## Critical role of DNAX accessory molecule-1 (DNAM-1) in the development of acute graft-versus-host disease in mice

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Acute graft-versus-host disease (GVHD) is a life-threatening complication following bone marrow transplantation; however, no effective molecular-targeting therapy has been determined. Here, we show that mice that received allogeneic splenocytes deficient in DNAX accessory molecule-1 (DNAM-1) had significantly milder GVHD and lower mortality than those that received allogeneic WT splenocytes. Donor CD8<sup>+</sup> T cells deficient in DNAM-1 showed significantly less proliferation and infiltration of the liver and intestines of recipient mice and produced less IFN- $\gamma$  after coculture with allogeneic splenocytes than WT CD8<sup>+</sup> T cells. Mice prophylactically treated with an anti-DNAM-1 antibody showed milder GVHD and lower mortality than those treated with a control antibody. Moreover, treatment with a single administration of the antibody after the overt onset of GVHD ameliorated GVHD and prolonged survival. Finally, we show that the anti-DNAM-1 antibody therapy also ameliorated the overt GVHD in lethally irradiated mice after MHC-matched, minor antigen-mismatched bone marrow transplantation. These results indicate that DNAM-1 plays an important role in the development of GVHD and is an ideal molecular target for therapeutic approaches to GVHD.

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llogeneic bone marrow transplantation (BMT) offers ther-A apy for a variety of hematological malignancies and for both inherited and acquired nonmalignant hematological disorders (1, 2). However, the outcome of treatment depends largely on the development of graft-versus-host disease (GVHD), a major and mortal complication of allogeneic BMT (3). Alloreactive donor T lymphocytes that recognize alloantigens in the host are primed by host antigen-presenting cells (APCs), are activated, and mount cellular immune responses against the recipient tissues as "non-self" damaging host tissues, typically in the liver, gastrointestinal tract, and skin (1, 2, 4). However, the underlying molecular mechanism is incompletely understood. Although immunosuppressants are widely used for prophylaxis and treatment of GVHD, no effective molecular targeting therapy specifically targeting the pathogenesis of GVHD has been determined, contributing to poor prognosis of patients with higher-grade GVHD (5-7).

Although the alloantigen-specific signal mediated by the T-cell receptor is essential for priming of the pathogenic T cells in GVHD development, a costimulatory signal is required for full activation of the T cells, which leads to the development of exacerbated GVHD (1–3, 8). A strategy to block costimulatory signaling has been established: several blocking mAbs and fusion proteins targeting the interaction between costimulatory molecules expressed on T cells and ligands on APCs have been effective in mouse GVHD models (8–12). However, most studies have focused on prophylaxis, not on therapy after the onset of GVHD. Although several blocking approaches have been used

in clinical trials, the effects have been inadequate and even deleterious (5–7, 13–16).

The leukocyte adhesion-molecule DNAX accessory molecule-1 (DNAM-1, also known as CD226) is a member of the Ig superfamily and is constitutively expressed on most CD4<sup>+</sup> T cells, CD8<sup>+</sup> T cells, natural killer (NK) cells, monocytes and macrophages, and platelets (17). The poliovirus receptor CD155 and its family member nectin-2 (CD112, also called poliovirus receptor-related family 2) are ligands for DNAM-1 (18, 19). CD155 and CD112 are broadly distributed on hematopoietic, epithelial, and endothelial cells in different amounts in many tissues, as well as in many types of tumors, in human and mouse (20-29). Interactions between DNAM-1 on CD8<sup>+</sup> T cells and NK cells and its ligands CD112 and CD155 on target cells augment cell-mediated cytotoxicity (19, 24, 25, 30). Thus, DNAM-1 is involved in a variety of T-lymphocyte functions for the elicitation of appropriate adaptive immune responses, raising the possibility that it is associated with the severity of several diseases, including cancers, autoimmune diseases, and GVHD.

Here, we explored the role of DNAM-1 in GVHD in a mouse model and assessed the feasibility of previously unexplored prophylactic and therapeutic approaches using blocking antibodies against DNAM-1.

