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Proatherogenic immune responses are regulated by the PD-1/PD-L pathway in mice

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T lymphocyte responses promote proatherogenic inflammatory events, which are influenced by costimulatory molecules of the B7 family. Effects of negative regulatory members of the B7 family on atherosclerosis have not been described. Programmed death–ligand 1 (PD-L1) and PD-L2 are B7 family members expressed on several cell types, which inhibit T cell activation via binding to programmed death–1 (PD-1) on T cells. In order to test whether the PD-1/PD-L pathway regulates proatherogenic T cell responses, we compared atherosclerotic lesion burden and phenotype in hypercholesterolemic PD-L1/2−/− LDLR+/− mice and LDLR+ mice. PD-L1/2 deficit led to significantly increased atherosclerotic burden throughout the aorta and increased numbers of lesional CD4+ and CD8+ T cells. Compared with controls, PD-L1/2−/− LDLR+/− mice had iliac lymphadenopathy and increased numbers of activated CD4+ T cells. Serum levels of TNF-α were higher in PD-L1/2−/− LDLR+/− mice than in controls. PD-L1/2−/− deficient APCs were more effective than control APCs in activating CD4+ T cells in vitro, with or without cholesterol loading. Freshly isolated APCs from hypercholesterolemic PD-L1/2−/− LDLR+/− mice stimulated greater T cell responses than did APCs from hypercholesterolemic controls. Our findings indicate that the PD-1/PD-L pathway has an important role in downregulating proatherogenic T cell response and atherosclerosis by limiting APC-dependent T cell activation.

Introduction
Atherosclerosis is a chronic inflammatory disease in which T cell-mediated immunity plays a significant component (1). Th cells are evident in the atherosclerotic plaque and have been shown to be important in the development of atherosclerosis (2, 3). T cell-mediated immune responses toward specific plaque antigens such as oxidized LDL or HSP-60/65 may have an important role in the propagation of the inflammatory process (4, 5). The development of atherosclerosis in animal models is enhanced by proinflammatory Th1 cells, through the secretion of IFN-γ (6, 7).

T cell responses to antigen are modulated by members of the B7 family of molecules expressed on APCs, which bind to signaling receptors of the CD28 family expressed on the T cells. Some of these molecules provide positive costimulatory signals that promote T cell activation, and other members of these families transduce inhibitory signals that limit the T cell response. The major positive costimulatory pathway involves B7-1 and B7-2 (also known as CD80 and CD86, respectively), which bind to CD28 on T cells (8). ICOS is another CD28 family member, which binds ICOS ligand and transduces signals that promote T cell activation.

Other members of the B7 family and their corresponding receptors on T cells have an important role in modulating or downregulating T cell activation (9). CTLA-4, which is a CD28 family member expressed on T cells, binds B7-1 and B7-2 and induces negative signals that block T cell activation. Programmed death–1 (PD-1) is a CD28 family member on T cells that binds programmed death–ligand 1 (PD-L1) and PD-L2 (also known as B7-H1 and B7-DC, respectively) on APCs and transduces signals that inhibit T cell immune-mediated responses (10, 11). PD-L1 and PD-L2 are highly expressed after antigen stimulation on APCs, specifically DCs, macrophages, and B cells (12). PD-1 on T cells is the only known receptor of these ligands (12, 13). Although there are reports suggesting that PD-L1 may have a stimulatory function (14), a receptor with such a function has yet to be found. Different B7/CD28 family members’ costimulatory and regulatory pathways may have distinct effects on atherosclerosis, depending on the pattern of expression and the state of the immune system. We have previously shown that the costimulatory molecules B7-1 and B7-2 support proatherogenic T cell responses and the absence of B7-1 and B7-2 significantly reduce the lesion development in hematoepoietically unmanipulated mice (15). Conversely, in the setting of radiated bone marrow chimeric LDLR+/− mice, ablation of the B7-1/2–CD28 pathway impairs Treg development and enhances atherosclerosis (16). Similarly, we have found that deficiency of ICOS results in decreased Treg function, enhanced immune responsiveness, and more atherosclerosis in LDLR+/− bone marrow chimeric mice (17).

