at each time point. Lumen diameters were determined by measuring the lumen perimeters of cross sections of frozen aortas.

To produce AAA by periaortic chemical injury, 8-week-old male mice were used for aorta CaCl2 injury as described (5). Aortic diameters were measured before CaCl2 application and 6 weeks after the injury.

To harvest AAA tissues, after measuring the aortic diameter and collecting the blood, we removed the aortic zone of maximal expansion, either directly freezing the tissue for protein extraction or OCT embedding for cross section preparation. Immunohistological analysis used 6-μm sections while in situ elastin zymography used 8-μm cross sections. All mouse experiments were performed under a protocol preapproved by the Animal Research Committee of Harvard Medical School.

**BMMC preparation and reconstitution.** WT (C57BL/6), IFN-γ−/− (congenic C57BL/6, N10), IL-6−/− (congenic C57BL/6, N11), and TNF-α−/− (C57BL/6*129/Sv, N1) mice were obtained from The Jackson Laboratory. TNF-α−/− mice were backcrossed to a congenic C57BL/6 background (N10). Bone marrow was isolated from these mice for BMMC differentiation for 5 weeks in the presence of recombinant murine IL-3 (PeproTech) and murine stem cell factor (PeproTech) as reported (21). At this time, the cell popula-
tions consisted of more than 95% mast cells as assessed by the presence of metachromatic granules in toluidine blue–stained cells and FACS analysis for the disappearance of progenitor cell marker stem cell antigen-1. To reconstitute BMMCs in vivo, the tail veins of 5-week-old male Kit<sup>W-sh</sup>/Kit<sup>W-sh</sup> mice were injected with BMMCs (1 × 10<sup>5</sup> cells/mouse). Mice were perfused with elastase to produce AAA 5 weeks after reconstitution.

**Mast cell activation and stabilization in vivo.** C48/80 activated the mast cells in WT mice, and DSCG stabilized the mast cells in WT mice. After mice (C57BL/6, male, 10 weeks old) recovered from elastase perfusion surgery (day 0), they received the first i.p. injection of C48/80 (Sigma-Aldrich, 4 mg/kg/d) or DSCG (Sigma-Aldrich, 25 mg/kg/d). Each animal received daily injection for the duration of the experiment thereafter (14 days).

**Mast cell content measurement.** Immunostaining with rabbit anti-mouse mMCP-4 antibody (1:50) determined the mast cell number on aortic sections (19). Antibody specificity was confirmed using WT mouse AAA sections immunostained with rabbit preimmune serum and Kit<sup>W-sh</sup>/Kit<sup>W-sh</sup> mouse aortic sections immunostained with the mMCP-4 antibody at the same antibody dilution. The total number of mMCP-4–positive cells on each section were counted manually and blindly, and the total area was determined using computer-assisted image analysis (Image-Pro Plus; MediaCybernetics). Immunohistochemistry, ELISA, and lesion characterization. Frozen sections (6 μm) were stained for macrophages (Mac-3), T cells (CD3), SMCs (α-actin), ECs (CD31), mast cells (mMCP-4), elastin (Verhoeff–van Gieson), and IFN-γ as described previously (19, 26). Lesion apoptotic cells were determined with the In situ Apoptosis Detection kit according to the manufacturer’s instructions (Chemicon International). Elastin degradation and lesion SMC content were graded according to the grading keys shown in Figure 2, C and D. To determine lesion IL-6 levels, frozen aortas were pulverized for protein extraction in a pH 7.5 buffer containing 50 mM NaAc and 1% Triton X-100. Protein concentration was determined with a BCA kit (Bio-Rad) and IL-6 levels measured by ELISA (R&D Systems). All immunostained cells were counted or calculated blindly by at least 2 observers.

**Cysteine protease active site labeling and gelatin gel in situ zymography.** Pulverized aortic tissues were lysed in a pH 5.5 buffer. The cysteine protease activity in tissue lysates was determined with [125I]JPM labeling (2 μg protein/lane) (29), and the MMP activity was assessed by in-gel gelatinase zymogram assay (5 μg protein/lane) as described (30). Equal protein loading (7 μg protein/lane) was confirmed by SDS-PAGE followed with Coomassie staining.