## Results

DNAM-1 Expression on Donor CD8<sup>+</sup> T Cells, but Not Recipient Cells, Is Involved in Development of Acute GVHD. To examine whether DNAM-1 is involved in the pathogenesis of GVHD, we transplanted splenocytes from WT or DNAM-1–deficient (KO) C57BL/6N (B6) (H-2<sup>b</sup>) mice into sublethally irradiated B6C3F1 (H-2<sup>b/k</sup>) mice. All of the mice that received WT splenocytes died by day 50 after transplantation. In contrast, the mice that received DNAM-1 KO splenocytes lived significantly longer (Fig. 1*A*). Histopathological examination showed that mononuclear cell infiltration and hepatocellular necrosis in the liver and villous atrophy in the small intestine were clearly milder in recipients of DNAM-1–deficient splenocytes (Fig. S14). Liver dysfunction, as determined by serum alanine aminotransferase (ALT) and

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The authors declare no conflict of interest.

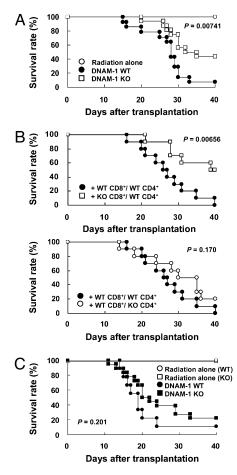
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**Fig. 1.** DNAM-1 expression on donor CD8<sup>+</sup> T cells is involved in exacerbation of acute GVHD. (A) After sublethal irradiation, B6C3F1 mice received splenocytes from DNAM-1 WT (n = 14) or KO B6 (n = 16) mice. B6C3F1 mice that received irradiation only are also shown (n = 5). Data are representative of two independent experiments. (B) After sublethal irradiation, B6C3F1 recipient mice received TCD-SP plus CD8<sup>+</sup> T cells from DNAM-1 WT or KO B6 mice and CD4<sup>+</sup> T cells from DNAM-1 WT or KO B6 mice and CD4<sup>+</sup> T cells from DNAM-1 WT or KO B6 mice (n = 10 in each group). Data are pooled from two independent experiments. (C) After sublethal irradiation, DNAM-1 WT (n = 9) or KO (n = 18) CBF1 mice received splenocytes from DNAM-1 WT B6 mice. DNAM-1 WT and KO CBF1 mice that received irradiation alone are also shown (n = 5 and 7, respectively). The experiments were independently performed four times and pooled data from all of the experiments are shown.

aspartate aminotransferase (AST) values, and the levels of serum IFN- $\gamma$  were significantly less elevated in the recipients of DNAM-1–deficient splenocytes (Fig. S1 *B* and *C*). These results indicate that DNAM-1 expressed on donor cells is involved in the development of GVHD.

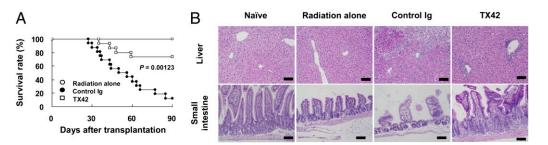
To next examine which cell type of donor cells expressing DNAM-1 plays a critical role in the development of GVHD, we prepared both CD4<sup>+</sup> and CD8<sup>+</sup> T cell-depleted splenocytes (TCD-SP) from WT B6 mice, and TCD-SP reconstituted with CD4<sup>+</sup> (derived from DNAM-1 WT or KO mice) and CD8<sup>+</sup> T cells (derived from DNAM-1 WT or KO mice). In contrast to B6C3F1 mice that received TCD-SP alone, recipient mice transplanted with TCD-SP reconstituted with any combinations of CD4<sup>+</sup> and CD8<sup>+</sup> T cells showed significantly higher levels of ALT and IFN- $\gamma$  (Fig. S1 *D* and *E*). Notably, serum levels of ALT and IFN- $\gamma$  were higher in mice that received TCD-SP reconstituted with DNAM-1 WT CD8<sup>+</sup> T cells than those that received DNAM-1 KO CD8<sup>+</sup> T cells-reconstituted TCD-SP, regardless of DNAM-1 expression on donor CD4<sup>+</sup> T cells (Fig.

S1 *D* and *E*). In accordance with these results, recipient mice transplanted with TCD-SP reconstituted with both DNAM-1 WT CD8<sup>+</sup> and WT CD4<sup>+</sup> T cells showed significantly higher mortality than those that received TCD-SP reconstituted with KO CD8<sup>+</sup> T cells and WT CD4<sup>+</sup> T cells (Fig. 1*B*, *Upper*), but there was no difference in mortality between recipient mice transplanted with TCD-SP reconstituted with DNAM-1 WT CD8<sup>+</sup> T cells and WT CD4<sup>+</sup> T cells and those transplanted with TCD-SP reconstituted with DNAM-1 WT CD8<sup>+</sup> T cells and WT CD4<sup>+</sup> T cells and those transplanted with TCD-SP reconstituted with DNAM-1 WT CD8<sup>+</sup> T cells (Fig. 1*B*, *Lower*). Taken together, these results indicate that DNAM-1 on donor CD8<sup>+</sup>, rather than CD4<sup>+</sup>, T cells plays a central role in the development of GVHD.