To our knowledge, the effects of negative regulatory pathways of the B7/CD28 family on atherosclerosis have not previously been explored. We hypothesized that PD-L1 and PD-L2 function to downregulate T cell responses to plaque antigens and thereby limit atherosclerosis. To test this hypothesis, we generated PD-L1/ PD-L2/LDLR triple-knockout mice by cross-breeding PD-L1/PD-L2 double-knockout mice with LDL receptor (LDLR) knockout mice, all on a C57BL/6 background. We then examined the influence of PD-L1/2 deficiency on the extent and phenotype of diet-induced atherosclerosis and plaque antigen–specific cell mediated responses in athero-prone LDLR-deficient mice. Our data demonstrate that the PD-1/PD-L pathway exerted a significant regulatory influence on the immune responses and atherosclerosis of hypercholesterolemic mice.

Nonstandard abbreviations used: acLDL, acetylated LDL; LDLR, LDL receptor; ORO, Oil red O; PD, programmed death; PD-L, programmed death–ligand.

Conflict of interest: The authors have declared that no conflict of interest exists.

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**Table 1**

<table>
<thead>
<tr>
<th>Mean weights and serum cholesterol</th>
<th>Weight, 6 wk (g)</th>
<th>Weight, 16 wk (g)</th>
<th>Total serum cholesterol, 16 wk (mg/dl)</th>
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</thead>
<tbody>
<tr>
<td>PD-L1/2−/-LDLR−/−</td>
<td>20.4 ± 0.6</td>
<td>29.8 ± 1.4</td>
<td>1,362 ± 78</td>
</tr>
<tr>
<td>LDLR−/−</td>
<td>20.7 ± 0.9</td>
<td>28.3 ± 1.6</td>
<td>1,396 ± 102</td>
</tr>
</tbody>
</table>

There was no significant difference between the groups when comparing weight or serum cholesterol (n = 14 per group).

**Results**

Weights and serum cholesterol levels. There was no significant difference in mean weight between the control LDLR−/− and PD-L1/2−/−LDLR−/− groups (Table 1). There were also no statistical differences in total serum cholesterol levels (Table 1) or cholesterol in specific lipoproteins (data not shown) between the 2 groups.

Quantitative analysis of atherosclerotic lesions. The absence of PD-L1 and PD-L2 led to markedly increased atherosclerotic burden in control LDLR−/− mice on a high-cholesterol diet. This was evident at 5 and 10 weeks of high-cholesterol diet feeding (Figure 1). At 5 weeks of diet, the PD-L1/2−/−LDLR−/− mice had a more than 2-fold increase in aortic sinus atherosclerotic burden compared with control mice. Mean lesion areas were 96,080 ± 22,320 μm² in control mice (n = 8) and 213,400 ± 23,260 μm² in PD-L1/2−/−LDLR−/− mice (n = 7; P < 0.01). Fractional lesion areas — the cross-sectional lesion area expressed as a fraction of the total aortic root sectional area — were 20.2% ± 3.9% in control mice (n = 8) and 49.2% ± 3.4% in PD-L1/2−/−LDLR−/− mice (n = 7; P < 0.001; Figure 1E). After 5 weeks of high-cholesterol diet, there were minimal atherosclerotic lesions in the aortic arch or descending aorta, and significant differences between the groups could not be ascertained at this age (data not shown). After 10 weeks of diet, there was a 2-fold increase in aortic root disease and 3-fold increases in aortic arch and descending aorta disease (Figure 1, F–I). The mean lesion areas in the aortic sinus were 299,300 ± 25,430 μm² in control mice (n = 13) and 569,300 ± 35,410 μm² in PD-L1/2−/−LDLR−/− mice (n = 23; P < 0.001). Fractional lesion areas were 47% ± 2.5% in control mice (n = 13) and 73.2% ± 2.3% in PD-L1/2−/−LDLR−/− mice (n = 23; P < 0.0001; Figure 1F). The mean lesion sizes in the aortic arch were 40,710 ± 9,115 μm² in control mice (n = 12) and 122,400 ± 16,820 μm² in PD-L1/2−/−LDLR−/− mice (n = 23; P < 0.01). Fractional lesion area increased from 20.7% ± 2.8% in control mice (n = 12) to 50.3% ± 3.4% in PD-L1/2−/−LDLR−/− mice (n = 23; P < 0.001; Figure 1G). The en face lesional areas in the descending aorta were 1.7% ± 0.2% in control mice (n = 6) and 5.5% ± 0.8% in PD-L1/2−/−LDLR−/− mice (n = 8; P < 0.01; Figure 1H). Oil red O (ORO) staining revealed that PD-L1/2 deficiency resulted in increased lipid content after 10 weeks of diet in the aortic sinus and arch. Aortic sinus lipid-positive areas were 25,080 ± 3,964 μm² in control mice (n = 13) and 56,610 ± 5,598 μm² in PD-L1/2−/−LDLR−/− mice (n = 23; P < 0.001). Aortic arch lipid-positive areas were 4,063 ± 1,057 μm² in control mice (n = 12) and 9,637 ± 1,605 μm² in PD-L1/2−/−LDLR−/− mice (n = 23; P < 0.05).