**Mast cell degranulation, aortic ring assay, and SMC apoptosis assay.** In vitro–differentiated BMMCs were degranulated with mouse anti-DNP IgE monoclonal antibody as described previously (54). Degranulation efficiency was calculated as follows: (experimental supernatant – control supernatant)/ (experimental supernatant + experimental cell lysates supernatant). An aortic ring assay was used to test mast cell function in angiogenesis. In brief, a 96–well plate was coated with 50 μl of Matrigel (BD Biosciences). A 1-mm–long mouse aortic ring was laid on the top of the Matrigel and covered with 100 μl of Matrigel. After solidification, 100 μl of RPMI (10% FBS) with or without 100 μl of degranulated BMNC supernatant was added to each well. After 7–10 days of culture, the aortas were photographed and the endothelial outgrowth was analyzed using Image-Pro Plus software. To confirm a microvesSEL phenotype of sprouts from the aortic ring, a Matrigel containing aortic ring and sprouts was embedded in OCT followed by sectioning and immunostaining with CD31.

Mouse aortic SMC apoptosis was induced with 50 μM of PDTC (Sigma-Aldrich) as previously described with or without degranulated mast cell supernatant prepared from the same amount of BMMCs (55). Apoptotic SMCs were detected using the in situ cell death detection kit according to the manufacturer’s instructions (Roche Applied Science). Data were presented as mean ± SE of percentage of apoptotic cells.

**In situ elastin zymography.** AHA lesion elastolysis activity was determined in 8-μm frozen sections using elastin conjugated with quenched fluorescein (DQ elastin; Invitrogen) as a substrate, which requires cleavage by elastinolytic enzymes to become fluorescent. In brief, DQ elastin (1 mg/ml in H<sub>2</sub>O) was mixed 1:10 with 1% low-melting agarose (Sigma-Aldrich). This mixture (20 μl) was added on top of each section, covered, and gelled at 4°C. Following incubation at 37°C (48 hours), fluorescence was examined under fluorescent microscopy. Cysteine protease activity was determined using an EDTA-containing pH 5.5 buffer (56) with or without E64d (20 μM). Zymographic images were acquired using identical shutter conditions. The percentage of fluorescence intensity of each cross section, excluding the media area due to medial elastin filament autofluorescence, was measured using computer-assisted image quantification (Image-Pro Plus software).

**Neutrophil and monocyte elastase activity assay.** Neutrophils from WT and Kit<sup>W-sh</sup>/Kit<sup>W-sh</sup> mice were prepared from bone marrow by gradient centrifugation. Bone marrow cells were overlaid on the top of 62.5% Percoll (Sigma-Aldrich) followed by centrifugation at 600 g for 30 minutes. Purified neutrophils were lysed in a pH 8.5 buffer containing 150 mM NaCl, 200 mM Tris HCl, and 1% Triton X-100. Monocytes from WT and Kit<sup>W-sh</sup>/Kit<sup>W-sh</sup> mice were isolated from blood by overlaying diluted blood (1:1 in PBS) on LSM Lymphocyte Monocyte Separation Medium (LLC; MP). Isolated monocytes were lysed in a pH 5.5 buffer containing 50 mM NaAc, 1% Triton X-100, and 12 μM dithiothreitol. Equal amounts of protein (60 μg) were added to a final volume of 100 μl with 50 μg of elastin conjugated with quenched fluorescein (DQ elastin; Invitrogen) followed by incubation at 37°C for 24 hours. Elastase activity was determined by measuring the OD<sub>492</sub>.

**Statistics.** The statistical significance of data sets from this study was examined with a nonparametric Mann-Whitney test. Data from the gel densitometry analysis were also analyzed using the nonparametric Mann-Whitney test due to the relatively small sample sizes and the non-Gaussian distribution of the data. P values of less than 0.05 were considered significant.

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