

Ikaros deficiency in host hematopoietic cells separates GVL from GVHD after experimental allogeneic hematopoietic cell transplantation

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Abbreviations: ⁵¹Cr, Chromium-51; ALL, acute lymphoblastic leukemia; Allo-HCT, allogeneic hematopoietic stem cell transplantation; APC, allophycocyanin; APCs, antigen-presenting cells; BC, blast crisis; BLI, bioluminescence imaging; BM, bone marrow; BMDCs, bone marrow derived dendritic cells; BMT, bone marrow transplantation; CML, chronic myeloid leukemia; CRT, calreticulin; CTL, cytotoxic T cell; DCs, dendritic cells; FACS, Fluorescence-activated cell sorting; FBS, fetal bovine serum; FITC, fluorescein isothiocyanate; GVHD, graft-versus-host-disease; GVL, graft-versus-leukemia; HCT, hematopoietic stem cell transplantation; ICAM-1, intracellular adhesion molecule 1; Ik DN, Ikaros dominant negative; Ik, Ikaros; luc+, luciferase+; mAbs, monoclonal antibodies; MACS, magnetic-activated cell sorting; MBL-2, moloney-murine sarcoma virus-induced MBL-2 lymphoma cells; mCRT, murine calreticulin; MHC, major histocompatibility complex; MiHAs, multiple minor histocompatibility antigens; MLR, mixed lymphocyte reaction; PBS, phosphate buffered saline; PE, phycoerythrin; SIRP- α , signal regulatory protein α ; TCD-BM, T cell depleted bone marrow; Tregs, regulatory T cells; TSA, tumor specific antigen; UCUCA, University Committee on Use and Care of Animals; WT, wild-type.

The graft-versus-leukemia (GVL) effect following allogeneic hematopoietic stem cell transplantation (allo-HCT) is critical for its curative potential. However, GVL is tightly linked to graft-versus-host disease (GVHD). Among hematological malignancies, acute lymphoblastic leukemia (ALL) is the most resistant to GVL, although the reasons for this remain poorly understood. Clinical studies have identified alterations in Ikaros (Ik) transcription factor as the major marker associated with poor outcomes in ALL. We have shown that the absence of Ik in professional host-derived hematopoietic antigen-presenting cells (APCs) exacerbates GVHD. However, whether Ik expression plays a role in resistance to GVL is not known. In this study we used multiple clinically relevant murine models of allo-HCT to explore whether Ik expression in hematopoietic APCs and/or leukemic cells is critical for increasing resistance to GVL and thus inducing relapse. We found that Ik deficiency in host APCs failed to enhance GVL despite increased GVHD severity. Mechanistic studies with bone marrow (BM) chimeras and tetramer analyses demonstrated reduced tumor-specific immunodominant (gag+) antigen responses in the [B6Ik^{-/-}→B6] group. Loss of GVL was observed when both the leukemia cells and the host APCs were deficient in Ik. We found that calreticulin (CRT) expression in host antigen-presenting dendritic cells (DCs) of Ik^{-/-} animals was significantly lower than in wild-type animals. Rescuing CRT expression in Ik^{-/-} DCs improved leukemic-specific cytotoxic T cell function. Together, our data demonstrate that the absence of Ikaros in host hematopoietic cells promotes resistance to GVL despite increasing GVHD and thus provides a potential mechanism for the poor outcome of Ik^{-/-} ALL patients.

Introduction

Allogeneic hematopoietic stem cell transplantation (allo-HCT) is the only curative therapy for many hematologic

malignancies. The critical factor for success of allo-HCT depends upon the degree of the graft-versus-leukemia (GVL) effect. Primary disease relapse is one of the major complications and a leading cause of death following allo-HCT.^{1,2} Although GVL is

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important for curative therapy, GVL is also tightly linked with graft-versus-host disease (GVHD), which is another major complication of allo-HCT.³ Thus, attempts to increase GVL are associated with exacerbating GVHD and uncoupling the beneficial benefit of GVL from the toxic effects of GVHD is imperative to improve curative therapy. Among hematologic malignancies acute lymphoblastic leukemia (ALL) is most resistant to GVL,⁴ the reasons for which remain poorly understood. We have previously demonstrated that professional host-derived hematopoietic antigen-presenting cells (APCs) are necessary for the induction of robust GVL.^{5,6} However, the cell autonomous molecular mechanisms of APCs that are critical for mediating GVL are poorly understood.

We have recently shown that *Ikaros* (*Ik*) deficiency in host hematopoietic APCs exacerbates GVHD.⁷ *Ik* is a transcription factor known to be important for the development of lymphoid cells^{8,9} and for the leukemogenesis of certain hematologic malignancies, such as ALL.¹⁰ Clinical data have demonstrated that the prognosis of ALL patients with *Ik* deficiency or mutation is extremely poor.^{11,12} Our recent study revealed that *Ik* deficiency in host hematopoietic APCs exacerbates acute GVHD in a Notch-dependent manner.⁷ Therefore, the *Ik*-Notch axis in host APCs is an important pathway for acute GVHD. Paradoxically, despite increasing GVHD in experimental allogeneic hematopoietic stem cell transplantation (allo-HCT), clinical data suggest that *Ik* deficiency or mutation in ALL patients is associated with a high risk for relapse, i.e., decreased GVL. Whether *Ik* plays an important role in mediating this GVL resistance by its expression in the host hematopoietic cells is not known. Here, we investigated the impact of *Ik* in host hematopoietic APCs and leukemic cells on GVL. We show that deficiency of *Ik* in host hematopoietic APCs did not increase GVL despite increased GVHD severity. Mechanistic studies demonstrated that the loss of GVL was more pronounced when both the leukemia cells and the host hematopoietic-derived APCs were deficient in *Ik*. Absence of *Ik* in host-derived APCs also decreased T-cell responses to tumor-specific antigen (TSA). Calreticulin (CRT) is expressed on the surface of APCs¹³ and is associated with phagocytic function¹⁴ and antigen presentation via MHC class I molecules.¹⁵ Therefore, CRT expression on APCs plays an important role in regulating immune responses. We found that CRT expression in dendritic cells (DCs) of *Ik*^{-/-} animals is significantly lower than that in WT mice and that forced expression of CRT in *Ik*^{-/-} DCs enhanced tumor-specific cytotoxic T cell (CTL) function. Together, our data demonstrate that the absence of *Ikaros* in host hematopoietic cells increased GVHD but decreased GVL and thus provide a potential mechanism for the increased relapse in *Ik*^{-/-} ALL after clinical allo-HCT.