Phenotypic analysis of atherosclerotic lesions. Analysis of the cellular content in atherosclerotic lesions in both groups of mice was determined by specific immunostaining in aortic sinus sections. The results of quantification of the positively stained areas are shown in Figure 2.

After 10 weeks of high-cholesterol diet feeding, there was a 3-fold increase in CD4+ T cells as well as a marked increase in CD8+ T cell infiltration in the intima of PD-L1/2−/−LDLR−/− mice compared with control mice (Figure 2). In addition, there was a 2-fold

![Image](2975)

**Figure 1**
Effect of PD-L1/PD-L2 deficiency on atherosclerotic burden. (A and B) Representative cross sections of the ORO-stained aortic sinuses are shown for female LDLR−/− (Control) (A) and PD-L1/2−/−LDLR−/− (PD-L1/2−/−) mice (B) after 10 weeks of high-cholesterol diet (original magnification, ×10). (C and D) Representative longitudinal sections of aortic arch are shown for female LDLR−/− (C) and PD-L1/2−/− mice (D) after 10 weeks of high-cholesterol diet (original magnification, ×100). (E–H) Quantitative analysis of lesions was performed on aortic roots taken after 5 weeks of high-cholesterol diet (E), and on aortic root (F), aortic arch (G), and descending aorta (H) after 10 weeks of high-cholesterol diet. Each data point represents the mean value obtained for each mouse; horizontal bars represent the mean value for each group. *P < 0.01, †P < 0.001, ‡P < 0.0001 versus control.
increase in macrophage content as well as a significant increase in smooth muscle cell content and collagen deposition in the lesions of the PD-L1/2−/−LDLR−/− mice (Figure 2). These differences were also significant when atherosclerosis was evaluated according to gender. These data clearly demonstrate that the absence of PD-L1/2 increases inflammatory cell infiltration in atherosclerotic lesions, and this is associated with more smooth muscle cell infiltration or proliferation and more collagen deposition.

Figure 2
Effect of PD-L1/2 deficiency on atherosclerotic phenotype. Representative sections of aortic sinus (adjacent to the ORO-stained sections in Figure 1) were immunohistochemically stained with antibodies specific for CD4 (A), CD8 (B), Mac3 for macrophages (C), and smooth muscle actin for smooth muscle cells (D). Collagen types I and III were stained by Picrosirius red, and the sections were analyzed by polarization microscopy (E). For each molecule, representative immunohistochemically stained sections (original magnification, ×200) and quantitative analysis of the staining for each experimental group are shown. Each data point represents the mean value determined for each mouse; horizontal bars represent the mean value for each group. Two-color immunofluorescence–stained sections from PD-L1/2−/−LDLR−/− mice are also shown using antibodies specific for CD3 (red) plus CD4 or CD8 (green). Double-stained cells appear yellow in the merged image. Sections are shown for each type of stain. P < 0.0001 versus control.
Expression of PD-L1 in hypercholesterolemic mice. PD-L1 is expressed on hematopoietic APCs (12, 18) and endothelium in inflammatory tissues, and this local expression may be important in the inhibitory function of this molecule (19, 20). Furthermore, PD-L1 is constitutively expressed on APCs, but is upregulated in states of immune activation (12), including during immune-mediated diseases (21). Because PD-L1/2 deficiency clearly increased atherosclerosis in LDLR-null mice, we asked whether PD-L1 expression is influenced by the hypercholesterolemic environment and whether it is expressed at the local level of the atherosclerotic lesion. We found PD-L1 expression by immunohistochemistry in aortic lesions (Figure 3A). Using 2-color immunofluorescence microscopy, we found PD-L1 on both DCs and macrophages (Figure 3B). Although PD-L1 is not abundantly expressed, and only some lesional macrophages are stained, PD-L1 was consistently found in most lesions, including those in the aortic sinuses and aortic arches.

