B7-H1 limits the entry of effector CD8+ T cells to the memory pool by upregulating Bim

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Abbreviations: BrdU, bromodeoxyuridine; CFSE, carboxyfluoresceinsuccinimidyl ester; CTL, cytotoxic T lymphocyte; KbOVA-tet, KbOVA257–264 tetramer; OVA, ovalbumin; Poly I:C, polyinosinic-polycytidylic acid

B7-H1, first identified by our laboratory in 1998,8 is expressed by a variety of tumor cells. B7-H1 is also constitutively expressed by macrophages and dendritic cells, its expression being upregulated upon cell activation.9 Classically, B7-H1 is thought to deliver inhibitory signals to CD8+ T cells by binding to PD-1, resulting in negative regulation of TCR signaling and apoptosis of activated T cells.10-12 Recently, CD80 (B7-1) was identified as an additional binding partner for B7-H1.13 The exact mechanism whereby B7-H1 leads to apoptosis of effector CD8+ T cells is unknown. Despite many studies of the B7-H1/PD-1 pathway in regulating activated T cells interacting with tumor cells during the effector phase of the immune response,13,26,28 the mechanism by which B7-H1 expressed by host cells regulates primary T-cell responses remains unresolved. Using B7-H1-deficient mice, we and others have reported that host B7-H1 limits the expansion and survival of effector CD8+ T cells.25,37 However, it is not clear whether increased accumulation of effector CD8+ T cells in B7-H1-deficient mice is due to decreased apoptosis or increased proliferation of effector CD8+ T cells in vivo. The role of B7-H1 in regulation of memory T-cell responses also remains unclear.

A primary CD8+ T-cell response is characterized by an expansion phase followed by a contraction phase, during which...
a majority of effector CD8$^+$ T cells are deleted. Importantly, a small population of antigen-specific effector CD8$^+$ T cells survives and develops into the memory population. The contraction phase must be tightly regulated: having too many cells survive contraction would be unfavorable for the host, and could lead to autoimmunity. Conversely, having too few cells survive contraction could compromise the establishment of a protective memory response.\textsuperscript{14}

The regulation of apoptosis in effector CD8$^+$ T cells has been the focus of numerous studies. While the contribution of the death-receptor pathway of apoptosis during contraction remains controversial,\textsuperscript{15-18} the involvement of the mitochondrial pathway of apoptosis is well established.\textsuperscript{19-21} Bcl-2 family members regulate the mitochondrial pathway of apoptosis, and some of these proteins have been characterized as the main regulators of apoptosis in activated T cells. Bcl-2, the most important anti-apoptotic molecule in activated T cells, functions by binding to the pro-apoptotic BH3-only molecule Bim.\textsuperscript{22} At the peak of activation, Bcl-2 levels significantly decline,\textsuperscript{23} allowing free Bim molecules to activate Bax/Bak-like molecules. Upon activation, Bax/Bak-like molecules form pores in the mitochondrial membrane, leading to the release of cytochrome $c$, activation of caspases and initiation of apoptosis.\textsuperscript{24} Fewer T cells undergo contraction and more memory T cells form in Bim-deficient mice,\textsuperscript{25,26} indicating that Bim could be a key regulator in limiting effector T-cell entry into the memory T-cell pool. How Bim levels are regulated in effector T cells following antigen-stimulation is not fully understood.

In this study, we sought to investigate the mechanism by which B7-H1 regulates apoptosis in effector CD8$^+$ T cells and subsequently to address the effect of B7-H1 on memory generation. We used a non-proliferative immunization approach and performed ex vivo assays to directly examine the levels of pro- and anti-apoptotic molecules in antigen-specific CD8$^+$ T cells. We found that more memory CD8$^+$ T cells were generated in B7-H1-deficient mice following immunization as compared with wild type (WT) mice. At the peak of the expansion phase, CD8$^+$ T cells expressed lower levels of Bim in B7-H1-deficient mice than in WT mice. In vitro assays revealed that stimulation by plate-bound B7-H1 led to Bim upregulation in activated CD8$^+$ T cells, and antibodies which inhibit the B7-H1/ PD-1 or B7-H1/CD80 interaction blocked this upregulation. Our results suggest that B7-H1 may be a previously unknown regulator of Bim expression, and B7-H1 may negatively regulate CD8$^+$ T-cell memory by enhancing the depletion of antigen-primed CD8$^+$ T cells through the upregulation of Bim.

Results



Figure 1. Kinetics of CD8$^+$ T-cell responses to antigen stimulation. Wild type (WT) and B7-H1-deficient (KO) mice were immunized (i.p.) with OVA plus poly I:C. K$^b$/OVA tetramer was used to identify antigen-specific CD8$^+$ T cells in spleen and liver at the indicated times after immunization. Data show the percentage of tetramer$^+$ CD8$^+$ T cells (mean ± SD of three mice per time point). One of two independent experiments is shown. *p < 0.05 compared with WT mice.