Results

Ikaros deficiency in host hematopoietic cells does not increase GVL despite increasing GVHD

We recently reported that *Ik* deficiency in host hematopoietic APCs exacerbates acute GVHD in a Notch-dependent manner

after experimental allo-HCT.⁷ In light of clinical observations that deficiency or mutation of *Ik* in ALL increases relapse,^{11,12} in the present study we explored whether *Ik* deficiency in host hematopoietic APCs also affects GVL responses. We first generated BM chimeras to evaluate the impact of *Ik* deficiency only in host hematopoietic APCs, without the confounding effects of *Ik* deficiency in either leukemia cells or in non-hematopoietic cell-derived APCs. Wild-type (WT) B6 Ly5.2 animals were lethally irradiated with 11 Gy and infused with 5×10^6 bone marrow (BM) cells and 5×10^6 splenocytes from syngeneic B6Ly5.1 WT or B6 *Ik*^{-/-} donors. The [B6→B6Ly5.2] or [*Ik*^{-/-}→B6Ly5.2] animals were then used as allo-HCT recipients 4 months after primary HCT. To test GVL responses, the chimeras received 9 Gy irradiation and were injected intravenously with 0.5×10^6 CD8⁺ T cells together with 5×10^6 BM cells from either syngeneic B6 or major histocompatibility (MHC) antigen-matched but multiple minor histocompatibility antigens (MiHA)-mismatched allogeneic C3H.SW donors. The model that we used for the experiments is a well-established donor CD8⁺ T-cell mediated murine GVHD model akin to most commonly performed clinical allogeneic bone marrow transplants (BMTs).^{5,16} Furthermore, syngeneic moloney-murine sarcoma virus-induced lymphoma (MBL-2) cells were injected concurrently with HCT. The tumors that we used express MHC class I but not class II and thus are susceptible only to CD8-mediated antitumor cytotoxicity. Therefore, in these experiments we analyzed only CD8⁺ T-cell responses that cause both GVHD and GVL in these models. Consistent with previous observations, *Ik* deficiency in host APCs exacerbated GVHD (Fig. 1A and as shown in our previous report⁷). However, despite increasing GVHD, there was no statistical difference in relapse mortality of the WT [B6→B6Ly5.2] and [*Ik*^{-/-}→B6Ly5.2] animals that received allogeneic T cells with MBL-2 tumor cells (Fig. 1B). The lack of increase in GVL was also observed at a lower dose (5×10^3 MBL-2 cells/mouse, Fig. 1C). To further confirm the effect on GVL, MBL-2 cells were transduced with lentivirus that contained a GFP and luciferase (*luc*⁺) construct to facilitate *in vivo* monitoring of the tumor burden by bioluminescence imaging (BLI) after allo-HCT. As shown in Fig. 1D, although the syngeneic WT [B6→B6Ly5.2] animals showed somewhat less tumor signal because of spinal cord infiltration, we confirmed that both allogeneic WT [B6→B6Ly5.2] and [*Ik*^{-/-}→B6Ly5.2] animals showed similar tumor growth kinetics and succumbed to the tumor burden even though the [*Ik*^{-/-}→B6Ly5.2] animals showed a greater severity of GVHD. Some syngeneic WT [B6→B6Ly5.2] animals showed hind-limb paralysis. To further rule out tumor and model artifacts, we used a different tumor model system and found a similar relapse mortality in other GVL with EL-4 tumor cells (1×10^4 cells/mouse; Fig. 1E).

Ikaros deficiency increases overall donor T-cell responses but not those directed against tumor-specific antigen

To determine the mechanisms underlying the comparable GVL effect despite significant differences in GVHD, we tested the expansion of total donor T cells and only T cells specific for tumor specific antigen (TSA). We used tetramer strategies of