We also examined the expression of PD-L1 and PD-L2 on DCs from the iliac lymph nodes, which may drain the descending aorta. Other lymph nodes were not examined. The fraction of PD-L1–expressing DCs in these nodes was increased in mice fed a high-cholesterol diet for 10 weeks compared with control diet–fed mice (Figure 4A). The expression of PD-L2 was not significantly different between the 2 groups. High-cholesterol diet feeding also led to a significant increase in the percentage of iliac lymph node DCs expressing the costimulatory molecule B7-1, and this increase was seen after 3 and 10 weeks of high-cholesterol diet. These data would suggest that these molecules are upregulated as the inflammatory events associated with atherosclerosis develop. We also examined the effect of high-cholesterol diet on the expression of PD-L1, PD-L2, and B7-1 in peritoneal macrophages. An increase in the number of PD-L1–expressing peritoneal macrophages was seen in LDLR–/– mice fed high-cholesterol diet for 10 weeks compared

Figure 3
Expression of PD-L1 in atherosclerotic lesions. (A) PD-L1 was detected by immunohistochemistry in aortic sinus lesions of control LDLR–/– mice after 10 weeks of high-cholesterol diet. The same tissues from PD-L1/2–/–LDLR–/– mice did not stain, and served as negative controls. Original magnification, ×400. (B) Two-color immunofluorescence stains for PD-L1 (green) and either CD11c or Mac3 (red) were performed on cholesterol aortic sinus lesions of LDLR–/– mice fed cholesterol-containing diet for 10 weeks. Top: PD-L1 staining alone. Bottom: Combined PD-L1 and MAC3 or PD-L1 and CD11c stains, in which superimposed stains appear yellow.

Figure 4
Hypercholesterolemia results in increased PD-L1, PD-L2, and B7-1 expression by APCs. LDLR–/– mice were fed control diet (con) or cholesterol-containing diet for 3 or 10 weeks before sacrifice and tissue harvesting. Iliac lymph node cell suspensions (A) and peritoneal macrophages (B) were stained with fluorescent antibodies specific for CD11c and F4/80, respectively, and both populations were also stained with fluorescent antibodies specific for PD-L1, PD-L2, and B7-1. Values represent the percentage of the CD11c+ or F4/80+ cells that stained positive for the indicated antibodies. *P < 0.05, †P < 0.01 versus control diet. n = 4 per group. Data are from 1 of 2 experiments with similar results. (C) ORO staining for lipid in peritoneal macrophages isolated from mice on cholesterol or control diets. Original magnification, ×100.
with control diet–fed mice (Figure 4B). An increase was also noted in PD-L2 expression after high-cholesterol diet feeding. There was no significant difference in macrophage B7-1 expression. The macrophages isolated from mice fed high-cholesterol diet were lipid-loaded foam cells, as confirmed by lipid staining (Figure 4C). These cells have been studied as accessible surrogates for the lipid-laden macrophages present in atherosclerotic lesions (22). We did not find any significant high-cholesterol diet–induced changes in the expression of PD-L1, PD-L2, or B7-1 on DCs or macrophages isolated from spleen (data not shown).

Effect of PD-L1/2 deficiency on T cell activation in vivo. In order to evaluate the changes in T cell activation in the absence of PD-L1/2 in the context of atherosclerosis, we examined iliac lymph nodes and spleens from PD-L1/2–/–LDLR–/– mice fed a high-cholesterol diet compared with LDLR–/– control mice fed the same diet. After 5 weeks of high-cholesterol diet, the iliac lymph nodes appeared larger in PD-L1/2–/–LDLR–/– mice than in control LDLR–/– mice. This was confirmed by increased average cell content in these lymph nodes (Figure 5A). The total number of CD4+ T cells in the lymph nodes increased in hypercholesterolemic PD-L1/2–/–LDLR–/– mice compared with controls. A higher fraction of the CD4+ T cells from PD-L1/2–/–LDLR–/– mice had an activation phenotype (CD25+CD62Llo) compared with CD4+ T cells from control mice (Figure 5B). Other lymph nodes were not examined. The PD-L1/2–/–LDLR–/– mice also had enlarged spleens with increased total cell numbers (Figure 5C), as well as a higher fraction of CD4+ T cells with an activated cell-surface phenotype (Figure 5D), similar to what was seen in the iliac lymph nodes. Differences in the number and activation phenotypes of CD4+ T cells were also observed after 10 weeks of high-cholesterol diet (data not shown).

Serum levels of cytokines. We looked for evidence of systemic immune activation in PD-L1/2–/–LDLR–/– mice by examining serum levels of pro- and antinflammatory cytokines after 10 weeks of high-cholesterol diet. We found a significant increase in TNF-α in the serum of PD-L1/2–/–LDLR–/– mice compared with the control mice (Figure 5E), but no significant differences in the serum levels of IFN-γ, IL-10, IL-12, IL-6, or MCP-1. When we examined the ratio of serum TNF-α in high-cholesterol diet–versus control diet–fed mice, we found that high-cholesterol diet–induced elevation of serum TNF-α was seen only in the setting of PD-L1/2 deficiency (Figure 5F).