We further examined whether DNAM-1 expressed on recipient cells is also involved in the pathogenesis of GVHD. We established a GVHD model in which splenocytes from WT B6 mice were transplanted into sublethally irradiated WT or DNAM-1 KO CBF1 (H- $2^{b/d}$ ) mice. One half of the mice died by day 20 after transplantation in both recipient groups and there was no significant difference in survival rates between the two groups (Fig. 1*C*), suggesting that DNAM-1 expression on recipient cells is not involved in development of acute GVHD.

Anti-DNAM-1 mAb Suppresses Development of Acute GVHD. To examine whether prophylactic treatment with an anti-DNAM-1 neutralizing mAb, TX42 (Fig. S24), suppressed the development of acute GVHD, we injected WT mice with TX42 or control antibody every other day from day -1 until day 17 after transplantation. Although most mice injected with control antibody died within 3 mo after transplantation, the mice injected with anti-DNAM-1 mAb lived significantly longer (Fig. 24). Moreover, TX42 dramatically ameliorated injury to the liver and small intestine, compared with control antibody, on day 25 after splenocyte transfer (Fig. 2B). In accordance with the histopathological analysis, TX42 prevented the elevation of serum ALT, AST, and IFN- $\gamma$  (Fig. S3 A and B). These results are similar to those in the recipients of the DNAM-1 KO splenocytes, suggesting that blockade of DNAM-1 signaling by using a neutralizing mAb in vivo prophylactically suppressed the development of acute GVHD.

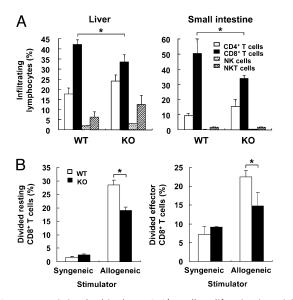
Anti-DNAM-1 mAb Suppresses Donor CD8<sup>+</sup> T-Cell Proliferation in Recipient Mice. To explore the role of DNAM-1 on donor cells in the pathogenesis of GVHD, we first used flow cytometry to examine the number of donor CD4<sup>+</sup> T cells (H-2K<sup>k-</sup>CD4<sup>+</sup>) and  $CD8^+$  T cells (H-2K<sup>k-</sup>CD8<sup>+</sup>) in the peripheral blood during the progression of acute GVHD (Fig. S4A). Although the numbers of donor CD4<sup>+</sup> T cells from DNAM-1 WT and KO mice in the peripheral blood of recipient mice were comparable (Fig. S4B, Left), CD8<sup>+</sup> T cells derived from DNAM-1 KO mice were present in significantly lower numbers than those from DNAM-1 WT mice on day 21 after transplantation (Fig. S4B, Right). Similar results were also observed in mice that received TX42 when compared with control antibody (Fig. S4C). Moreover, although the frequencies of total donor-derived hematopoietic cells in the liver and small intestine, the major target organs of alloreactive CD8<sup>+</sup> T cells in GVHD, was comparable between two groups that received DNAM-1 WT or KO splenocytes (Fig. S4D), the populations of donor  $CD8^+$  T cells infiltrating into these organs of recipients of DNAM-1 KO splenocytes were significantly smaller than those in recipients of WT splenocytes (Fig. 3A). These results indicate that blocking DNAM-1 signaling limits the proliferation of donor  $CD8^+$  T cells in vivo. Most donor  $CD8^+$  T cells, as well as  $CD4^+$  T cells from DNAM-1 WT and KO mice, differentiated into CD44<sup>high</sup>CD62L<sup>low</sup> effector T cells at day 21 after transplantation (Fig. S5), suggesting that there were fewer effector  $\overline{CD8}^+$  T cells in proportion to the total CD8<sup>+</sup> T cells in recipients of DNAM-1 KO vs. WT splenocytes or in mice treated with TX42. Taken together with the results