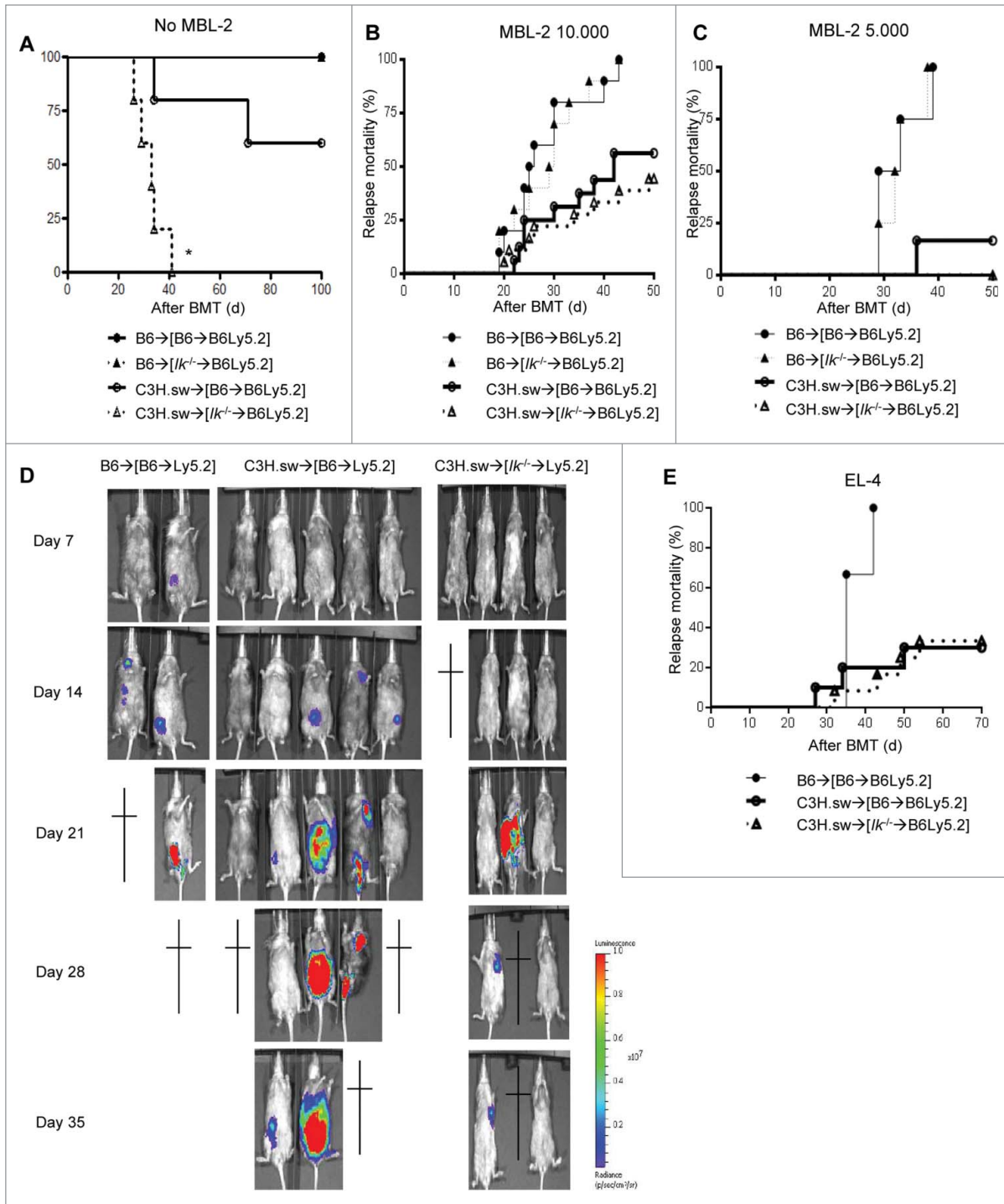


Figure 1. Ikaros deficiency in host APCs does not increase GVL responses regardless of enhanced GVHD in experimental HCT. WT B6 Ly5.2 animals were lethally irradiated with 11 Gy and infused with 5×10^6 bone marrow (BM) cells and 5×10^6 splenocytes from syngeneic Ly5.1 WT B6 or $Ik^{-/-}$ B6 donors. Four months later these [B6→B6Ly5.2] or [Ik^{-/-}→B6Ly5.2] chimeras received 9 Gy irradiation and 1×10^6 CD90⁺ T cells together with 5×10^6 BM cells from either syngeneic B6 or allogeneic MHC-matched or multiple miHA-mismatched C3H.sw donors concurrently with syngeneic MBL-2 tumor at the same time as allo-HCT. (A) Overall survival data of GVHD study. (●) B6→[B6→B6Ly5.2], (▲) B6→[Ik^{-/-}→B6Ly5.2], (◐) C3H.sw→[B6→B6Ly5.2], (◑) C3H.sw→[Ik^{-/-}→B6Ly5.2]. Data shown are one representative dataset (n=3–5/each group) of 5 independent experiments. (B, C) Tumor mortality data for MBL-2 at 10,000 cells/mouse (n=10–18/group) (B) and 5,000 cells/mouse (n=4–10/group). (C) (●) B6→[B6→B6Ly5.2], (▲) B6→[Ik^{-/-}→B6Ly5.2], (◐) C3H.sw→[B6→B6Ly5.2], (◑) C3H.sw→[Ik^{-/-}→B6Ly5.2]. Data are combined from 2 or 3 independent experiments. (D) Tumor growth was monitored using bioluminescence imaging (BLI) after allo-HCT (n=2–5). Representative data from 3 independent experiments are shown. (E) Tumor mortality data of the same model using a different syngeneic tumor, EL-4 (10,000 cells/mouse, n=3–12/group). (●) B6→[B6→B6Ly5.2], (◐) C3H.sw→[B6→B6Ly5.2], (▲) C3H.sw→[Ik^{-/-}→B6Ly5.2]. Data are combined from 2 independent experiments.

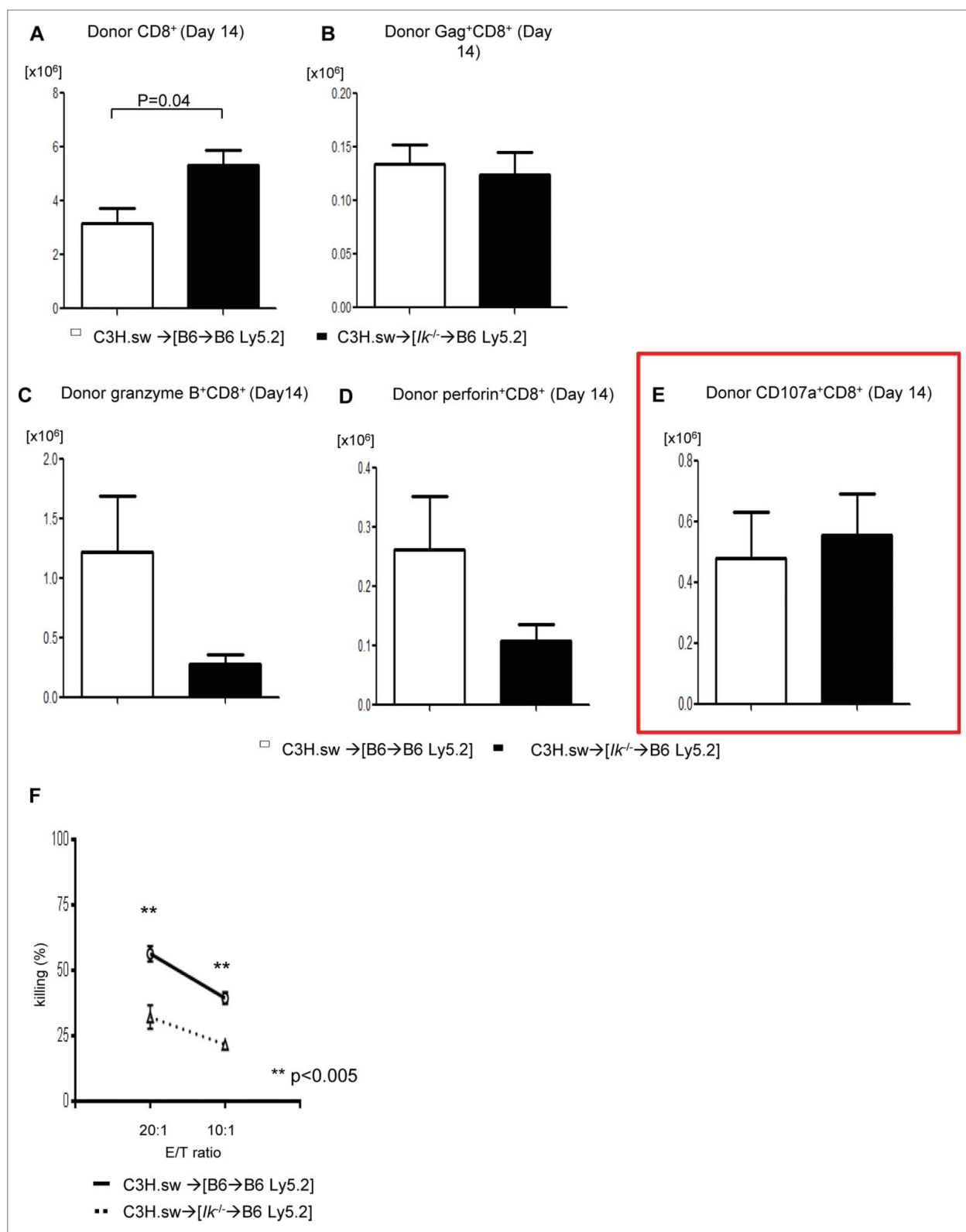


Figure 2. GVL response of host antigen-presenting cells with Ikaros deficiency is equivalent to that of WT cells. Donor T cells were isolated from spleen (n = 3–4) at day 14 following allo-HCT and analyzed for (A) donor CD8⁺ T-cell expansion (n = 3–4), (B) donor-derived MBL-2 specific gag⁺CD8⁺ T-cell expansion (n = 3–4), and expression of (C) granzyme B, (D) perforin, and (E) CD107a on donor CD8⁺ T cells (n = 3–4). (F) ⁵¹Cr-release assay using donor CD8⁺ T cells at day 14 after allo-HCT against MBL-2 tumors. One representative dataset from 3 independent experiments is shown. Data are given as mean + standard deviation.