Effect of PD-L1/2 deficiency on APC activation of T cells. We examined the effects of PD-1 ligand deficiency on antigen-specific activation of T cells, using a well-characterized TCR transgenic T cell model system. Naive CD4+ OVA-specific TCR transgenic T cells were isolated from OT-II mice (23) and were stimulated in vitro with peritoneal macrophages from mice lacking PD-L1/2. We first evaluated differences without lipid loading, and then further evaluated whether differences were evident after loading the macrophages for 5 hours with acetylated LDL (acLDL) prior to stimulating T cells as previously described (24). As expected, peritoneal macrophages isolated from PD-L1/2–/– mice caused increased CD4+ T cell proliferation compared with control mice (Figure 6A). Importantly, this was also evident after lipid loading. This increase occurred when T cells were stimulated with the polyclonal stimulus anti-CD3 as well as with 5 μg/ml of cognate OVA peptide antigen. However, at a higher antigen concentration (10 μg/ml) plus lipid loading, there was no significant influence of PD-L1/2 deficiency. This is consistent with previous data showing that the downregulating effect of PD-L1/2 on CD4+ T cells is greatest at lower antigen concentrations (11). Because DCs have been implicated in T cell responses in atherosclerosis (25, 26), we also performed studies with PD-L1/2–/–deficient DCs. As seen with the macrophages, there was an increased T cell proliferative response using DCs isolated from PD-L1/2–/– deficient mice compared with control mice, with or without lipid loading. This was evident at a minimal DC:T cell ratio of 1:10 (Figure 6B). However, at a DC:T cell ratio of 1:1, this effect of PD-L1/2 deficiency was not apparent (data not shown), again suggesting that the inhibitory effect of PD-L1/2 on T cell activation can be overridden by

Figure 5
PD-L1/2 deficiency results in enhanced immune responses in hypercholesterolemic mice. (A and C) The number of total cells and cells in the CD4+ T subset was evaluated by FACS staining of cell suspensions prepared from the iliac lymph nodes (A) and spleens (C) of PD-L1/2–/–LDLR–/– and LDLR–/– mice after 5 weeks of high-cholesterol diet. (B and D) The expression of T cell activation markers CD25 and CD62L in cells from the iliac lymph nodes (B) and spleens (D) from mice of each experimental group were determined by FACS analysis. Percentages were evaluated on gated CD4+ T cells. n = 6 per group. (E) Serum levels of the proinflammatory cytokine TNF-α at time of sacrifice were determined. n = 10–14 per group. (F) Ratio of the serum TNF-α level in mice fed high-cholesterol diet to that of mice fed control diet in both PD-L1/2–/–LDLR–/– and LDLR–/– mice. n = 3 per genotype and diet group. Data are mean ± SEM. *P < 0.05, **P < 0.01 versus control.
stronger antigenic stimuli. IFN-γ secretion in these T cells was also increased in response to activation by DCs from PD-L1/2−/− mice (Figure 6D). This was evident even at high stimuli that did not show difference in proliferative response (DC/T cell ratio of 1:1; data not shown), suggesting that the effector function of the T cells is tightly regulated by PD-L1/2 even when proliferation is overridden by strong activation stimuli.

We also analyzed the influence of PD-L1/2 on T cell activation when mixed splenic APCs were freshly isolated from hypercholesterolemic mice. In these experiments, the T cell–depleted splenocytes were isolated from PD-L1/2−/−LDLR−/− or LDLR−/− control mice after 10 weeks of high-cholesterol diet. These APCs were cultured with CD4+ T cells isolated from hypercholesterolemic LDLR−/− mice in order to ensure the presence of oxidized LDL–specific T cells. We found these CD4+ T cells had a significantly greater proliferative response to anti-CD3 in the presence of PD-L1/2−/−LDLR−/− APCs compared with those from LDLR−/− mice (Figure 7A). The increase in the proliferation response was also seen in response to oxidized LDL (Figure 7B), although the proliferative response was relatively weak. The increased response was also evident when analyzing IL-2 and IFN-γ secretion by T cells stimulated with PD-L1/2−/− APCs plus anti-CD3 or OVA peptide (Figure 7, D and E).