**Fig. 2.** Anti–DNAM-1 mAb suppressed the development of acute GVHD. (A) After sublethal irradiation, B6C3F1 mice received splenocytes from WT B6 mice. The recipient mice were i.p. injected with anti–DNAM-1 (TX42) (n = 14) or control antibodies (n = 16) every other day from day –1 until day 17 after transplantation. B6C3F1 mice that received irradiation only are also shown (n = 10). Data were pooled from three independent experiments. (*B*) The liver and small intestine of mice on day 25 after transplantation were stained with H&E and histologically analyzed. The organs of three mice in each group were analyzed and representative data of a mouse in each is shown. (Scale bars, 100  $\mu$ m.)

shown in Figs. 1 to 3, these results suggest that DNAM-1 on donor CD8<sup>+</sup> T cells plays an important role in cell proliferation and infiltration in the liver and small intestine, and is thus involved in the development of acute GVHD.

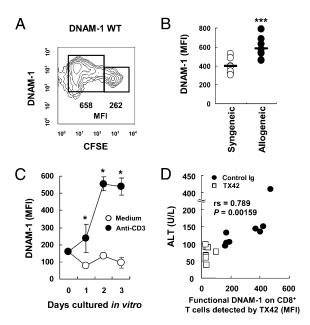
DNAM-1 Costimulation Promotes Proliferation of and IFN-γ Production by Alloreactive CD8<sup>+</sup> T Cells. DNAM-1 mediates a costimulatory signal in cytotoxic T cells and promotes cytotoxicity against target cells expressing DNAM-1 ligands (30, 31). To examine whether DNAM-1 is also involved in the costimulatory effect on proliferation of donor CD8<sup>+</sup> T cells after priming in vivo, we transplanted 5-(and 6)-carboxyfluorescein diacetate succinimidyl ester (CFSE)-labeled splenocytes T cells from B6 mice into sublethally irradiated B6C3F1 or B6 mice. In contrast to CD8<sup>+</sup> T cells transplanted into syngeneic mice, most DNAM-1 WT CD8<sup>+</sup> T cells transplanted into allogeneic mice had divided at 3 d after



**Fig. 3.** DNAM-1 is involved in donor CD8<sup>+</sup> T-cell proliferation in recipient mice after transplantation. (*A*) After sublethal irradiation, B6C3F1 mice received splenocytes from DNAM-1 WT or KO B6 mice. The infiltrating cells in the liver and small intestine in recipient mice (n = 3) on day 14 after transplantation were separated, and each donor-derived (H-2K<sup>k-</sup>) lymphocyte subset was determined by flow cytometry. Data are representative of two independent experiments. (*B*) Resting CD8<sup>+</sup> T cells purified from naive DNAM-1 WT or KO B6 mice (*Left*) and donor effector CD8<sup>+</sup> T cells purified from B6C3F1 mice that received WT or KO B6 splenocytes (*Right*) were labeled with CFSE, cocultured with mitomycin C-treated syngeneic (B6) or allogeneic (B6C3F1) splenocytes for 3 d, and analyzed by flow cytometry. Data are representative from three independent experiments with similar results. \*P < 0.05. Error bars show SD.

transplantation (Fig. S6A), suggesting that the cell division resulted from alloantigen recognition. However, fewer DNAM-1 KO CD8<sup>+</sup> T cells transplanted into allogeneic mice divided than did WT  $CD8^+$  T cells (Fig. S64), indicating that DNAM-1 mediates a costimulatory signal for proliferation in CD8<sup>+</sup> T cells in vivo. In vivo injection with TX42 mAb on day -1 and day 1 after transplantation suppressed cell division of donor CD8<sup>+</sup> T cells (Fig. S6B). To verify these results in vitro, we stimulated CFSE-labeled resting CD8+ T cells from DNAM-1 WT or KO mice with the anti-CD3 mAb plus an anti-DNAM-1 or control Ig. DNAM-1 signaling enhanced the proliferation of resting CD8<sup>+</sup> T cells only when CD8<sup>+</sup> T cells were simultaneously stimulated with anti-CD3 mAb (Fig. S6 C and D). We also performed the same proliferation assay using donor-derived effector CD8<sup>+</sup> T cells sorted from recipient B6C3F1 mice that had been infused with donor B6 splenocytes. As in the resting CD8<sup>+</sup> T cells, the proliferation of effector CD8<sup>+</sup> T cells was augmented by DNAM-1 costimulation (Fig. S6E). The costimulatory effect of DNAM-1 on proliferation was also observed in cocultures of resting or effector CD8<sup>+</sup> T cells with allogeneic stimulators, such as whole splenocytes (Fig. 3B) and either CD11c<sup>+</sup> or CD11c<sup>-</sup> cells purified from the spleen (Fig. S6F). Addition of a neutralizing anti-CD155 mAb (TX56) in the assay significantly suppressed the alloantigen-specific CD8<sup>+</sup> T-cell proliferation (Fig. S6G), suggesting that CD155 expression on the spelnocytes was involved in DNAM-1-mediated costimulation in CD8<sup>+</sup> T cells. Similar to the costimulatory function of DNAM-1 in the proliferation of CD8<sup>+</sup> T cells, DNAM-1 also mediated a costimulatory signal for IFN-y production by resting and effector  $CD8^+$  T cells (Fig. S6 H and I). These results demonstrate that DNAM-1 costimulation of alloreactive CD8<sup>+</sup> T cells promotes the activation and proliferation of and IFN-y production by alloreactive  $CD8^+$  T cells.