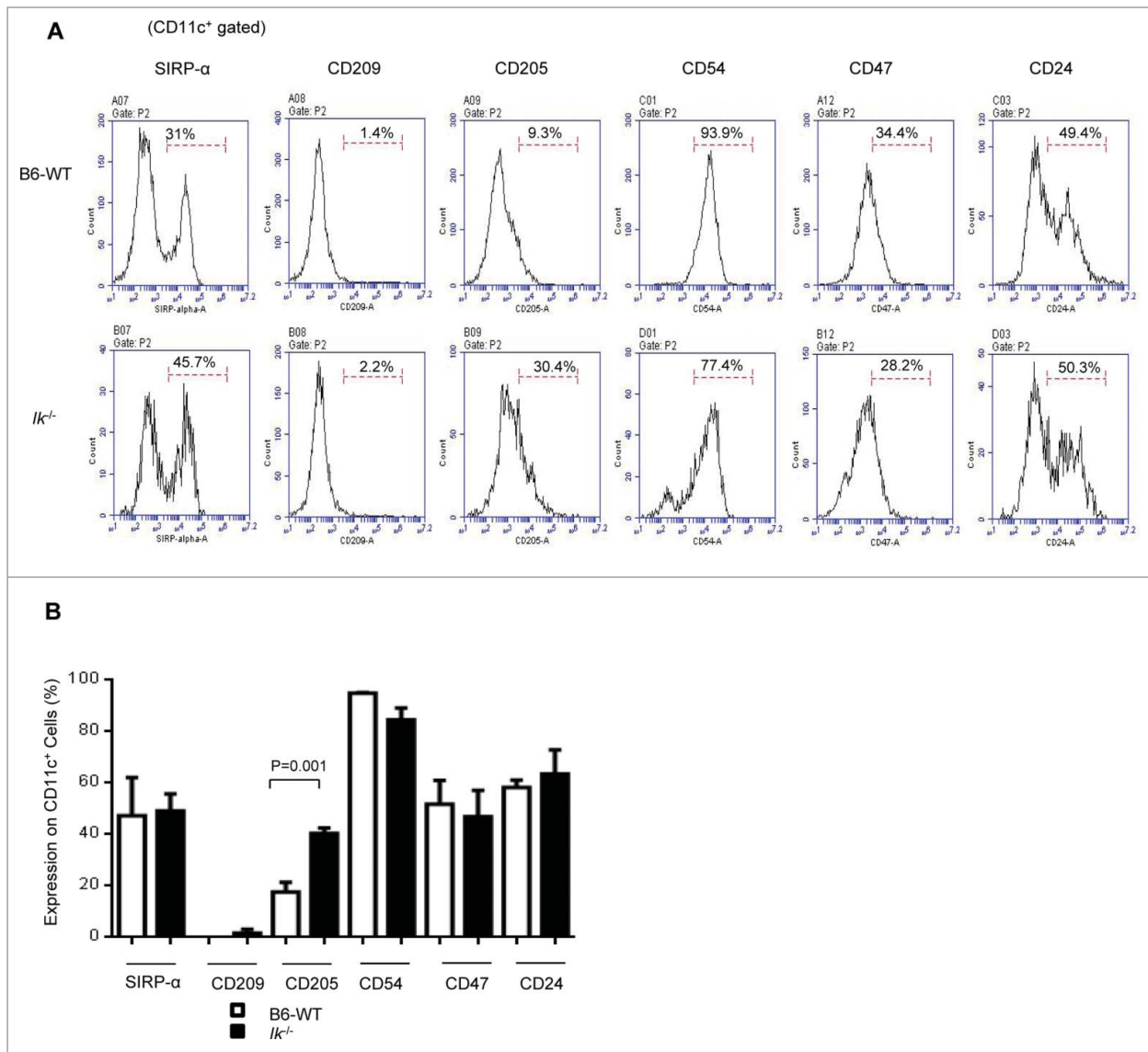


Figure 3. Expression of antigen-presenting molecules in splenic dendritic cells. The expression of antigen-presenting molecules (SIRP- α , CD209, CD205, CD54, CD27, and CD24) on CD11c⁺ cells in the spleen was analyzed for naive *Ik*^{-/-} and WT B6 animals that were not transplanted (n = 5–7/group). (A) Representative histogram. (B) Frequency of these antigen-presenting molecules on CD11c⁺ DCs.

donor CD8⁺ T cells to detect MBL-2 TSA *gag*-specific T-cell expansion.⁶ Using a similar model, we analyzed the expansion of *gag*-specific CD8⁺ donor T cells on day 21 following allo-HCT. We found that *Ik* deficiency in APCs caused greater allogeneic T-cell expansion (Fig. 2A) but similar TSA responses (Fig. 2B). To further assess the functionality of the TSA-specific donor T cells, we examined the cytotoxicity of donor T cells against MBL-2 tumor. We found similar expression of granzyme B, perforin, and CD107a (Fig. 2C–E) in donor T cells 14 d after allo-BMT in both groups. However, when specifically tested for their ability to kill tumor cells using the ⁵¹Cr release assay we found a reduced cytotoxicity of splenic donor T cells against MBL-2 in both groups on day 21 (Fig. 2F). These data demonstrated that *Ik* deficiency in host hematopoietic APCs enhances overall donor

T-cell responses but does not concomitantly increase TSA-specific responses.