Discussion

This study demonstrates a profound influence of the PD-L1/2 molecules on T cell responses and atherosclerosis in hypercholesterolemic mice. Without inflammatory challenges, PD-L1/2−/− mice do not show overt manifestations of dysregulated immunity for many months. However, in the setting of infections, allografts, or pre-existing susceptibility to autoimmunity, the absence or blockade of PD-L1, PD-L2, or PD-1 result in enhanced T cell responses and disease (18, 27–30). Our results indicate that hypercholesterolemia and the resulting arterial pathology represent another systemic immune challenge that is regulated by the PD-L1/2 pathway. We chose to study the effects of combined deficiency of PD-L1 and PD-L2 because the differential roles of these molecules remain unclear, and redundant functions might minimize the net effect of a single gene deficiency. Furthermore, our previous work with cultured mouse endothelial cells and mouse hearts indicates that PD-L1 plays a dominant role in regulating T cell responses to...
cardiovascular antigens (20). Nonetheless, differential effects of PD-L1 and PD-L2 have been demonstrated in some mouse disease models, albeit in strain-specific manners. These diseases include cutaneous Leishmaniasis (28), EAE, and diabetes (29). This raises the possibility that combined deficiency of PD-L1 and PD-L2 may obscure opposing effects of each molecule. The marked effects of combined PD-L1 and PD-L2 deficiency on APC function and on atherosclerosis we observed in this study support the conclusion that there are not opposing influences of these 2 molecules of comparable magnitude on the immune responses that affect atherosclerosis, at least in the C57BL/6 strain.

Our finding that deficiency of PD-L1/2 increased atherosclerosis is comparable to the exacerbating effects of deficiency of this pathway in other immune-mediated diseases such as EAE and insulin (18, 19, 31). Specifically, the data from this study would suggest that the effect of PD-L1/2 deficiency is mediated largely through dysregulated T cell activation by APCs. We demonstrated this in vitro with normal and lipid-loaded DCs and macrophages as well as with APCs freshly isolated from atherosclerotic mice. In all cases, there was more proliferation in the absence of PD-L1/2; more importantly, there was increased secretion of IFN-γ, a cytokine that has previously been shown to have a crucial role in the propagation of atherosclerosis (7). An alternative way in which PD-L1/2 deficiency might lead to enhanced proatherogenic T cell responses is by impairment of Treg development or function. B7 family molecules have been shown to support natural Treg-mediated suppression of proatherogenic immune responses (16, 17). Although PD-L1 has been implicated in the development of type 1 Tregs (32), there are no data clearly indicating a role for PD-L1 or PD-L2 in natural Treg suppression. Furthermore, using an in vitro assay, we did not observe an impairment in the suppressive function of natural Tregs isolated from PD-L1/2-deficient mice (data not shown).

The site where PD-L1/2 may be inhibiting proatherogenic T cell responses remains unclear. We show here that hematopoietically derived APCs from spleen and paraaortic lymph nodes relied on PD-L1/2 to limit T cell activation. This has also been shown in other models of immune-mediated diseases (18). However, there are data to suggest that PD-L1 may have an inhibitory function within tissue inflammatory sites (19). We have previously shown that PD-L1 is expressed on endothelial cells and inhibits T cell activation (20). In addition, PD-L1 is expressed on microvascular endothelial cells (21, 33) as well as on pancreatic islet cells (31) and tubular epithelium (34). Although we found PD-L1 in the neointima of atherosclerotic lesions, this likely reflects expression on lesional DCs or macrophages. It is possible that the increased number of T cells we observed in the lesions of PD-L1/2-deficient mice is a result of increased recruitment, increased local proliferation, or both. The possibility that PD-L1/2 on APCs within the plaque may regulate local effector T cell proliferation or effector responses requires further study.

Although a significant body of literature supports the hypothesis that CD4+ T cell responses promote atherogenesis (5), the effects of hypercholesterolemia are not uniformly conducive to enhanced T cell-mediated immunity (35, 36). Furthermore, the role of different T cell subsets in atherogenesis is complex (37), with many questions unresolved. Our finding that CD8+ T cell infiltration in atherosclerotic lesions was markedly enhanced in PD-L1/2-deficient mice is of interest, because CD8+ T cells are usually scarce in lesions compared with CD4+ T cells (1). Recent studies have shown that PD-1 is upregulated on CD8+ T cells in the setting of chronic viral infections, leading to an “exhausted” phenotype characterized by functional impairment of the T cells (28, 38–40). Blockade of PD-1 can reverse this exhausted phenotype (38, 39). The CD8+ T cell infiltration in atherosclerotic lesions of PD-L1/2-deficient mice suggests that a similar mechanism of reversible inhibition of CD8+ T cell responses exists in the chronic inflammatory milieu of atherosclerosis.

