



Stem cell transplantation

The mechanistic study behind suppression of GVHD while retaining GVL activities by myeloid-derived suppressor cells

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Abstract

Graft-versus-host disease (GVHD) is a major barrier to the widespread use of allogeneic hematopoietic stem cell transplantation (allo-HSCT) for treating hematologic malignancies. Myeloid-derived suppressor cells (MDSCs) have been recognized as crucial immunosuppressive cells in various pathologic settings. Here, we investigated whether the unique functional properties of MDSCs could be harnessed to control allo-HSCT-associated GVHD. Using multiple murine GVHD/GVL models including both MHC-mismatched and miHA-mismatched, we demonstrated that treatment with CD115+ MDSCs efficiently suppressed GVHD but did not significantly impair graft-versus-leukemia (GVL) activity, leading to 80 and 67% protection in treated mice in GVHD and GVL models, respectively. The mechanism for this dissociation of GVHD from GVL, specifically the emergence of donor-derived NKG2D⁺ CD8 T cells with a memory phenotype in MDSC-treated recipient mice, was identified. NKG2D expression on donor T cells was required for eradication of allogeneic lymphoma cells. Furthermore, long-term surviving MDSC recipients that exhibited cytolytic activities against allogeneic leukemia cells had a significantly increased percentage of T regulatory cells and, more importantly, NKG2D⁺ CD8 T cells. These findings indicate that MDSCs can be used as a novel cell-based therapy to suppress GVHD while maintaining GVL activities through selective induction of NKG2D⁺ CD8 memory T cells.

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Introduction

Allogeneic hematopoietic stem cell transplantation (allo-HSCT) is a potentially curative immunotherapy for a variety of hematopoietic malignancies and non-malignancies [1] and has been increasingly used to treat solid tumors [2]. However, utilization and full realization of the therapeutic merit of allo-HSCT procedures has been limited by the occurrence of life-threatening graft-versus-host disease (GVHD) [3, 4]. Despite the involvement of natural killer (NK) cells in certain allo-HSCT settings [5], T cells present in the grafts play a central role in the pathogenesis of GVHD [6, 7]. Furthermore, donor-derived T cells are also the principal effector cells in graft-versus leukemia (GVL) [8], the specific anti-tumor activity of allo-HSCT. Due to this dual role for donor T cells, approaches currently employed to diminish GVHD, for example, T-cell depletion and non-specific immunosuppressive drugs, are often associated with increased incidences of graft failure, tumor recurrence, opportunistic infection, and/or drug resistance [7, 9, 10]. Thus, developing new strategies to dissociate the suppression of GVHD from GVL remains a major challenge in allo-HSCT.

In recent years, there has been increasing interest in myeloid-derived suppressor cells (MDSCs) owing to accumulating evidence suggesting a critical role for MDSCs in the initiation, progression, and metastasis of tumors as well as in other pathologic conditions, including pathogenic infections, inflammation, and traumatic stress in animal models and human patients [11–19]. MDSCs can function to suppress T-cell responses both *in vitro* and *in vivo* via diverse mechanisms, e.g., production of nitric oxide (NO), reactive oxygen species (ROS), expression of arginase 1 and inducible nitric oxide synthase (iNOS), and/or secretion of interleukin (IL)-10 and transforming growth factor beta (TGF- β) [18, 20–23].

Although MDSCs may hamper the success of immune-based cancer therapy, multiple immunosuppressive properties of MDSCs, on the other hand, may endow them with great therapeutic potential in the fields of auto-immune disease and transplantation, where immune responses need to be limited. This concept has been supported by recent studies proposing a potential role for MDSCs in the GVHD treatment [24, 25]. However, the effect of MDSCs on GVL activities in allo-HCST recipients remains to be determined. We previously identified a major subset of MDSCs expressing the myeloid markers Gr-1, F4/80, and CD115 in tumor-bearing mice, and demonstrated that, in comparison to MDSCs usually defined as a Gr-1⁺CD11b⁺ population, CD115⁺Gr-1⁺F4/80⁺ cells not only display stronger suppressive capabilities but also induce the development of CD4⁺CD25⁺Foxp3⁺ T regulatory cells (Tregs) in tumor-bearing mice [26]. In this report, we demonstrate that upon adoptive transfer, CD115⁺Gr-1⁺F4/80⁺ MDSCs freshly isolated from tumor-bearing mice or *in vitro*-generated MDSC could efficiently suppress GVHD while maintaining GVL effects in both MHC-mismatched and miHA-mismatched GVHD/GVL models. The MDSCs mediated induction of NKG2D⁺ CD8 T cells is the key driver for the retention of GVL activities after allo-HSCT with MDSC treatment.