Anti-DNAM-1 mAb Suppresses the Up-Regulation of DNAM-1 Expression on Donor CD4<sup>+</sup> and CD8<sup>+</sup> T Cells in Recipient Mice. To further investigate the role of DNAM-1 in the pathogenesis of GVHD, we analyzed its expression on donor T cells in recipient mice. B6C3F1 recipient mice received CFSE-labeled splenocytes from WT B6 mice, and the expression of DNAM-1 on donor CD8<sup>+</sup> T cells was analyzed by flow cytometry 3 d after transplantation. The mean fluorescence intensity of DNAM-1 on proliferating donor CD8<sup>+</sup> T cells was significantly higher than that on nondividing CD8<sup>+</sup> T cells in recipient mice (Fig. 4A). To examine whether the up-regulation of DNAM-1 expression on CD8+ T cells was induced by alloantigen recognition, B6 or B6C3F1 recipient mice were transplanted with CFSE-labeled splenocytes from WT B6 mice. Three days after transplantation, DNAM-1 expression on donor CD8<sup>+</sup> T cells in the spleen of allogeneic recipient mice was significantly higher than that in syngeneic recipients (Fig. 4B). In vitro stimulation of CD8<sup>+</sup> T cells with an anti-CD3 mAb for 3 d also up-regulated DNAM-1

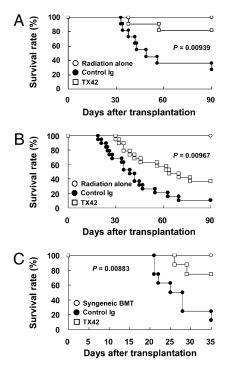


**Fig. 4.** DNAM-1 expression on donor T cells was up-regulated in recipient mice. (*A* and *B*) After sublethal irradiation, B6C3F1 (allogeneic) or B6 (syngeneic) recipient mice received CFSE-labeled splenocytes from WT B6 mice, and CFSE<sup>+</sup> donor CD8<sup>+</sup> T cells in the spleen were analyzed by flow cytometry for the expression and mean fluorescence intensity (MFI) of DNAM-1 on day 3 after transplantation. (C) CD8<sup>+</sup> T cells from B6 mice were stimulated with plate-coated anti-CD3 and analyzed for the expression of DNAM-1 on CD8<sup>+</sup> T cells by flow cytometry. (*D*) The correlation between DNAM-1 expression on donor CD8<sup>+</sup> T cells on day 21 and ALT values on day 28 in recipients injected with control lg or TX42 was statistically evaluated. Data are representative of three independent experiments with similar results, respectively. \*\*\*P < 0.005; \*P < 0.05. Error bars show SD.

expression (Fig. 4*C*). These results suggest that DNAM-1 expression on alloreactive T cells was up-regulated during proliferation in recipient mice. Of note, DNAM-1 on these donor T cells in recipient mice treated with TX42 was significantly less detected by TX42 mAb than in mice treated with control antibody on days 7 to 21 (Fig. S2*C*), probably because of masking of DNAM-1 by injected TX42 in vivo (Fig. S2*B*). Because TX42 mAb is a neutralizing antibody, the masked DNAM-1 on CD8<sup>+</sup> cells is not functional. Taken together, these results suggest that development of acute GVHD is associated with up-regulated expression of DNAM-1 on donor T cells and anti–DNAM-1 mAb TX42 suppresses DNAM-1 function in vivo.