The PD-1/PD-L pathway is emerging as a major regulator of T cell responses to chronic infectious and autoimmune diseases. Our data demonstrate, for the first time to our knowledge, that...
this pathway affects the systemic T cell response to hypercholesterolemia and the associated development of arterial disease. This knowledge should be considered for the development of immunomodulatory approaches to atherosclerosis.

Methods

Mice. PD-L1–/–PD-L2–/– mice on a C57BL/6 background (10) were bred with LDLR–/– mice to produce the triple-knockout PD-L1/2–/–LDLR–/– mice used in this study. LDLR–/– mice backcrossed 10 times onto a C57BL/6 background (The Jackson Laboratory) were used as control mice in these studies. OT-II mice, which express a class II MHC (I-Eα) restricted transgenic TCR specific for OVA peptide (OVA251-264), were provided by W.R. Heath and F. Carbone (Walter and Eliza Hall Institute of Medical Research, Melbourne, Victoria, Australia) (23). All mice were housed and bred in accordance with the institutional guidelines and were approved by the institutional review board of Brigham and Women’s Hospital and Harvard Medical School.

Aortic atherosclerotic lesion analysis. Atherosclerotic lesions were analyzed in the aortic root, aortic arch, and descending aorta as previously described (17). Alternate cryosections (5 μm thick) throughout the aortic sinus (total distance covered, approximately 200 μm) were taken for analysis. From the area in which 3 aortic valve cusps are clearly seen, alternate sections were collected for quantification. Longitudinal cryostat sections of the aortic arch, and formalin fixed, pinned-out en face preparations of the descending aorta were prepared as described previously (15, 41). Aortic lesions were stained with ORO according to the method of Paigen et al. (42). Images of sections were captured digitally (SV Micro) and quantified using IMAGE-PRO PLUS software (Media Cybernetics). For aortic root sections, the plaque lesion area and percent of the total cross-sectional vessel wall area were quantified, and the results were expressed as the mean of 6 sections per mouse. Quantification of aortic arch and abdominal aortic lesions was performed as described (15, 41). Measurements and evaluation of the atherosclerotic lesions were performed in a blinded fashion.

Immunohistochemical and immunofluorescence staining. For immunohistochemical analysis, serial cryostat sections of aortic sinus adjacent to the ORO-stained sections were stained with the respective molecule-specific antibodies, as described previously (21, 43). Antibodies for immunohistochemistry included rat anti-mouse Mac-3 for macrophage identification (1:1000; BD Biosciences—Pharmingen), and anti-CD4+ for T cells (1:120; BD Biosciences—Pharmingen). For mouse smooth muscle cell–actin staining, primary antibody (FITC-conjugated α-actin, 1:500; Sigma-Aldrich) was applied, followed by anti-FITC biotin-conjugated secondary antibody (1:400; Sigma-Aldrich). For PD-L1 staining we used purified rat anti-mouse B7-H1 mAb clone 10F.9G2 (33). Quantitative analysis of lesional content of macrophages, smooth muscle cells, and collagen was determined by computer-assisted image analysis (15, 17) and expressed as percentage of intimal area in order to normalize for overall differences between the study groups. Quantification of CD4 staining was accomplished by counting individual positively stained lesional cells, which are easily resolved, in the aorta-sinus sections.

For immunofluorescence staining, serial cryostat sections (6 μm) of aortic roots were fixed in acetone (–20°C, 5 minutes), air-dried, and blocked with 1% BSA in PBS. All primary antibodies were from BD Biosciences—Pharmingen except anti–PD-L1 (see above). For colocalization of CD3+ T cells with CD4+ or CD8+ subsets, sections were stained with rat anti-CD4 or rat anti-CD8 antibodies, followed by biotinylated rabbit anti-rat (mouse absorbed; Vector Laboratories) secondary antibody and then streptavidin-FITC (1:100; Amersham Corp.). Secondary stains were performed with rhodamine-conjugated rat anti-mouse CD3. For colocalization of PD-L1 and either CD11c or Mac3, sections were first stained with rat anti-mouse PD-L1 (described above) and then incubated with biotinylated rabbit anti-rat secondary antibody (Vector Laboratories), followed by FITC-streptavidin (Amersham Corp.). After blocking with an avidin/biotin blocking kit (Vector Laboratories), the sections were subsequently stained with hamster anti-mouse CD11c or Mac3, followed by biotinylated anti-hamster cocktail and Texas red–conjugated streptavidin (Vector Laboratories).