Materials and methods

Animals and cell isolation

C57BL/6, C3H.SW, BALB/c, and congenic (Thy-1.1) BALB/c mice were purchased from the National Cancer Institute (Frederick, MD) and the Jackson Laboratory (Bar Harbor, ME). MDSCs from Lewis lung carcinoma-bearing C57BL/6 mice were prepared as described [27]. The *in vitro*-derived MDSCs were generated as previously described [28, 29]. Briefly, C57BL/6 bone marrow (BM) cells were cultured in RPMI containing 10% FBS, 40 ng/

ml GM-CSF and 40 ng/ml IL-6 (Peprotech) for 5 days. BM cells from naive C57BL/6 mice underwent negative selection using anti-Thy-1.2 and the Thy-1.2 negative cells were used as T-cell depleted bone marrow (TCDBM). Donor T cells were isolated from splenocytes of naive C57BL/6 mice using a negative selection kit (R&D systems, Minneapolis, MN).

Mixed lymphocyte reaction and MDSC suppression assay

In all, 2×10^5 C57BL/6 splenocytes (responder) were co-cultured with an equal number of irradiated (25 Gy) BALB/c splenocytes or non-irradiated BALB/c splenocytes (stimulator) in the absence or presence of various numbers of CD115⁺ MDSC isolated from C57BL/6 tumor-bearing mice. [³H]-Thymidine was added for the last 8 hours of 4-day or 5-day culture. For Tregs induction, C57BL/6 and BALB/c splenocytes (5×10^6 /well) were co-cultured with 1.25×10^6 purified MDSCs for 5 days. Foxp3 expression was measured by flow cytometry.

GVHD and GVL models

The establishment of GVHD models was previously described [30, 31]. Briefly, BALB/c or C3H.SW mice (8–14-weeks-old, female) were lethally irradiated and randomly distributed into each group. Within 24 hours after irradiation, recipients were left untreated or reconstituted via tail vein injection with donor-derived cells, C57BL/6 TCDBM (5×10^6 /mouse), purified C57BL/6 splenic T cells (5×10^5 /mouse for Balb/c recipients and 2×10^6 /mouse for C3H.SW recipients) and MDSCs (3×10^6 /mouse for Balb/c recipients and 6×10^6 /mouse for C3H.SW recipients) isolated from C57BL/6 tumor-bearing mice or from *in vitro* cultures. Mice that received MDSC on Day 0 were given two additional injections of MDSCs (3 or 6×10^6 /mouse) on days 4 and 10 after transplantation. Animals were monitored daily for GVHD symptoms and overall survival. For survival rate analysis, at least 22 mice were used for each *in vivo* experiment. For histopathological analysis, specimens obtained at day 21–30 were fixed in formalin and tissue sections were stained with hematoxylin and eosin. The pathologist was blinded to the group allocation during the analysis.

In the experiments designed for expansion and activation of donor T cells, MDSC-treated recipients were given MDSCs once on day 0, and mice were sacrificed on days 7 or 14 after transplantation. In the GVL experiments, recipients were co-transplanted with A20 cells (1×10^5 /mouse) unless otherwise specified. Animals found to have hepatic or lymphoid tumor nodules at post-mortem were categorized as death due to tumor. Mice that died without tumors

but with clear signs of GVHD were considered deaths due to GVHD.