**DNAM-1 Ligand Expression in Target Organs in Acute GVHD.** We next examined the expression of DNAM-1 ligands *Cd112* and *Cd155* in target organs in acute GVHD. *Cd112* was expressed predominantly in the liver, large and small intestines, and kidney (Fig. S7). Although *Cd155* was expressed at the highest levels in the heart and kidney, the liver and large and small intestines also expressed a significant amount of *Cd155* (Fig. S7). These results suggest that up-regulation of DNAM-1 on alloreactive CD8<sup>+</sup> T cells and constitutively high expression of the ligands, particularly CD112, in the liver and intestines are important in the pathogenesis of GVHD. Indeed, the levels of functional DNAM-1 expression detected by TX42 mAb on donor CD8<sup>+</sup> T cells were significantly correlated with ALT values (Fig. 4D), suggesting that DNAM-1 expression on CD8<sup>+</sup> T cells plays a critical role in the exacerbation of GVHD.

Therapeutic Treatment with a Single Administration of Anti–DNAM-1 mAb Ameliorates GVHD. We next investigated whether treatment with DNAM-1 had a therapeutic effect on overt GVHD. Anti– DNAM-1 mAb TX42 or control Ig was injected i.p. every week into recipient mice from day 14, when GVHD had become overt, to day 77. The mice treated with TX42 mAb showed significantly lower mortality, a dramatic improvement of liver dysfunction (as determined by serum ALT and AST values), and decreased IFN- $\gamma$  in the sera after treatment (Fig. 5A and Fig. S8 A and B). Injection of TX42 mAb significantly decreased functional DNAM-1 on donor-derived CD8<sup>+</sup> T cells, as detected by staining with TX42 mAb (Fig. S8C). The amount of functional DNAM-1 on CD8<sup>+</sup> T cells was associated with the progression of GVHD, because there was a positive and significant correlation between the amount of functional DNAM-1 on donor-derived CD8<sup>+</sup> T cells and ALT values (Fig. S8D). Furthermore, the numbers of donor-derived CD8<sup>+</sup> T cells in the recipients were decreased after treatment with TX42 (Fig. S8E). Therefore, the blockade of DNAM-1 signaling by administration of a blocking mAb against DNAM-1 in a therapeutic setting could inhibit the expansion of alloreactive effector CD8<sup>+</sup> T cells even after the onset of GVHD. To assess the feasibility of the therapeutic approach with anti-DNAM-1 mAb, we examined whether a single, rather than multiple, administration of anti-DNAM-1 mAb was also effective for the therapy of GVHD. The mice treated with TX42 mAb only on day 14 also showed significantly lower mortality and longer survival (Fig. 5B).



**Fig. 5.** Anti–DNAM-1 mAb ameliorated overt acute GVHD. (*A* and *B*) B6C3F1 mice received splenocytes from WT B6 mice after sublethal irradiation. The recipient mice were injected i.p. with 300  $\mu$ g anti–DNAM-1 (TX42) (n = 11) or control antibodies (n = 11) every week from day 14 until day 77 after transplantation (*A*), or 1.0 mg TX42 (n = 19) or control antibodies (n = 10) or control antibodies (n = 8) once on day 14. Recipient B6 mice that received transplantation with bone marrow cells and T cells after lethal irradiation (Syngeneic BMT) are also shown (n = 6). Data are representative of two independent experiments with similar results in *A*. Data are pooled from two independent experiments in *B* and C.

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Treatment with Anti–DNAM-1 mAb Ameliorates GVHD in Lethally Irradiated Mice After MHC-Matched, Minor Antigen-Mismatched BMT. Finally, we investigated the involvement of DNAM-1 in the development of GVHD in a more clinical-relevant GVHD model after minor-mismatched BMT. Bone marrow cells and T cells from B6 mice (H-2<sup>b</sup>) were transplanted into lethally irradiated minor-mismatched C3.SW-H2<sup>b</sup>-Sn/J (C3) mice (H-2<sup>b</sup>). Most recipient mice that were treated with control Ig on day 14 died by day 35 after BMT. In contrast, treatment of the recipient mice with a single administration of TX42 mAb on day 14 significantly prolonged the survival (Fig. 5*C*). These results indicate that the administration of a neutralizing mAb against DNAM-1 is highly potential for therapy, as well as prophylaxis for GVHD in a clinical-relevant setting.