Expression of antigen-presenting molecules on CD11c⁺ DCs in spleen

To evaluate the mechanism underlying the lack of increasing GVL responses despite the exacerbated GVHD between *Ik* and WT chimeras, we next determined the effect of *Ik* deficiency on the expression of antigen-presenting molecules on DCs. We found that only CD205 expression was significantly higher in *Ik*-deficient animals than in WT; the expression of other antigen-presenting molecules, such as signal regulatory protein (SIRP)- α , CD209, CD54/intercellular adhesion molecule 1 (ICAM1),

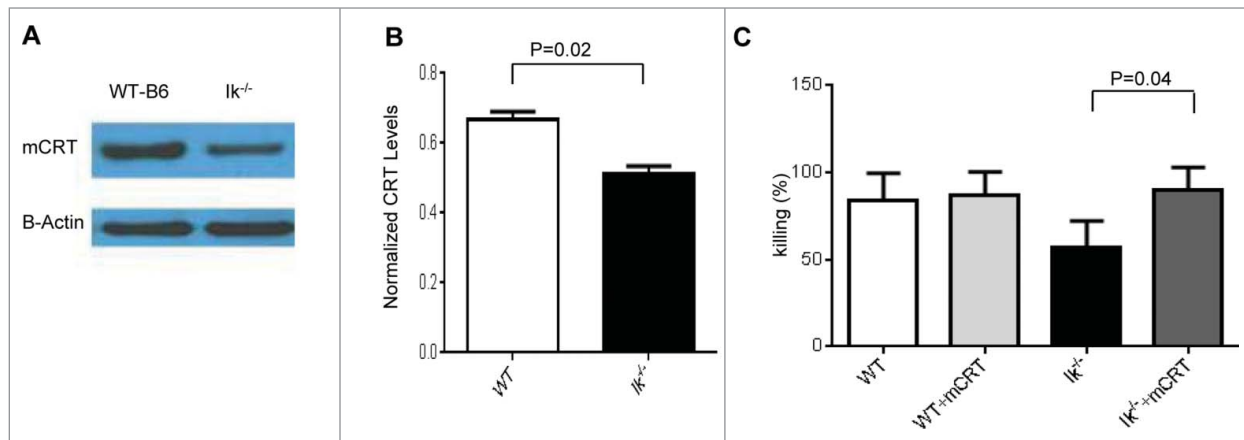


Figure 4. Calreticulin expression in dendritic cells from *Ik*^{-/-} animals and forced expression-enhanced CTL responses. **(A, B)** Spleen CD11c⁺ DCs were isolated from either B6-WT or *Ik*^{-/-} animals using CD11c-micobeads and calreticulin expression was measured by western blotting. **(A)** The absence of *Ik* decreased expression of CRT. **(B)** Normalized CRT level relative to actin expression. **(C)** BMDCs were transduced with CRT and empty vector control and used as stimulators for generating CTLs in bulk mixed-lymphocyte reaction (MLR). These generated CTLs were used as effectors against MBL-2 tumors for the ⁵¹Cr release assay.

CD47, and CD24 in the *Ik*-deficient animals was similar to that in WT animals (Fig. 3A, B).

We recently demonstrated that TSA cross-presentation on professional hematopoietic APCs is required for optimal GVL responses.⁶ Therefore, to determine whether the decreased TSA response is due to defects in cross-presentation on *Ik*^{-/-} APCs, we examined the expression of CRT, which plays an important role in regulating phagocytic function and regulating antigen cross-presentation of TSA.¹⁷ We first hypothesized that CRT expression would be decreased in *Ik*^{-/-} APCs. We examined the expression of CRT in spleen DCs isolated from either B6 WT or *Ik*^{-/-} animals and found reduced expression of CRT in *Ik*^{-/-} DCs (Fig. 4A–B). To determine whether the reduction of CRT in *Ik*^{-/-} DCs was critical for the lack of CTL responses against TSA, mCRT was transduced into BM-derived DCs from *Ik*^{-/-} animals and CTL function was tested with a ⁵¹Cr release assay. We found that *Ik*^{-/-} DCs with restored expression of CRT exhibited completely recovered CTL function (Fig. 4C). These data suggested that Ikaros in DCs might regulate CTL function in a CRT-dependent manner.

Ikaros deficiency in both hematopoietic APCs and leukemia cells contributes to lack of an optimal GVL response

Finally we determined whether absence of *Ik* only in the leukemic cells, and not in professional hematopoietic APCs, contributes to GVL resistance. To this end, we used the same BMT model but added p185 (*Ik* wild-type tumors), or *Ik*6 (*Ik* dominant negative tumors) at the time of BMT. Similar to the MBL-2 tumor model, allogeneic [*Ik*^{-/-}→B6Ly5.2] animals showed greater GVHD without concomitant GVL responses compared to allogeneic WT[B6→Ly5.2] animals when they received p185 tumors (Fig. 5A). Allogeneic T cells co-transferred with p185 from allogeneic [*Ik*^{-/-}→B6Ly5.2] animals on day 14 after allo-HCT demonstrated comparable cytotoxicity against p185 to that of allogeneic WT[B6→Ly5.2] animals (Fig. 5B). Conversely,

allogeneic WT [B6→Ly5.2] chimeras that received *Ik*6 showed better GVL responses than allogeneic [*Ik*^{-/-}→B6Ly5.2] animals (Fig. 5C). Allogeneic T cells co-transferred with *Ik*6 also effectuated greater apoptosis in T-cell cytotoxic killing assays (Fig. 5D). These data suggest that *Ik* deficiency in both non-leukemic and leukemic cells collectively contributes to GVL resistance (Table 1).

Discussion

Relapse of primary disease and GVHD are the greatest obstacles to improving the long-term outcome after allo-HCT.³ The GVL response is necessary to prevent relapse; however, GVL is tightly linked with GVHD therefore increasing GVL comes at the cost of exacerbating GVHD. Thus, separating GVL from GVHD remains the central issue in allo-HCT. Clinical data suggest that certain diseases such as ALL demonstrate greater relapse rates after allo-HCT, i.e., decreased GVL despite that fact that the patients suffer from severe GVHD. The reason for the reduced GVL in ALL could be multifactorial. Moreover, whether the decrease in GVL is due to defects in antigen presentation in addition to leukemia-intrinsic properties remains unknown. Here, we explored whether *Ik* deficiency in host hematopoietic cells contributes to GVL resistance. Although both host and donor APCs, specifically DCs as they are the most potent APCs, are important regulators of GVHD,^{16,18,19} the role of APCs in GVL is poorly understood. We have recently demonstrated that cross-presentation on professional hematopoietic APCs is critical for optimal GVL responses.^{5,6} Evidence suggests that professional APCs play an important role in generating effective CTLs for elimination of recipient tumor cells.^{20,21} Donor T cells that recognize alloantigens are critical for mediating GVHD and GVL and tumor-specific antigens also contribute to GVL. Recent data suggest that DCs may play an essential role in inducing