Antibodies and tumor cells lines

All fluorochrome-labeled and purified mouse antibodies and corresponding isotype controls were purchased from commercial source and listed in Table S1. Flow cytometric surface staining was performed as described [18]. Intracellular staining for Foxp3 and granzyme B was performed per manufacturer's instructions (Mouse Regulatory T cell Staining Kit, eBioscience). For intracellular staining of interferon (IFN) γ , splenocytes isolated from each group ($n=3$) were individually cultured for 6 hours in the presence or absence of PMA (20 ng/ml) and ionomycin (1 μ g/ml), with the addition of monensin for the last 4 hours. Data were acquired on a FACSAria II (BD Biosciences) and analyzed using Flowjo software (Tree Star, Inc., Ashland, OR). A20, YAC-1 and EL4 tumor cell lines were purchased from the American Type Culture Collection. L1 (BALB/c line 1 lung carcinoma) and MCA-26 (BALB/c-derived colon carcinoma) are maintained in our laboratory. Periodic mycoplasma detection tests were performed in all cell lines used in this study.

Cytotoxic T lymphocyte (CTL) assay

Using Thy-1 as the marker, effector T cells were purified following 5-day MLR culture in the absence or presence of MDSC or directly from pooled splenocytes of treated mice ($n=3$ mice) then normalized for H-2K^b⁺ CD8⁺ T-cell numbers based on FACS data. The purified effector T cells were then co-cultured for 4 hours with target cells (A20, YAC-1, EL4, L1, and MCA-26, 1×10^4 /well) at various ratios. Anti-natural-killer group 2, member D protein (NKG2D) and anti-CD3 (5 μ g/ml, Biolegend, San Diego, CA) were added during the CTL assays. Supernatants were collected from each well for measurement of lactate dehydrogenase (LDH) release (cytotoxicity assay kit, Promega, Madison, WI). Specific killing was calculated using the following formula: % cytotoxicity = $100 \times (\text{experimental release} - \text{effector spontaneous release} - \text{target spontaneous release}) / (\text{total target release} - \text{target spontaneous release})$.

Statistical analysis

Statistical differences in animal survival were analyzed by log-rank test. Differences between two groups were compared using unpaired *t*-test. Differences between three or more groups were compared using one-way ANOVA. All experiments were repeated at least three times. All the statistical analysis was performed with GraphPad Prism, version 6.05. $P < 0.05$ was considered statistically significant.

Study approval

All animal studies were approved by the IACUC at the Center for Comparative Medicine and Surgery of the Icahn School of Medicine at Mount Sinai and at Comparative Medicine at Houston Methodist Research Institute.

Results

MDSCs suppress allo-immune response in vitro

We first assessed the functional activity of MDSCs in an allogeneic mixed lymphocyte reaction (MLR). As shown in Fig. 1a, CD115⁺ MDSCs isolated from tumor-bearing mice exerted potent suppressive activity in a dose-dependent manner, and almost completely suppressed the allo-specific proliferation of C57BL/6 T cells, when used at high ratios of MDSCs versus C57BL/6 splenocytes (Fig. 1a). In contrast, CD115⁻ cells did not show suppressive activity even at a 1:1 ratio. Interestingly, the addition of MDSCs to the MLR cultures at a 1:4 ratio of MDSCs versus C57BL/6 splenocytes led to a marked augmentation of CD25⁺Foxp3⁺Tregs within the C57BL/6 CD4 T-cell compartment (Fig. 1b). These results indicate that MDSCs are able to significantly inhibit the proliferation of allo-reactive T cells in vitro, a mechanism, which may be partially mediated by Treg induction.

MDSCs effectively alleviate GVHD following allo-HSCT

We next sought to determine whether MDSCs would be effective in suppressing the allo-immune response in vivo, thus preventing GVHD following allo-HSCT. We adopted a well-characterized murine GVHD model in which MHC-mismatched mouse strains BALB/c (H-2^d) and C57BL/6 (H-2^b) mice were designated as recipients and donors, respectively. While lethally irradiated mice without treatment died within 10 days, all mice treated with TCDBM alone were healthy and survived for more than 100 days (Fig. 1c). As expected, mice that had received TCDBM plus donor-type T cells developed severe signs of GVHD as evidenced by loss of hair, hunched posture, diarrhea, and death in most of the animals between 15 and 25 days after transplantation. In contrast, when given MDSCs derived from tumor-bearing mice, along with TCDBM plus T cells, recipients were largely protected from GVHD lethality and approximately 80% of the animals survived for more than 100 days.