More memory T cells are generated in the absence of B7-H1. We compared the kinetics of CD8$^+$ T-cell responses in the spleen and liver of WT and B7-H1-deficient C57BL/6 mice following immunization with ovalbumin (OVA) protein and polyinosinic:polycytidylic acid (poly (I:C)) as adjuvant. We observed an increased number of CD8$^+$ T cells at the peak of the immune response (day 7-post immunization) in the spleen and liver of B7-H1-deficient mice as compared with WT mice. During the contraction phase (days 7 to 14-post immunization) there was a significant delay in the reduction of antigen-specific CD8$^+$ T cells in the spleen and liver of B7-H1-deficient mice as compared with WT mice. On day 40 following immunization, more antigen-specific memory CD8$^+$ T cells were detected in B7-H1-deficient mice as compared with WT mice (Fig. 1). These data suggest that host B7-H1 may regulate the extent of expansion and contraction of effector CD8$^+$ T cells, thus influencing the size of the memory CD8$^+$ T-cell pool in both lymphoid and non-lymphoid tissues.

We examined to what extent B7-H1 regulates the generation of memory CD8$^+$ T cells in immunized mice by using K$^b$/OVA$^{257-264}$ tetramer (K$^b$/OVA-tet) to detect antigen-primed memory CD8$^+$ T cells in the spleen on day 4 after in vivo restimulation (OVA protein, administered on day 40 after primary immunization). Day 4 was selected for analysis because at this time point it is possible to distinguish a recall response (which takes 7 d to establish). Thus, naïve mice did not show a significant increase of antigen-specific CD8$^+$ T cells on day 4 after immunization (Fig. 2A). The frequency of K$^b$/OVA-tet$^+$ CD8$^+$ T cells increased more than 2-fold in immunized B7-H1-deficient mice (0.38%) as compared with WT mice (0.16%; $p < 0.05$; Fig. 2A). This increase was reflected in the absolute cell numbers ($p = 0.001$; Fig. 2B). In addition to having increased numbers of memory CD8$^+$ T cells, an increased...
A hallmark of memory CD8+ T cells is their rapid recall response to cognate antigens, so we next examined if the increased memory pool in B7-H1-deficient mice would lead to a more protective recall response. We injected B16-OVA melanoma tumor cells (engineered to express OVA) into immunized WT and B7-H1-deficient mice. Intravenously injected B16-OVA tumor cells form metastases in the lung, and antitumor immunity can be monitored by counting the number of tumor foci. On day 4 following intravenous injection of 5 × 10^5 B16-OVA tumor cells, the frequency of functional memory CD8+ T cells in the lungs of WT and B7-H1-deficient mice was determined by intracellular staining for IFN-γ. We detected ~4 to 5-fold more IFN-γ+ CD8+ T cells in the lungs of B7-H1-deficient mice as compared with WT mice (p < 0.01; Fig. 3A). On day 21-post tumor injection the number of tumor metastases in the lungs of naïve B7-H1-deficient mice was comparable to that of naïve percentage of memory CD8+ T cells capable of producing multiple cytokines was detected in the spleens of B7-H1-deficient mice (0.73% IFN-γ/TNF-α, 0.17% IFN-γ/IL-2) as compared with WT mice (0.24% IFN-γ/TNF-α, 0.07% IFN-γ/IL-2; p < 0.05; Fig. 2C). We also performed an in vivo CTL assay to measure cytolytic activity of the memory CD8+ T cells. On day 4 after in vivo re-stimulation we injected OVA peptide- or control peptide-pulsed target cells (syngeneic splenocytes labeled with either high or low CFSE) into immunized WT and B7-H1-deficient mice. Four hours following cell injection, we analyzed the remaining CFSE positive cells in the spleen. Memory CD8+ T cells in the B7-H1-deficient mice lysed more OVA-peptide pulsed target cells (33.5%) than those in WT mice (9.3%, p < 0.01; Fig. 2D). Collectively, these data suggest that B7-H1 negatively regulates the generation of memory CD8+ T cells in immunized mice.

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the concept that the CD11a high CD8 + T-cell population represents true OVA-specific CD8+ T cells. Nearly 80–90% of OVA-induced CD11ahigh CD8+ T cells were reactive against undefined antigen epitopes of the OVA protein (Fig. 4C). Therefore, the CD11ahigh CD8+ T-cell population could be used to represent a majority of the antigen-primed CD8+ T cells during primary T-cell responses. In the following studies we used CD11a high as a marker to track antigen-specific CD8+ T cells.

We examined the proliferation of effector CD8+ T cells following immunization by staining cells for Ki67, a nuclear protein associated with cell proliferation.28 The percent of Ki67+ cells increased in CD11ahigh CD8+ T cells from B7-H1-deficient mice (9.32%) as compared with WT mice (7.5%), but this increase was not statistically significant (Fig. 5A). We also monitored proliferation by performing a BrdU incorporation assay to measure the ongoing proliferation of CD8+ T cells following immunization. Again, we found that the percentage of BrdU+ CD11ahigh CD8+ T cells was similar between WT (6.05%) and B7-H1-deficient mice (5.59%; Fig. 5B). Ki67+ or BrdU+ cells were mainly detected in the CD11ahigh CD8+ T cells but not in CD11alow CD8+ T cells, suggesting that CD11ahigh CD8+ T cells are proliferating following antigen-stimulation (Fig. 5A and B). Our results suggest that the observed increased population of antigen-primed CD8+ T cells in B7-H1-deficient mice is not due to an increased proliferation of this cell compartment, as compared with WT mice.