Serum cholesterol analysis. Overnight fasting blood from individual mice collected after 8 weeks of cholesterol-enriched diet was allowed to clot; the serum fraction was microcentrifuged and stored at –20°C until specific assays were performed. Total serum cholesterol levels, triglycerides, and plasma lipoproteins were analyzed by an online dual enzymatic method for simultaneous quantification of cholesterol and triglycerides by HPLC at SkyLight Biotech Inc. according to the procedure as described by Usui et al. (44) and expressed in mg/dl.

Serum cytokine level. Serum from mice at the end of the experiment was analyzed for cytokine levels using flow cytometry–based cytokine bead assays (BD Biosciences—Pharmingen) (45).

Isolation of DCs and macrophages. DCs were isolated from minced and collagenase-treated spleens, using a commercially available kit that relies on anti-CD11c–coated magnetic beads (Miltenyi Biotec). The purity of these cells was greater than 90% CD11c+ by FACS analysis, and greater than 90% of the CD11c+ cells were also positive for class II MHC. Peritoneal macrophages were isolated by peritoneal lavage with cold PBS. The purity of these cells was +90% F4/80+ by FACS analysis.

Lipid loading of DCs and peritoneal macrophages. Isolated DCs and peritoneal macrophages were incubated for 5 hours in tissue culture medium with added acLDL (100 μg/ml; Biomedical Technologies Inc.) and the acyl-CoA cholesterol acyltransferase (ACAT) inhibitor Compound 58035 (3-[decyldimethylsilyl]-N-[2-(4-methylphenyl)-1-phenylethyl] propanamide; 10 μg/ml) (24).

In vitro and ex vivo assays of CD4+ proliferation and cytokine secretion. For isolation of CD4+ T cells from experimental mice on high-cholesterol diet or OT-II TCR transgenic mice, spleens were removed from the mice and CD4+ T cells were isolated by anti-CD4 Microbeads (Miltenyi Biotec). The cells were stimulated in 96-well cultures (1 × 105 cells/well) with plate-bound anti-CD3e (145-2C11; BD Biosciences—Pharmingen), human copper-oxided LDL (Biomedical Technologies Inc.), or OVA peptide (OVA251-264) plus irradiated T cell–depleted spleen cells (1 × 105 cells/well) as APCs. Medium alone was used as a control. For in vitro stimulation with peritoneal macrophages or DCs, the T cells were stimulated at the ratios indicated in Results and in the figure legends. Culture supernatants were removed at 48 hours and analyzed by flow cytometry–based cytokine bead assays (BD Biosciences—Pharmingen) of culture supernatants for IFN-γ and IL-2 cytokines (45). Cultures were assayed for proliferation after 64 hours by [3H]-thymidine uptake (1 μCi/well), added 16 hours before harvest.

Flow cytometry. We performed 2- or 3-color flow cytometry by standard protocol. Briefly, 0.5–1 × 106 cells from spleens were incubated in 100 μl staining buffer (PBS with 1% BSA and Fcγ III/II Receptor blocking antibody (2.4G2; BD Biosciences—Pharmingen) for 10 minutes. The cells were then stained with a mixture of PE, FITC, PE-Cy5+, or APC-conjugated mAb (0.5 μg/sample) for 15 minutes, washed twice, then fixed in PBS with 1% formaldehyde. The analysis was performed on a FACSCalibur flow cytometer (BD Biosciences) with CellQuest software (BD Biosciences). All procedures were performed on ice until analysis. The mAbs for flow cytometry were as follows: APC-conjugated F4/80 (Caltag/Invitrogen), APC-conjugated CD11c (BD Biosciences), PE-conjugated PD-L1 (BD Biosciences), FITC-conjugated PD-L2 (BD Biosciences), FITC-conjugated B7-1 (BD Biosciences), PE-conjugated anti-CD25 (PC61; BD Biosciences), and FITC-conjugated anti-CD4 (L3T4; BD Biosciences). The fraction of positive-staining cells was determined relative to isotype control–stained cells.

Statistics. All statistical analyses were performed using Prism software. Differences between mice were analyzed by Student’s t test and expressed as mean ± SEM or by the Mann-Whitney test (for nonparametric data).

Differences among multiple groups were determined by the Kruskal-Wallis test and the Dunn’s multiple-comparison test. A P-value of 0.05 or less was considered significant for all analyses.

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