A striking feature of GVHD pathology is the occurrence of inflammation and tissue damage in specific target organs. Histological examination of liver, small intestine, and skin

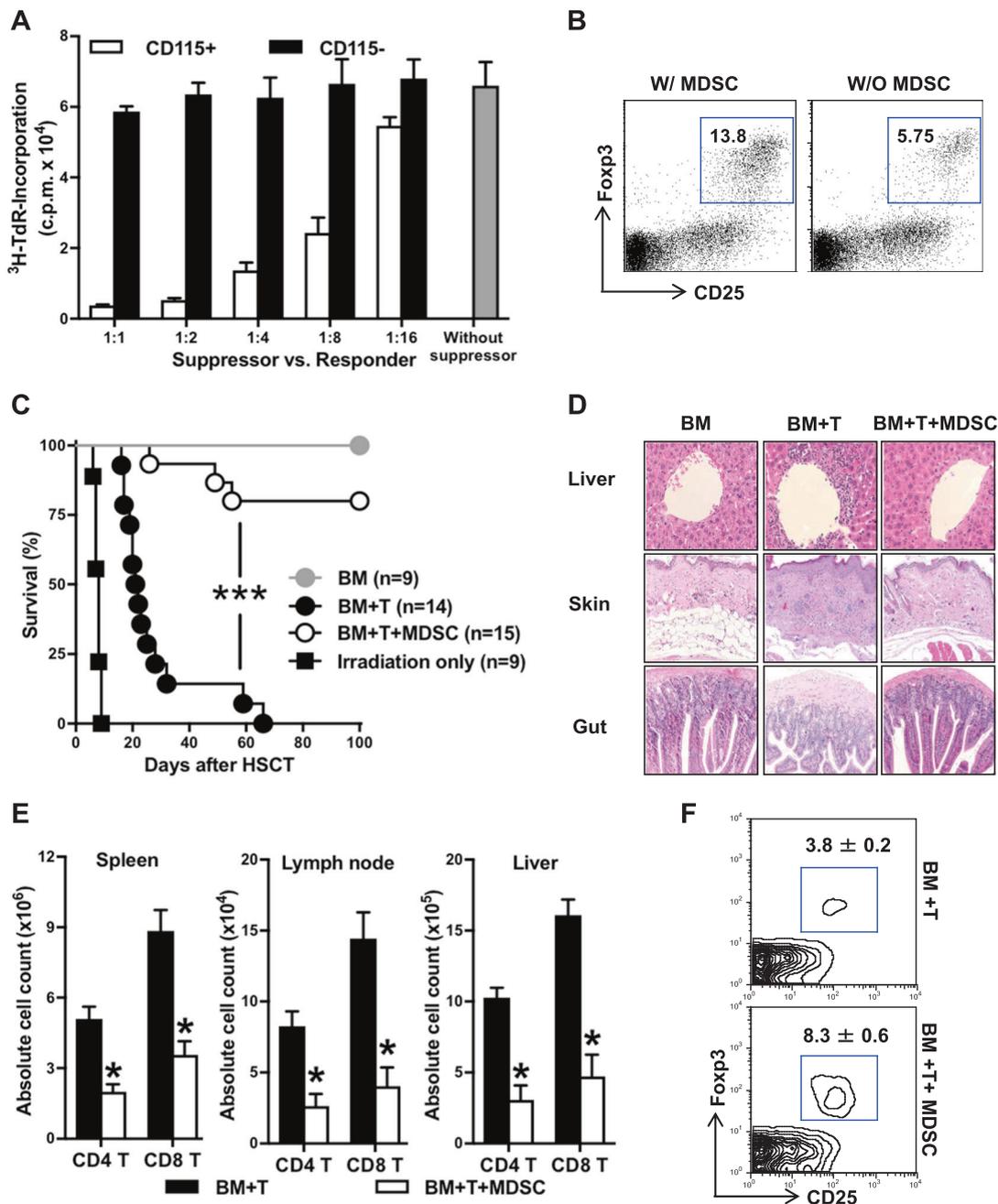


Fig. 1 MDSCs induce Tregs and suppress allogeneic T-cell responses in vitro and in vivo. **a** C57BL/6 splenocytes (responders, 2×10^5 /well) were co-cultured with an equal number of irradiated BALB/c splenocytes in the absence or presence of indicated ratios of CD115⁺ (suppressors) or CD115⁻ (control) cells from tumor-bearing C57BL/6 mice. T-cell proliferation was determined by measuring [³H]-thymidine incorporation during the last 8 h of a 4-day culture. **b** C57BL/6 and BALB/c splenocytes (5×10^6 /well) were cultured in the presence or absence of MDSCs isolated from tumor-bearing mice (1.5×10^6 /well) for 5 days. Treg population (CD4⁺CD25⁺Fcγp3⁺) was measured by flow cytometry. **c** The Balb/c GVHD model and MDSCs treatment was described in Methods. Survival curve of recipient mice with indicated treatments. ****P* < 0.0001, BM + T + MDSC versus

BM + T recipients. Data shown were combined from three independent experiments with reproducible results. **d** Histopathological analyses of GVHD target tissues. Liver, gut, and skin tissues from the mice received BM, BM + T, or BM + T + MDSC were collected on day 21 after transplantation and hematoxylin and eosin staining of tissue sections was performed for histopathological analysis. Representative photomicrographs are shown. **e** Absolute numbers of CD4 and CD8 T cells in spleen, lymph node and liver at day 7 post-treatment. Data are presented as mean ± S.D. (*n* = 3). **P* < 0.05. **f** Splenic Treg populations in indicated groups were analyzed by gating on H2K^bCD4⁺ cells on day 7 post-treatment. Data are presented as mean ± S.D. (*n* = 3 mice)