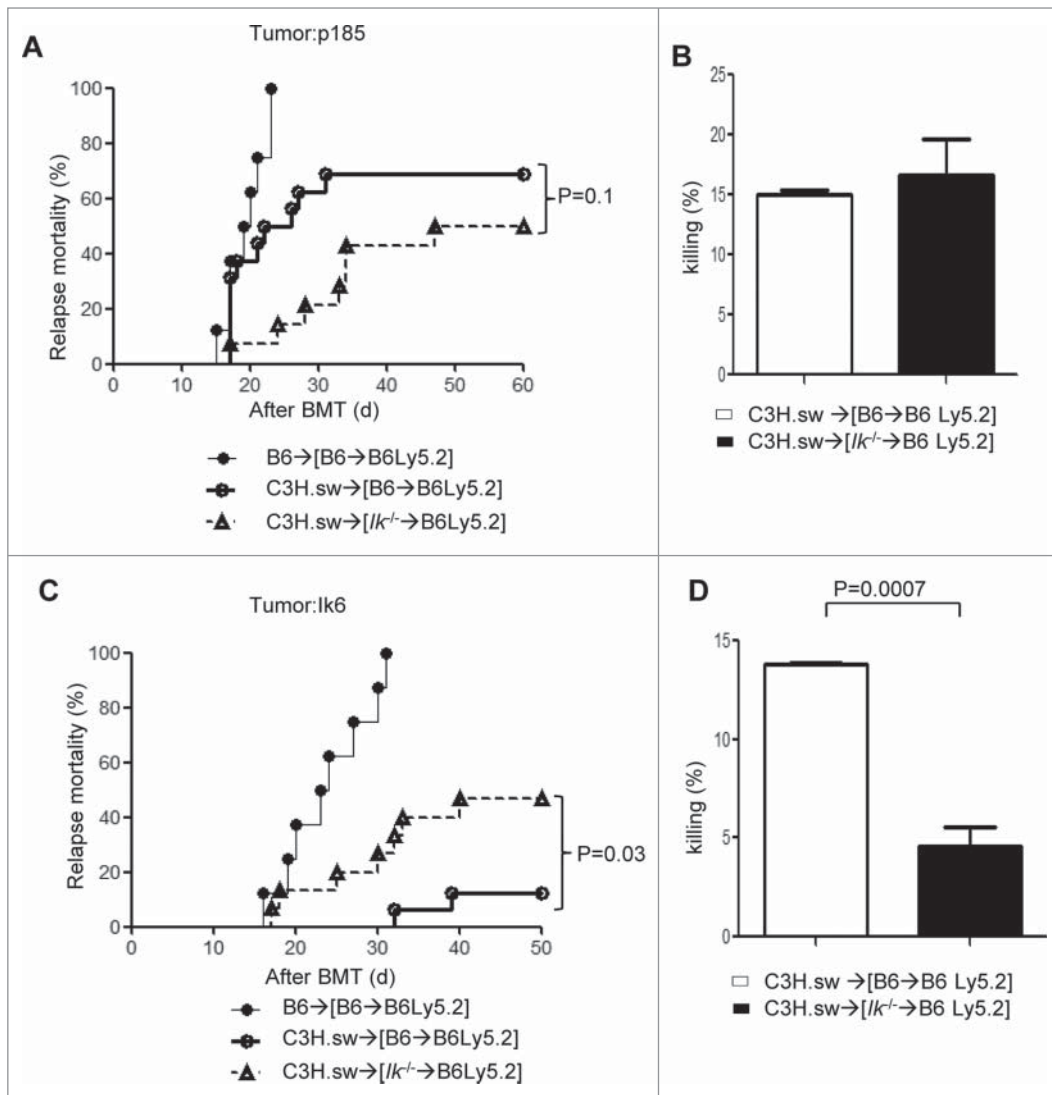


Figure 5. Ikaros deficiency in both hematopoietic APCs and leukemia cells ameliorates GVL responses. WT B6Ly5.2 animals were lethally irradiated with 11 Gy and infused with 5×10^6 BM cells and 5×10^6 splenocytes from syngeneic Ly5.1 WT B6 or *Ik*^{-/-} B6 donors. These animals [B6→B6Ly5.2] or [IkDN^{+/-}→B6Ly5.2]) were used as recipients 4 months later. Mice were irradiated with 9 Gy and transplanted with 1×10^6 CD90⁺ T cells together with 5×10^6 BM cells from either syngeneic B6 or allogeneic MHC-matched or multiple miHA-mismatched C3H.sw donors concurrently with syngeneic P185 (Ik WT tumor) or Ik6 (Ik DN tumor). **(A)** Tumor mortality data for P185 200/mouse. (●) B6→[B6→B6Ly5.2] (n = 8), (◐) C3H.sw→[B6→B6 Ly5.2] (n = 16), (◑) C3H.sw→[Ik^{-/-}→B6Ly5.2] (n = 14). Data are combined from 3 independent experiments. **(B)** CTL assay. Donor CD8⁺ T cells were isolated from spleen (n = 3–4) at day 14 following allo-HCT and used as effector T cells against P185 tumor for ⁵¹Cr-release assay. One representative dataset from 3 independent experiments is shown. Data are given as mean ± standard deviation. **(C)** Tumor mortality data of Ik6 200/mouse. (●) B6→[B6→B6Ly5.2] (n=8), (◐) C3H.sw→[B6→B6 Ly5.2] (n = 16), (◑) C3H.sw→[Ik^{-/-}→B6Ly5.2] (n = 15). Data are combined from 3 independent experiments. **(D)** CTL assay. Donor CD8⁺ T cells were isolated from spleen (n = 3–4) at day 14 following allo-HCT and used as effector T cells against Ik6 tumor for ⁵¹Cr-release assay. One representative dataset from 3 independent experiments is shown. Data are given as mean ± standard deviation.

Our previous data suggest that deficiency of *Ik* in host APCs aggravated the severity of GVHD in a Notch-dependent manner.⁷ Herein, we demonstrated that the absence of *Ik* in host APCs did not concomitantly enhance GVL, irrespective of severe GVHD, in [*Ik*^{-/-}→B6 Ly5.2] animals of the MHC-matched multiple MiHA mismatched C3H.sw→B6 model. We also used a different tumor (EL-4) and confirmed that *Ik*-deficient host APCs demonstrated equivalent GVL responses to WT chimeras. GVT responses are usually tightly linked with GVHD severity. Thus, this uncoupling of GVHD and GVL at the level of antigen presentation is a novel observation.