We next asked if decreased apoptosis of antigen-primed CD8+ T cells could contribute to the observed increased population of antigen-primed CD8+ T cells in immunized B7-H1-deficient mice. As discussed earlier, the Fas/Fas ligand death receptor pathway is implicated in regulation of T-cell contraction, so we measured the surface expression levels of Fas and Fas ligand on effector CD8+ T cells on day 7 after immunization. Expression of Fas and Fas

**Figure 3.** Enhanced memory CD8+ T-cell recall responses and improved antitumor immunity in the lung in the absence of B7-H1. On day 35 after immunization, immunized or naïve WT and B7-H1-deficient mice were injected (i.v.) with 5 × 10^5 B16-OVA tumor cells. (A) Percentage and absolute numbers of IFNγ+ CD8+ T cells in the lung of immunized mice (mean ± SD, n = 3) on day 4 after tumor injection. *p < 0.01 compared with WT mice. (B) Metastatic tumor foci in the lung tissue were identified and counted on day 20 after tumor injection (mean ± SD, n = 5). One of two independent experiments is shown. N.S.: not significant.

WT mice (p = 0.43; Fig. 3B). Fewer tumor metastases formed in the lungs of immunized WT mice as compared with naïve WT mice (p = 0.001). Significantly, tumor metastases were completely rejected in the lungs of immunized B7-H1-deficient mice (Fig. 3B), suggesting that a more efficient CD8+ T-cell memory population is established in the absence of B7-H1.

Bim expression is reduced in antigen-primed CD8+ T cells in the absence of B7-H1. We investigated which mechanisms could be responsible for the increased population of memory CD8+ T cells in B7-H1-deficient mice by examining the proliferation and apoptosis of antigen-primed CD8+ T cells following immunization. We used CD11a as a surrogate activation marker.27 An advantage of this method is that CD11a^high CD8+ T cells represent antigen-primed CD8+ T cells that are responsive to undefined antigen epitopes not recognized by tetramers. CD11a^high CD8+ T cells were detected at low levels in the spleens of naïve WT and B7-H1-deficient mice (Fig. 4A). On day 7 after immunization, the percentage of CD11a^high CD8+ T cells increased more than 2-fold in the spleens of B7-H1-deficient mice (41.5%) as compared with immunized WT mice (17.2%; p < 0.01; Fig. 4A and B), consistent with the results obtained by tetramer staining (Fig. 1). 7–15% of CD11a^high CD8+ T cells from WT and B7-H1-deficient mice were specific for the known H-2Kb-restricted OVA257–264 epitope based on tetramer staining, and CD11a^low CD8+ T cells did not contain tetramer+ cells (Fig. 4C), suggesting that all antigen-specific CD8+ T cells are found in the CD11a^high CD8+ T-cell population. In addition, the CD11a^high CD8+ population from both WT and B7-H1-deficient mice, but not the CD11a^low CD8+ T-cell population, produced IFNγ and underwent degranulation (indicated by CD107a surface expression) following ex vivo re-stimulation (Fig. 4D). As T-cell responses against diverse epitopes are coordinately regulated, these data further support the concept that the CD11a^high CD8+ T-cell population represents true OVA-specific CD8+ T cells. Nearly 80–90% of OVA-induced CD11a^high CD8+ T cells were reactive against undefined antigen epitopes of the OVA protein (Fig. 4C). Therefore, the CD11a^high CD8+ T-cell population could be used to represent a majority of the antigen-primed CD8+ T cells during primary T-cell responses. In the following studies we used CD11a^high as a marker to track antigen-specific CD8+ T cells.

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ligand was detected at similar levels in WT and B7-H1-deficient mice (Fig. S1). These results suggest that the observed increased population of effector CD8+ T cells is not due to a change in Fas-induced apoptosis in B7-H1-deficient mice. We also investigated the mitochondrial pathway for apoptosis by analyzing levels of Annexin V and tetramethylrhodamine ethyl ester (TMRE) staining. TMRE is a fluorescent marker that is incorporated into intact mitochondria, and cells undergoing apoptosis show reduced TMRE staining as compared with live cells.29 Fewer antigen-primed CD11a^{high} CD8+ T cells were undergoing apoptosis (TMRE^{low} Annexin V) in B7-H1-deficient mice (3.4%) as compared with WT mice (6.7%, p < 0.05; Fig. 5C and D). These results suggest that decreased levels of mitochondrial apoptosis may contribute to the observed increased population of antigen-primed CD8+ T cells in B7-H1-deficient mice.