## Discussion

Given that alloreactive CD8<sup>+</sup> T cells primed by host APCs presenting alloantigens directly mediate host-tissue injuries in GVHD (2, 3), it is important that this study has clarified the role of DNAM-1-mediated costimulation of pathogenic CD8<sup>+</sup> T cells in the development of GVHD. Previous works have revealed that DNAM-1 plays several important roles in modulating cellular immunity, including (i) the enhancement of cytotoxic T lymphocyte (CTL) (including alloantigen-specific CTL)- and NK cell-mediated cytotoxicity against target cells expressing CD155 or CD112 and cytokine secretion, such as IFN- $\gamma$ , and (*ii*) the proliferation and differentiation of naïve CD4<sup>+</sup> T cells toward Th1 cells in cooperation with leukocyte function-associated antigen-1 (LFA-1) (17, 19, 30, 32). A previous report demonstrated that DNAM-1 promoted CD8<sup>+</sup> T-cell proliferation in response to the presentation of superantigens or OVA peptide antigens by nonprofessional APCs but not dendritic cells (31). In contrast, we demonstrated that nonprofessional (CD11c<sup>-</sup>) as well as professional (CD11c<sup>+</sup>) APCs in the spleen were involved in the costimulatory effect of DNAM-1 in  $CD8^+$  T cells. This discrepancy may be explained by the molecular and functional differences between the alloantigens and the OVA peptide antigen or superantigens. Although emerging evidence supports the involvement of DNAM-1 in pathogenic T cell-mediated immune diseases, as yet the mechanisms responsible have not been identified. Here we have demonstrated that GVHD exacerbation is caused by augmentation of the functions of pathogenic  $CD8^+$  T cells by DNAM-1 costimulation, in which the signaling promotes the proliferation of alloreactive T cells immediately after priming, as well as the subsequent clonal expansion of the T cells in the effector phase in vivo and in vitro. In addition, the enhanced production of IFN- $\gamma$  by CD8<sup>+</sup> T cells dependent on DNAM-1 costimulation potentially contributes to the exacerbation, a result of the immunomodulating effect of IFN-y on Th1-biased promotion of cellular immunity (33). Thus, our results provide definitive evidence of the critical involvement of DNAM-1 in the pathogenesis of GVHD and of the immunological mechanisms of the exacerbation of the disease.

Most conventional molecular targeting strategies using neutralizing mAbs for prophylaxis for GVHD focus on blocking the interaction between receptors expressed on pathogenic T cells and the ligands on APCs, especially dendritic cells, that leads to priming of T cells in the draining lymph nodes in patients with GVHD (e.g., CD28 and CTLA-4-B7, CD40L-CD40, and LFA-1-ICAM) (5, 9–12). The expression of the ligands for these costimulatory receptors is restricted to professional APCs or hematopoietic cells. In contrast, we have shown that two DNAM-1 ligands (CD112 and CD155) are widely distributed not only on hematopoietic cells, but also on nonhematopoietic cells in many tissues in mice, including the liver and intestines, the major target organs of GVHD (20–22, 26–29, 34, 35). Previous reports demonstrated that the liver expresses *Cd155* much more than the other organs in human (20, 35). Remarkably, we demonstrated that DNAM-1 expression was up-regulated on  $CD8^+$  T cells after priming and was maintained at high levels on effector T cells in recipient mice. Together, these results suggest that DNAM-1 is involved in effector phase as well as priming phase of alloreactive  $CD8^+$  T cells that directly attack the target organs expressing DNAM-1 ligands in the host. This finding may be one of the reasons why the treatment with anti–DNAM-1 antibody is effective in the therapy as well as prophylaxis for GVHD.

Patients that received BMT are at the high risk of infectious diseases and relapse of hematological and nonhematological tumors under a long-term immunosuppressive state. Although the role of DNAM-1 in immune response against infectious diseases remains to be elucidated, we previously demonstrated that DNAM-1 plays an important role in tumor immune surveillance (30). Administration of a long-term overabundant dose of a neutralizing mAb against DNAM-1 in a clinical application might be deleterious due to the impairment of graft-versusleukemia/tumor effect in patients; however, attenuation of GVHD with concomitant potent graft-versus-leukemia/tumor effect might be achieved by the regulation of dose and timing of administration of a neutralizing mAb against DNAM-1. In fact, we demonstrated that only a single dose of anti-DNAM-1 mAb dramatically ameliorated GVHD. These results encourage us to test clinical trials of this antibody therapy for GVHD in patients.