Host MHC class II⁺ APCs play a critical role in CTL responses in mixed chimera models.²² However, the role of APC subsets and their intrinsic molecular mechanisms in GVL are poorly understood. Using *Batf3*^{-/-} animals, we recently found that host CD8⁺ DCs play an important role in mediating optimal GVL responses. We also found that host APCs stimulated by TLR3 agonist enhanced the GVL effect without concomitant aggravation of GVHD.⁶ It was also recently shown that absence of the Ik–Notch pathway in host APCs increased GVHD.⁷ These data suggest

that host type APCs are required for optimal GVL responses following allo-HCT. These observations, together with our current findings, indicate that *Ik* expression in host APCs is a critical regulator of GVHD and GVL. We found that *Ik*^{-/-} DCs highly expressed CD205 (DEC205), which plays an important role in antigen capture and presentation²³ and is predominantly expressed

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Table 1. Summary of GVL experiments. WT and *Ik*-deficient host APCs showed an equivalent GVL effect for P185, *Ik*-WT tumor. However, *Ik* deficiency in both hematopoietic APCs and leukemia cells ameliorates GVL responses

APC/Tumor	P185 (<i>Ik</i> WT)	<i>Ik6</i> (<i>Ik</i> DN mutation)
WT APC	++	++
<i>Ik</i> ^{-/-} APC	++	+

in CD8 α ⁺ DCs.²⁴ CD8⁺CD205⁺ DCs also regulate T-cell immunity and homeostasis by increasing the generation of regulatory T cells (Tregs).²⁵ Enhancement of CD205 function in DCs with tumor antigen-specific vaccine increased tumor-specific CTLs and resulted in tumor regression in some patients with solid tumors, such as melanoma.²⁶ On the other hand, the expression of antigen-presenting related receptors other than CD205 in *Ik*^{-/-} DCs was comparable to that of WT DCs. Our previous data also demonstrated that expression of co-stimulatory molecules such as CD80, CD86, CD40, CD83, and PDL1 was much lower in splenic *Ik*^{-/-} DCs compared with WT DCs,⁷ suggesting that enhancing allo-stimulatory function of *Ik*^{-/-} DCs contributes to GVHD but does not increase GVL is not in receptors including CD205, and co-stimulation molecules dependent manner. The effects of *Ik* deficiency are not dependent on CD205 and the co-stimulatory molecules.

To further explore why *Ik*-deficient non-leukemic hematopoietic APCs cannot enhance GVL in spite of greater GVH responses, we focused on calreticulin expression in DCs. DCs express CRT on their surface^{13,14,27} and this CRT expression facilitates TSA cross-presentation via MHC class I molecules to cytotoxic T cells.^{15,27-30} CRT plays an important role in protein folding and the maintenance of MHC class I assembly pathways,^{31,32} and CRT expression on the cell surface enhances cellular phagocytic uptake.¹⁴ We examined CRT expression on both WT and *Ik*^{-/-} BMDCs by western blotting and found that *Ik*^{-/-} BMDCs showed lower CRT expression than WT cells (Fig. 4A-B). Therefore, we next examined whether forced increase in CRT expression by gene transduction rescues CTL function in *Ik*^{-/-} BMDCs and found a significant increase in cytotoxic killing by T cells primed by CRT transduced *Ik*^{-/-} BMDCs, compared with WT (Fig. 4C). Therefore, the mechanism underlying the lack of concomitant increased TSA expression may be related to a lack of increasing cross-presentation capacity as a result of a deficiency on CRT expression. Future studies will determine the direct mechanism of the molecular regulation of CRT by *Ik* and explore the intrinsic mechanisms by which *Ik* in host hematopoietic-derived APCs separately regulates GVHD and GVL, as well as whether other molecular mechanisms such as Notch-dependent signaling play an important role in GVL responses.

Our observations provide novel insights into clinical observations that are particularly germane to ALL. Among all leukemias, ALL with *Ik* deficiency or mutation has a poor outcome and is at high risk of relapse after allo-HCT.³³ Although the lack of sufficient GVL in these patients is relatively well established, available evidence suggests that these patients have a similar incidence and severity of GVHD.

Ik is not only an essential transcription factor as a tumor suppressor for leukemogenesis, such as in ALL^{10,34} and for blast crisis (BC) of chronic myeloid leukemia (CML),³⁵ but *Ik* (*Ikzf1*) deficiency and mutation in leukemic cells is also associated with a poor prognosis in ALL.^{11,12,36,37} However, whether *Ik* deficiency in leukemic cells alone contributes to resistance, i.e., by increasing relapse, has not been explored. In our study using *Ik* DN tumor cell lines (*Ik6*), we determined whether the absence of *Ik* in leukemic cells makes them more resistant to GVL, or whether the resistance is due to primarily to its absence in the non-leukemic host hematopoietic cells. We found that when allogeneic [*Ik*^{-/-}→B6Ly5.2] animals, in which *Ik* was deficient in the host non-leukemic hematopoietic cells, were co-transplanted with *Ik6*, the recipients showed greater tumor relapse and diminished cytotoxic responses against the tumor targets, compared to allogeneic WT[B6→B6Ly5.2] animals. These data suggest that *Ik* deficiency in leukemic cells and in non-leukemic hematopoietic APCs collectively contributes to GVL resistance.

In conclusion, we demonstrate that the absence of *Ik* in both host hematopoietic APCs and leukemic cells enhances GVL resistance despite an increased severity of GVHD. Therefore, strategies that overcome the impact of *Ik* deficiency by targeting both hematopoietic-derived APCs and hematopoietic leukemic cells are needed.

Materials and Methods

Mouse

C57BL/6 (B6, H-2^b, CD45.2⁺), C3H.sw (H-2^b), BALB/c (H-2^d), and B6 Ly5.2 (H-2^b, CD45.1⁺) mice were purchased from the Jackson Laboratory (Bar Harbor, ME) and National Cancer Institute (Frederick, MD). *Ik*^{-/-} mice were provided by Dr. Winandy (Boston University, Boston, MA) and backcrossed to B6 for μ (Boston 8 generations.³⁸ BMT recipient mice were housed and maintained as in Supplemental methods. All animals were cared for under regulations reviewed and approved by the University Committee on Use and Care of Animals, based on University Laboratory Animal Medicine guidelines.

Generation of bone marrow chimeras

Bone marrow (BM) chimeras ([B6→B6 Ly5.2] and [*Ik*^{-/-}→B6 Ly5.2]) were generated as described previously.^{5,7} Briefly, B6 Ly5.2 WT animals were subjected to 11-Gy total-body irradiation (TBI;¹³⁷Cs Source) on day -1 and then injected intravenously with 5 × 10⁶ BM cells and 5 × 10⁶ whole spleen cells from WT B6 or the *Ik*^{-/-} donor mice on day 0. Donor hematopoietic chimerism was confirmed using the CD45.2 monoclonal antibody 3–4 months after BMT (donor type >95.0%).

Bone marrow transplantation

BMT was performed as described previously.^{5,7,18} Briefly, splenic T cells from donors were enriched while the BM was depleted of T cells by autoMACS (Miltenyi Biotec, Bergisch Gladbach, Germany). [B6→B6Ly5.2] and [*Ik*^{-/-}→B6Ly5.2]

animals received 9 Gy TBI (^{137}Cs source) on day -1 and were injected intravenously with 0.5×10^6 CD8 $^+$ T cells and 5×10^6 T-cell depleted BM (TCD-BM) from either syngeneic B6 or allogeneic C3H.sw donors on day 0. Survival was monitored daily and recipients' body weight and GVHD clinical scores³⁹ were measured weekly.