We next looked for alterations in the expression of apoptosis-regulating molecules in effector CD8+ T cells. We measured intracellular levels of the pro-apoptotic molecule Bim, and the anti-apoptotic molecules Bcl-2 and Bcl-xL in CD11a^{high} CD8+ T cells freshly isolated from the spleen on day 7 after immunization of naive mice. We observed lower intracellular expression levels of Bim in CD11a^{high} CD8+ T cells from B7-H1-deficient mice than in the same cells obtained from WT mice (p < 0.001; Fig. 6A and B), while the expression levels of Bcl-2 and Bcl-xL were comparable in WT and B7-H1-deficient mice (Fig. 6A). The expression of Bim, Bcl-2 and Bcl-xL were comparable in CD11a^{low} CD8+ T cells from B7-H1-deficient and WT mice (Fig. 6A). We also analyzed intracellular expression levels of Bim in CD11a^{high} CD8+ T cells isolated from the liver on day 7 after immunization of naive mice. We observed lower intracellular expression levels of Bim in CD11a^{high} CD8+ T cells from B7-H1-deficient mice than in the same cells obtained from WT mice (Fig. 6C), while the expression levels of Bcl-2 and Bcl-xL were comparable in WT and B7-H1-deficient mice (Fig. 6C). Finally, we examined intracellular expression levels of these proteins in CD11a^{high} CD8+ T cells isolated from the spleen of naive mice, and observed no significant differences in B7-H1-deficient vs. WT mice in the expression levels of Bim (Fig. 6D), Bcl-2 or Bcl-xL (data not shown). Our data suggests that the downregulation of the pro-apoptotic molecule Bim may contribute to the observed increased population of antigen-primed effector CD8+ T cells in B7-H1-deficient mice.

To exclude the possibility that the downregulation of Bim in B7-H1-deficient mice would be due to an intrinsic change in B7-H1-deficient T cells, we performed transfer experiments in which we injected naive OT-1 CD8+ T cells (Thy1.1+) into WT or B7-H1-deficient mice (Thy1.2-). Following transfer of the OT-1 CD8+ T cells, host mice were immunized with OVA plus poly I:C. On day 7 after immunization we measured the intracellular levels of Bim, Bcl-2, and Bcl-xL in the transferred OT-1 CD8+ T cells freshly isolated from spleen and liver. OT-1 CD8+ T cells transferred into B7-H1-deficient hosts expressed lower levels of Bim in both the spleen and liver as compared with the OT-1 CD8+ T cells transferred into WT hosts (Fig. 7). The expression of Bcl-2 and Bcl-xL in OT-1 CD8+ T cells transferred into WT or B7-H1-deficient mice was comparable (Fig. 7). These data suggest that the downregulation of Bim in B7-H1-deficient mice is not due to an intrinsic change in B7-H1-deficient T cells, but rather to host B7-H1 interacting with one of its binding partners on CD8+ T cells.

Next, we used antibodies that block the interaction between B7-H1 and PD-1 or between B7-H1 and CD80 to examine if blocking either of these pathways would impact Bim expression levels. On days 1 and 3 after immunization of WT mice with OVA plus poly I:C, we injected an anti-PD-1 antibody (G4) that only blocks PD-1 binding to B7-H1,30 or an anti-B7-H1 antibody (43H12) that only blocks B7-H1 binding to CD80.31
day 7 after immunization we compared Bim expression levels in CD11a<sup>high</sup> CD8<sup>+</sup> T cells between groups with or without antibody blockade. Antibodies blocking the interaction between B7-H1 and PD-1 or between B7-H1 and CD80 both reduced the expression of Bim in primed CD8<sup>+</sup> T cells as compared with control antibodies, whereas the expression of Bcl-2 and Bcl-x<sub>L</sub> remained unaffected (Fig. S2). Our results suggest that the downregulation of Bim in B7-H1-deficient mice might be due to a lack of interaction between B7-H1 and its binding partners PD-1 and CD80.

The influence of PD-1 signaling on memory generation was addressed previously by Allie et al. They demonstrated that after an acute viral infection more central memory T cells accumulate in the lymphoid organs of PD-1-deficient mice as compared with WT mice, indicating that PD-1 signaling negatively regulates memory T-cell generation. We addressed the relevance of CD80 signaling in the regulation of memory generation by transferring equal numbers of CD80-deficient OT-1 and WT OT-1 naïve CD8<sup>+</sup> T cells into CD45.1<sup>+</sup> mice. One day after T-cell transfer, host mice were immunized with OVA plus poly I:C. On day 21 after immunization the frequency and phenotype of the transferred CD80-deficient and WT OT-1 CD8<sup>+</sup> T cells was analyzed. On day 21 after immunization, a 2-fold increased percentage of CD80-deficient OT-1 CD8<sup>+</sup> T cells as compared with WT OT-1 CD8<sup>+</sup> T cells was detected in the spleen (Fig. S3A), indicating that the transferred CD80-deficient OT-1 CD8<sup>+</sup> T cells generated more memory T cells as compared with WT OT-1 CD8<sup>+</sup> T cells. Surface staining confirmed that these cells had a central memory phenotype (CD44<sup>hi</sup> CD62L<sup>hi</sup>, Fig. S3B). We investigated the recall response of the memory population generated from transferred cells by injecting the hosts with OVA plus poly I:C on day 30 after the initial immunization, and 3 d later the frequency and phenotype of the transferred cells was analyzed. An increased percentage of CD80-deficient OT-1 CD8<sup>+</sup> T cells as compared with WT OT-1 CD8<sup>+</sup> T cells was detected in the spleen (p = 0.013; Fig. S3C and D). Surface staining confirmed
that these cells had an effector memory phenotype (CD44hi CD62Llo, Fig. S3E). Taken together, these data demonstrate that Cd80−/− OT-1 CD8+ T cells generated more memory T cells as compared their WT counterparts, indicating that CD80 expressed by CD8+ T cells may negatively regulate memory T-cell generation.