The feasibility of conventional strategies for blocking costimulatory molecules in most previous work was examined in prophylactic but not therapeutic approaches in mouse GVHD models (9-11, 36, 37). In sharp contrast, our administration of a neutralizing mAb against DNAM-1 after the clear onset of GVHD ameliorated disease by suppressing alloreactive effector CD8<sup>+</sup> T-cell proliferation, IFN- $\gamma$  production and, probably, cytotoxicity against recipient tissue cells. This report is unique in using a neutralizing mAb against DNAM-1 for the blockade, and it validates the efficacy of this strategy for the amelioration of GVHD in both MHC-mismatched transplantation and MHCmatched, minor antigen-mismatched BMT in mice. Our GVHD model used MHC-mismatched mouse as a donor, which might be clinically irrelevant. However, because BMT from one locus of HLA-mismatched donor is frequently performed in patients at present, an important point in the present study is that DNAM-1 blocking is able to overcome even a high-grade GVHD induced by MHC-mismatched transplantation. Nonetheless, further works aimed at the association of DNAM-1 and its ligands with GVHD pathogenesis induced by transplantation with mismatched minor histocompatibility antigens will be also required for the clinical application of DNAM-1-targeting therapy for GVHD.

## **Materials and Methods**

Materials and methods for mice, antibodies, ELISA, assessment of GVHD, infiltrating donor lymphocytes, proliferation assays, and quantitative RT-PCR used here are described in *SI Materials and Methods*.

**GVHD Model.** Recipient B6C3F1 mice were sublethally irradiated with 500 cGy by X-ray (Hitachi Medical Corporation). Fifty-million splenocytes from DNAM-1 WT or KO B6 mice were i.v. infused into each recipient B6C3F1 mouse. For prophylaxis, 100  $\mu$ g TX42 was injected i.p. into the recipient mice 1 d before transplantation (day –1) and then every other day for nine additional doses (total 10 times, 1.0 mg per mouse). For therapy, 300  $\mu$ g TX42 was injected i.p. therapy, 300  $\mu$ g TX42 was injected 1 beginning on day 14 and then every week for nine additional doses (total 10 times, 3.0 mg per mouse), or 1.0 mg TX42 was injected i.p. once on day 14.

In some experiments, B6C3F1 recipient mice were transplanted with 3.8  $\times$  10<sup>7</sup> TCD-SP alone or the same number of TCD-SP plus 7.0  $\times$  10<sup>6</sup> CD4<sup>+</sup> T cells (derived from DNAM-1 WT or KO mice) and 5.0  $\times$  10<sup>6</sup> CD8<sup>+</sup> T cells (derived from DNAM-1 WT or KO mice) after sublethal irradiation with 500 cGy by X-ray. TCD-SP were purified from WT B6 mice by negative selection with biotinylated anti-CD4 and CD8 mAbs, followed with Dynabeads MyONE streptavidin (Invitrogen). TCD-SP contained T cells at less than 3%, as determined by flow cytometry. CD4<sup>+</sup> and CD8<sup>+</sup> T cells were purified from the spleens of DNAM-1 WT or KO B6 mice by negative selection with

biotinylated anti-B220, CD11b, CD11c, Gr-1, and DX5 (CD49b) mAbs and either biotinylated CD8 or CD4 mAbs, followed with Dynabeads MyONE streptavidin. Purities of CD4<sup>+</sup> and CD8<sup>+</sup> T cells were more than 85%, as determined by flow cytometry.

To examine whether DNAM-1 on recipient cells is involved in development of GVHD, we generated DNAM-1 WT or KO CBF1 mice (H-2<sup>b/d</sup>) by crossing  $Cd226^{+/-}$  BALB/c mice with  $Cd226^{+/-}$  B6 mice. These mice received 1  $\times$  10<sup>7</sup> splenocytes from WT B6 mice after sublethally irradiation (600 cGy).

For a GVHD model after minor-mismatched BMT, recipient C3 mice were lethally irradiated with 900 cGy 1 d before BMT (day -1), and then  $5 \times 10^{6}$  bone marrow cells and  $4 \times 10^{6}$  spleen T cells from B6 mice were i.v. infused into each recipient mouse on day 0.

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**Statistical Analyses.** To analyze survival, we used Kaplan-Meier estimation with the statistical analysis system-type log-rank test. The correlation between DNAM-1 expression levels and ALT values was evaluated with Spearman's rank-order correlation coefficient. All other statistical analyses were performed with the two-tailed Mann-Whitney *U* test. *P* < 0.05 was considered statistically significant.

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