Induction of leukemia and lymphoma

Tumors (MBL-2 or EL-4, H-2 b) were introduced during BMT at 2 different doses as described previously.^{5,6} MBL-2 is a Moloney-murine leukemia virus-induced T-cell lymphoma^{40,41} whereas EL-4 is a chemically-induced T-cell lymphoma.⁴² Both are of B6 origin (H-2 b) and are extensively used as models of acute leukemia and lymphoma. P185 BCR-ABL1 Arf null (WT for *IKZF1*) and the dominant negative isoform Ik6 were kindly provided by Dr. Mullighan (St. Jude Children's Research Hospital, Memphis, TN). To observe GVL responses, we used a lower dose of tumor cells (MBL-2 0.5–1 $\times 10^4$ /mouse) because we had already established that with injection of this dose the animals cannot reject the tumor and injection results in tumor infiltration and engraftment of the liver and spleen with characteristic nodule formation or hind-limb paralysis caused by spinal cord infiltration.^{5,6,43,44} The mice that showed less tumor signal among [B6 \rightarrow [B6 \rightarrow B6Ly5.2]] animals demonstrated severe hind-limb paralysis induced by the tumor and were euthanized by the criteria established by the UCUCA protocol. Moreover, we additionally confirmed the cause of death in the allogeneic animals, regardless of BLI, by postmortem examination for tumor and histopathology. We attributed death to tumor if tumor was present at necropsy. Death was attributed to GVHD only if no tumor was evident by flow cytometry or autopsy and there was histologic evidence of GVHD. Mice surviving beyond the observation period of BMT were sacrificed for histologic evaluation to determine leukemia- and lymphoma-free survival.

Luciferase $^+$ MBL-2 cell line

MBL-2 cells were transduced with a third-generation lentivirus coexpressing GFP and firefly luciferase (Luc) as previously described.^{6,45}

Bioluminescence imaging

Bioluminescence imaging was performed with a cryogenically cooled CCD camera (IVIS, Caliper Life Sciences). Acquisition and analysis of images were performed as previously described.⁴⁵ All animals were imaged 10 min after intraperitoneal (i.p) injection with 100 μL of 40 mg/mL firefly D-luciferin (Biosynth International Inc.). Animals were imaged for 30 s to 5 min, depending on the signal strength. All animals were maintained under isoflurane anesthesia in a 37°C environment.

FACS analysis

FACS analyses was performed as described previously.^{6,7} Briefly, to analyze chimerism and donor T-cell expansion, splenocytes from transplanted mice were resuspended in FACS wash buffer (2% FBS in PBS) and stained with conjugated monoclonal antibodies (mAbs) as follows: fluorescein isothiocyanate (FITC)-

conjugated mAbs to mouse CD45.2 and CD229.1 (Biolegend); phycoerythrin (PE)-conjugated mAbs to CD24, CD47, CD54, CD107a, and CD205 (Biolegend), granzyme B, CD172a (signal regulatory protein α (SIRP- α) and CD209 (eBioscience); allophycocyanin (APC)-conjugated mAbs to CD4, CD229.1, and perforin; and PerCPcy5.5-conjugated mAb to CD8 (Biolegend). Cells were stained and fixed with 1 \times BD FACSTM Lysing Solution (BD Bioscience), and analyzed using BD AccuriTMC6 cytometer (BD Bioscience).

Analysis of MBL-2-specific T-cell responses

Donor MBL-2 specific CD8 $^+$ T cells were analyzed on day 21 after BMT with the immune dominant PE-conjugated peptide tetramer CCLCLTVFL epitope, a *gag*-encoded protein of the Friend/Mononey/Rauscher (FMR) retrovirus that is recognized in the context of H-2 b .^{40,41} An H-2D b restricted influenza peptide (D b PA, SLENFRAYV) was used as negative control. Tetramers were made by the NIH tetramer core facility (Atlanta, GA). Briefly, splenocytes from transplanted mice were resuspended in FACS wash buffer (2% FBS in PBS), stained with conjugated monoclonal antibodies, and incubated with PE-conjugated gag specific tetramer. Cells were washed twice with FACS wash buffer, fixed with 1 \times BD FACSTM Lysing Solution (BD Biosciences), and analyzed using BD AccuriTMC6 cytometer (BD Bioscience).

^{51}Cr release assay

Splenic CD8 $^+$ T cells (4×10^6 /ml) were isolated from transplanted mice 21 d after allo-BMT, purified with a CD8 $^+$ T-cell isolation kit (Miltenyi Biotec), and used as effector cells. MBL-2, P185 BCR-ABL1 Arf null, and Ik6 tumor cells were labeled by incubating 2×10^6 cells with 2 MBq of Na $_2^{51}\text{CrO}_4$ (PerkinElmer Life) for 2 h at 37 °C in a 5% CO $_2$ atmosphere and were used as target cells. After washing, 4×10^3 labeled targets were resuspended and added to triplicate wells at varying effector-to-target ratios and then incubated for 4 h. Maximal and minimum release was determined by the addition of Triton-X (MP Biomedicals) or media alone to targets, respectively. Supernatants were transferred to a Luma plate (PerkinElmer) after 4 h and ^{51}Cr activity was determined using an autogamma counter (Packard).

Calreticulin expression and overexpression transfection to DCs

Spleens were harvested from either B6-WT or *Ik* $^{-/-}$ animals and incubated with collagenase D (1 mg/mL, Roche Diagnostics, Indianapolis, IN) in a 37°C/5% CO $_2$ incubator for 45 min. Cells were mashed and purified with CD11c microbeads (Miltenyi Biotec) and auto MACS (Miltenyi Biotec). Calreticulin expression was detected by western blotting.

For overexpression transfection into DCs, we used the plasmid pMSV-mCRT (a gift from Dr. Malini Raghavan, Department of Microbiology and Immunology, the University of Michigan Medical School), which contains full-length mouse wild-type calreticulin cDNA. The pMSV-mCRT plasmid was transfected into Oneshot Top 10 competent cells (Life

Technologies) and transfectants were selected by ampicillin. Positive strains were expanded and purified using an Endofree Plasmid Purification kit (Qiagen). Purified pMSV-mCRT plasmids and empty vector control were transfected into DCs from WT-B6 or *Ik*^{-/-} animals in a 6-well plate (at 80% confluency) using X-treme GENE HP DNA Transfection Reagent (Roche) according to the manufacturer's instruction. After incubation at 37°C for 48 h, the transfected cells were collected for assays.

Statistical analysis

The Mann-Whitney *U* test was used for statistical analysis of *in vitro* data, and the Wilcoxon rank test was used to analyze survival data. A *p* value <0.05 was considered statistically significant.

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Disclosure of Potential Conflicts of Interest

No potential conflicts of interest were disclosed.

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