**B7-H1 enhances Bim expression in activated CD8+ T cells.** We investigated how B7-H1 could regulate Bim levels in activated CD8+ T cells by incubating pre-activated WT CD8+ T cells with plate-bound B7-H1 fusion protein for 48 h in the presence of TCR stimulation (anti-CD3 antibody). We analyzed Bim expression by western blot and found increased expression levels in CD8+ T cells cultured in the presence of B7-H1 fusion protein compared with a control fusion protein (Fig. 8A). We also analyzed Bim expression by intracellular flow cytometry and observed that B7-H1 fusion protein dramatically increases the levels of Bim protein in CD8+ T cells compared with a control fusion protein (p < 0.02; Fig. 8B and C). In the absence of anti-CD3 antibodies, Bim levels did not increase upon incubation with B7-H1 fusion protein (data not shown), suggesting that B7-H1 provides a co-stimulatory signal for Bim upregulation. Accordingly, the absolute number of live cells was also reduced in CD8+ T cells cultured in the presence of B7-H1 fusion protein compared with a control protein (p < 0.01; Fig. 8D). We observed increased levels of cells undergoing apoptosis (TMRElow Annexin V+) in cultures of activated CD8+ T cells exposed to the B7-H1 fusion protein and anti-CD3 (12.4%) as compared with cells cultured with control fusion protein and anti-CD3 (4.1%, Fig. 8E). The induction of apoptosis by B7-H1 fusion protein was lost in CD8+ T cells isolated from Bim-deficient and Bcl-2 transgenic mice (Fig. 8E), suggesting that B7-H1-induced T-cell apoptosis may be dependent on the Bim-mediated mitochondrial pathway of apoptosis.

To examine which receptor of B7-H1 is involved in mediating Bim upregulation, we incubated pre-activated WT CD8+ T cells with plate-bound B7-H1 fusion protein pre-blocked with anti-B7-H1 (10B5 or 43H12) or anti-PD1 (G4) antibodies. The 10B5 antibody blocks the interaction of B7-H1 with both PD-1 and CD80. Both 10B5 and G4 antibodies completely blocked Bim upregulation induced by B7-H1 fusion protein, while 43H12 only partially, but significantly, did so (Fig. 8F). None of the antibodies used in this experiment
had effects on Bim expression levels in cells cultured with control fusion protein, indicating that their effect on Bim expression levels is due to blocking the interaction between B7-H1/PD-1 or B7-H1/CD80, and not due to a non-specific effect. These results suggest that B7-H1 may use PD-1 or CD80 on CD8+ T cells to deliver co-stimulatory signals for the upregulation of Bim.

We next investigated the mechanism by which B7-H1 regulates Bim expression levels. We analyzed the mRNA levels of Bcl211, which encodes the Bim protein, by quantitative real-time PCR analysis on mRNA isolated from pre-activated CD8+ T cells exposed to B7-H1 fusion protein or control fusion protein and anti-CD3 for 24 h. Incubation of pre-activated CD8+ T cells with B7-H1 fusion protein did not increase the levels of Bcl211 (Fig. 9A), indicating that the B7-H1-mediated upregulation of Bim does not result from transcriptional regulation. The degradation of Bim is tightly regulated, one mechanism being the activation of Akt followed by Akt-mediated Bim phosphorylation and degradation. The level of Akt activation in CD8+ T cells after B7-H1 engagement was measured by intracellular flow cytometry for phosphorylated-Akt (Ser473). CD8+ T cells cultured with B7-H1 fusion protein exhibited decreased levels of phosphorylated Akt as compared with CD8+ T cells cultured with a control fusion protein (p < 0.01; Fig. 9B and C). As phosphorylation of Akt at Ser473 is regulated by activation of mTOR,34,35 we next examined whether B7-H1 regulates phosphorylation of mTOR in vitro. Unexpectedly, there was no difference in levels of phospho-mTOR in CD8+ T cells cultured with B7-H1 fusion protein and cells cultured with control fusion protein (Fig. 9B and C).

Discussion

In this study, we report that B7-H1 negatively regulates the establishment of CD8+ T-cell memory by inducing the upregulation of Bim, resulting in apoptosis of effector CD8+ T cells. Accordingly, B7-H1-deficient mice develop enhanced memory CD8+ T-cell populations following immunization, resulting in improved protective immunity. Bim expression levels are reduced in CD8+ T cells when there is a lack of B7-H1 signaling, either in B7-H1-deficient mice, when WT CD8+ T cells are transferred.
into B7-H1-deficient hosts, or when B7-H1 signaling is inhibited by in vivo administration of blocking antibodies. It appears that both PD-1 and CD80 receptors are involved in the B7-H1-mediated upregulation of Bim. Our results reveal a new mechanism by which B7-H1 regulates CD8+ T-cell responses in vivo, and provide a new direction to pursue in dissecting the signaling events downstream of the receptors for B7-H1.

Recent studies in human patients have indicated that tumor-infiltrating CD8+ T cells express PD-1, resulting in impaired function and compromised antitumor immunity.36,37 Analysis of a variety of patient tumor tissues revealed that a majority of tumors express B7-H1, and this tumor-associated B7-H1 increases the apoptosis of infiltrating T cells.11 B7-H1 expression on renal cell carcinoma cells has also been shown to be associated with poor prognosis.38 These findings demonstrate the clinical relevance of B7-H1 signaling in patients with cancer, motivating our studies of how B7-H1 regulates T-cell memory generation and apoptosis. Bim is a key regulator of T-cell apoptosis during the contraction phase of CD8+ T-cell responses.39 In a study of patients with chronic HCV infection, Larrubia et al. reported increased levels of Bim expression in CD8+ T cells that upregulated PD-1.40 This and other recent studies also reported that blockade of the B7-H1/PD-1 pathway leads to improved functioning of virus-specific CD8+ T cells.41-43 However, none of these studies addressed the interaction between the B7-H1/PD-1 pathway and Bim expression levels as we have done here.

We observed that B7-H1-deficient mice exhibit delayed contraction following antigen immunization (Fig. 1), and this difference could not be attributed to increased proliferation of antigen-primed CD8+ T cells (Fig. 5A and B). We did observe a lower frequency of TMRElow Annexin V+ CD8+ T cells in B7-H1-deficient mice, which prompted us to examine the mitochondrial pathway of apoptosis (Fig. 5C). Both our in vivo and in vitro studies revealed that B7-H1 may upregulate Bim levels in antigen-primed or TCR-activated CD8+ T cells (Figs. 6–8). The B7-H1-mediated induction of apoptosis in activated CD8+ T cells appears to be dependent on Bim expression, because B7-H1 ligation did not induce apoptosis in activated CD8+ T cells isolated from Bim-deficient or Bcl-2-transgenic mice (Fig. 8E). Thus, our studies demonstrate that B7-H1 may induce apoptosis of activated CD8+ T cells via regulation of Bim. In support of this finding, Kryczek et al. reported that Notch/Bcl-2 signaling mediates resistance to apoptosis in human tumor-associated memory T cells.44 Altogether, these studies demonstrate the importance of the Bcl-2 family of pro- and anti-apoptotic molecules in the regulation of antitumor memory T-cell responses.

Our group has recently published a study revealing a cell-intrinsic role for B7-H1 expressed by CD8+ T cells in upregulating the anti-apoptotic molecule Bcl-xL.45 Our data presented here clearly shows that regulation of Bim expression levels in CD8+ T cells is an extrinsic function of B7-H1. This was demonstrated by transferring congenically marked naive WT OT-1 CD8+ T cells into WT or B7-H1-deficient hosts, followed by immunization of the mice and analysis of the resulting effector CD8+ T cells. As shown in Figure 7, WT CD8+ T cells expressed lower levels of Bim upon transfer into B7-H1-deficient hosts as compared with transfer into WT hosts, due to a lack of extrinsic B7-H1 signaling. B7-H1 regulation of Bim expression is likely mediated through B7-H1 interacting with both of its binding partners expressed by CD8+ T cells, PD-1 and CD80, because in vivo antibody blockade of either B7-H1/PD-1 or B7-H1/CD80 interactions resulted in reduced Bim expression levels in CD8+ T cells (Fig. S2). This function of B7-H1 appears to depend on TCR signaling, because in the absence of anti-CD3 stimulation the B7-H1 protein failed to enhance Bim levels in pre-activated CD8+ T cells (data not shown).

Work investigating signaling events downstream of PD-1 have revealed that PD-1 ligation results in inhibition of phosphoinositide-3-kinase (PI3K) activity, leading to reduced activation of Akt.12 Akt inhibits Bim activity by phosphorylating Bim, leading to its degradation.13 Hence, it is likely that B7-H1 signaling through PD-1 inhibits Akt, resulting in enhanced expression of Bim in activated CD8+ T cells. Accordingly, our data shows that B7-H1 engagement of pre-activated CD8+ T cells leads to decreased phosphorylation of Akt (Fig. 9B). Phosphorylation of

Figure 9. B7-H1 co-stimulation inhibits activation of Akt. Pre-activated CD8+ T cells were stimulated with plate-bound B7-H1 or control fusion protein (Fc). After 24 h of stimulation, CD8+ T cells were harvested and used for analysis. (A) Analysis of Bcl211f transcript levels by real-time qPCR using the comparative C_T method. GAPDH served as the internal control gene. Graph shows fold change (mean ± SD, n = 4). (B) Phosphorylation of Akt and mTOR was analyzed by intracellular staining of CD8+ T cells with anti-phospho-Akt and anti-phospho-mTOR Abs. Numbers show percentage of positive stained cells. (C) Bar graph of average MFI of phospho-Akt and phospho-mTOR expression (mean ± SD, n = 3). N.S.: not significant.
Akt following PI3K activation is mediated by PDK1 at Thr308 and mTOR at Ser473. We observed down-regulation of phospho-Akt at Ser473 (Fig. 9B) but not at Thr308 (data not shown), suggesting that B7-H1 may use the mTOR pathway to regulate Akt activation. Recently, Francisco et al. reported that B7-H1 ligation reduces the phosphorylation of mTOR at Ser4248 during the induction or regulatory T cells. However, we could not detect changes in the phosphorylation of mTOR Ser4248 in CD8+ T cells cultured with a B7-H1 fusion protein (Fig. 9B). This discrepancy might be due to the different T-cell subsets studied (CD4+ vs. CD8+) and/or differences in the duration of the culture (18 h vs. 48 h). Nevertheless, the regulation of Akt activation downstream of B7-H1 signaling in T cells warrants further investigation.

Protective immunity is dependent on the generation of durable memory T-cell responses. However, the signals required to induce the differentiation of functional long-lived memory T cells are not fully understood. Most studies of T-cell memory generation support a linear model. According to this model, upon antigen stimulation naïve T cells undergo a massive level of expansion, during which they differentiate into effector T cells. Following clearance of the initiating antigen stimulus, effector T cells enter a contraction phase, during which a majority of them die. The cells that survive the contraction phase further differentiate to make up the antigen-specific memory T-cell population. In support of this model, it has been shown that the transition from naïve to effector to memory T cells is a gradual process, and that a large portion of memory T cells arise from effector T-cell progenitors, as determined by their inherent markers.

Recent studies indicate that negative T-cell regulators affect the differentiation of memory T cells. Blockade of CTLA-4 inhibitory signals results in more memory T cells generated following bacterial infection. In PD-1-deficient mice, Allie et al. observed that more central memory T cells accumulate in the lymphoid organs after acute viral infections. Those studies did not address the molecular mechanisms by which negative regulators influence the process of memory T-cell development. Our studies suggest that B7-H1-mediated upregulation of the pro-apoptotic molecule Bim in effector T cells may be a key factor in limiting effector T-cell entry into the memory pool. In line with previous studies unveiling more memory T cells in Bim deficient mice, our results show that, in the absence of B7-H1, Bim levels are downregulated in effector CD8+ T cells and apoptosis of effector CD8+ T cells is reduced. Thus, more effector CD8+ T cells are allowed to enter the memory pool. Because Bim upregulation by B7-H1 is dependent on PD-1 (Fig. S2), our data may explain why more memory CD8+ T cells are generated in PD-1-deficient mice after acute infection.

In addition, our studies show that, in the absence of B7-H1, a robust antitumor immune response can be established that prevents the formation of tumor metastases in the lung (Fig. 3B). As tumor metastases frequently occur in peripheral tissues, the generation or maintenance of a functional memory T-cell response in the periphery will provide a first line of protection against tumor metastases. Although we did not directly address to what extent B7-H1 influences the maintenance of memory T cells in peripheral tissues, our data strongly indicate that enhancing primary T-cell responses may promote effector T cells to enter the memory T-cell pool. In this regard, B7-H1 may function as a checkpoint in limiting the number of effector T cells that enter the memory pool. Future studies are needed to address which subsets of effector T cells are more sensitive to B7-H1 signals and whether TCR affinity may affect the sensitivity of effector T cells to B7-H1 regulation. This knowledge is important in vaccine design for treatment of cancer and chronic infections for which the B7-H1/PD-1 blockade is being evaluated in Phase I clinical trials.

In summary, our findings suggest that B7-H1 signaling through both PD-1 and CD80 on antigen-primed CD8+ T cells negatively regulates the establishment of memory populations by upregulating expression of the pro-apoptotic molecule Bim. Our results add to the understanding of how memory T-cell generation is controlled by inhibitory molecules, and also have important implications for rational vaccine design and the development of immunotherapies targeting tumor metastases.

Materials and Methods

Mice, cell lines and reagents. Female CD45.2+ C57BL/6 mice were purchased from Taconic Farms and CD45.1+ congenic C57BL/6-Ly5.1 mice were purchased from National Cancer Institute. OT-1 TCR (Thy 1.1+) transgenic mice were provided by T. Tian (Harvard University, Boston, MA). B7-H1-deficient C57BL/6 mice were provided by L. Chen (Yale University, New Haven, CT). Bcl211−/− mice and Cd80−/− mice were purchased from Jackson Laboratory. Cd80−/− mice were crossed into WT OT-1 mice and produced Cd80−/− OT-1 mice. Bcl-2 transgenic mice were provided by V. Shapiro (Mayo Clinic, Rochester). Mice were maintained under pathogen-free conditions and used at 8–12 weeks of age. B16-OVA murine melanoma cells were provided by R. Vile (Mayo Clinic, Rochester, MN), and were cultured in RPMI 1640 medium (Cellgro) with 10% FBS (Life Technologies), 1 U/mL penicillin, 1 μg/mL streptomycin and 20 mM HEPES buffer (all from Mediatech). Hamster anti-mouse B7-H1 mAb (10B5) and PD-1 (G4) was obtained from hybridoma cells provided by L. Chen. Hamster anti-mouse B7-H1 mAb (43H12) was provided by K. Tamada (John Hopkins University). Studies were conducted in accordance with the National Institutes of Health guidelines for the proper use of animals in research and with local Institutional Animal Care and Use Committee approval.

Flow cytometry analysis. Class I MHC (KoOVA peptide SIINFEKL) tetramer and negative control tetramer were purchased from Beckman Coulter. Fluorochrome-conjugated Abs against CD8, CD11a, Fas (CD95), Fas ligand, CD90.1 (Thy 1.1), CD90.2 (Thy 1.2), CD107a, IFNγ, IL-2 and TNFα were purchased from BD Biosciences, BioLegend, or eBiosciences. To detect intracellular cytokine levels cells were incubated with GolgiPlug (BD Biosciences) for 4 h prior to analysis.
Cells were stained for surface antigens, and then incubated in Fixation Buffer (BioLegend) for 20 min at room temperature, followed by permeabilization in Permeabilization Wash Buffer (BioLegend). Fixed and permeabilized cells were then stained with Abs for 20 min at room temperature. Abs to Akt, Bcl-xL, Bcl-2, Bim and mTOR and fluorochrome-conjugated secondary Abs were purchased from Cell Signaling (Danvers, MA). To detect the intracellular levels of Akt, Bcl-xL, Bcl-2, Bim and mTOR, T cells were first stained for surface antigens, then fixed with 2% paraformaldehyde for 10 min at 37°C, followed by permeabilization with ice-cold methanol for 30 min. After blocking with 15% rat serum for 15 min, cells were stained with Abs for 1 h at room temperature. After staining, cells were washed three times with incubation buffer before analysis. At least 100,000 viable cells were live gated on FACScan or FACScanCalibur (BD Biosciences) instrumentation. Flow cytometry analysis was performed using FlowJo software (Tree Star).

**T-cell immunization, activation, apoptosis assay and proliferation assay.** Mice were immunized by i.p. injection of 0.5 mg ovalbumin (OVA, from Sigma-Aldrich) and 50 μg poly (I:C) (Sigma Aldrich). For in vitro T-cell activation and apoptosis assay, purified CD8+ T cells were labeled with CFSE (Invitrogen-Molecular Probes) and incubated with OVA peptide257–264 (Mayo Clinic Core Facilities) at 0.2 μg/mL for 72 h. Apoptosis of CD8+ T cells was analyzed by staining using Annexin V (BD Biosciences) and TMRE (tetrathymothidamine ethyl ester, Invitrogen/Molecular Probes T-669). Proliferation was also measured by detection of BrdU incorporation and Ki67 staining. Immunized mice were injected i.p. with 0.8 mg/mL BrdU (BD Biosciences) on days 4 through 6 following immunization. On day 7 after immunization BrdU incorporation was determined by intra-nuclear staining with anti-BrdU (B9285, Sigma-Aldrich) and anti-Ki67 (556027, BD Biosciences).

**In vivo CTL assay.** For the in vivo CTL assay, OVA257–264 peptide-pulsed or control peptide-pulsed spleen cells (as target cells) from syngeneic mice were labeled with a high dose of CFSE (5 μM) or low dose of CFSE (0.5 μM), mixed at 1:1 (2.5 × 10^6 of each) before injection. Target cells were i.v. injected into immunized mice on day 4 after re-challenge with cognate antigen protein. The CTL activity was determined 4 h after target cell transfer. Specific lysis is calculated using the following formulas: ratio = (% CFSEhigh/% CFSElow), % specific lysis = [1 − (ratio primed/ratio unprimed)] × 100%

**Tumor studies.** Mice were inoculated i.v. with 5 × 10^5 B16-OVA tumor cells on day 25 after immunization. On day 21-post tumor injection mice were sacrificed and the lung tissue was perfused with PBS. The number of tumor foci on the lung tissue was counted.

**T-cell transfer experiments.** Purified CD8+ T cells (1 × 10^6) from Thy1.1+ OT-1 transgenic mice were i.v. injected into Thy1.2+ WT or B7-H1-deficient recipient mice followed with immunization with OVA plus poly I:C. On day 7 after immunization, transferred CD8+ T cells were identified by their expression of Thy1.1 and used for detection of intracellular expression of Bim, Bcl-2 and Bcl-xL. Equal numbers of Cd80+ (CD45.2+) and WT OT-1 (Thy1.1+, CD45.2+) CD8+ T cells (10^4 of each) were i.v. injected into CD45.1+ mice followed with immunization of OVA and poly I:C. The transferred OT-1 CD8+ T cells in the spleen were identified by flow cytometry.

**In vitro T-cell activation and culturing with fusion proteins.** Spleen cells were harvested from naïve mice and pre-activated with ConA (5 μg/mL, L7647, Sigma-Aldrich) for 48 h. Following activation, CD8+ T cells were purified (EasySep CD8+ T-cell negative selection kit, Stem Cell Technologies) and incubated with plate-bound anti-CD3 (BD Biosciences) and B7-H1 Fc fusion protein or control Fc protein (R&D Systems). Cultures were maintained for indicated time periods, and then cells were harvested for analysis.

**Western blotting.** Cells were lysed with NETN buffer (0.5% NP40, 150 mM NaCl, 50 mM Tris and 1 mM EDTA). Cell lysates were boiled and run on SDS-PAGE gels (BioRad), transferred to nitrocellulose membrane (Millipore), and blotted using standard procedures.

**Quantitative RT-PCR.** Total RNA was isolated from purified CD8+ T cells (RNaseq Kit, Qiagen), and reverse transcribed (iScript cDNA synthesis kit, BioRad). Samples were analyzed for Bim transcript levels using Bcl211 primers (Qiagen) and QuantiFast SYBR Green PCR Master Mix (Qiagen) on an iCycler (BioRad), GAPDH levels were used to normalize data by the comparative C_{T} method.

**Statistical analysis.** A two-sided, unpaired or paired Student’s t-test was used to assess statistical differences in experimental groups. A p value < 0.05 was considered statistically significant.

**Disclosure of Potential Conflicts of Interest**

No potential conflicts of interest were disclosed.

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**Supplemental Materials**

Supplemental materials may be found here:
www.landesbioscience.com/journals/oncoimmunology/article/20850

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