harvested on day 21 after allo-HSCT revealed dense lymphocyte infiltration in all examined tissues obtained from TCDBM plus T-cell-treated mice, particularly in the periportal area of the liver and dermis layer of the skin. The intestines from these mice manifested atrophy of villi and mucous membrane and crypt destruction. On the contrary, tissue samples isolated from MDSC-treated mice showed no significant histological changes (Fig. 1d and Fig. S1A), almost indistinguishable from those recovered from TCDBM-only recipients. These observations establish that MDSCs are highly effective in suppressing the development of GVHD, thus resulting in a great improvement in animal survival after allo-HSCT.

To gain insight into the regulation of GVHD by MDSCs, we further investigated the cellular mechanisms underlying this process. It is well established that upon recognition of host antigens, including MHC and miHAs (minor histocompatibility) antigens, donor-derived T lymphocytes become activated and undergo rapid and extensive expansion in the early course of GVHD following allo-HSCT [7]. Accordingly, MDSCs may function to suppress GVHD by restraining the expansion of allo-reactive T cells. To examine this possibility, we performed additional transplantation experiments. On day 7 after adoptive transfer, secondary lymphoid organs and livers were isolated from representative mice in each group and donor-derived T-cell subsets were profiled by flow cytometry. As determined by cell percentage (Fig. S1B) and absolute cell number (Fig. 1e), the proliferation of both donor-derived (H-2k^b positive) CD4 and CD8 T cells was significantly inhibited in MDSC recipients when compared to the group without MDSC treatment. The spleens of these mice (Fig. 1e, left panel) exhibited 46% and 55% decreases in the donor CD4 and CD8 subsets, respectively. The reductions were even more pronounced in mesenteric lymph nodes (66% and 71%, respectively, Fig. 1e, middle panel) and liver (69% and 70%, respectively, Fig. 1e, right panel). Similar results were also obtained with samples harvested on day 14 after transplantation (Fig. S1C). Taken together, these data suggest that upon adoptive transfer, MDSCs remain suppressive *in vivo*, thus causing suppression of allo-reactive T-cell expansion in both lymphoid organs and GVHD target organs.

Since MDSCs induced Treg development in MLR (Fig. 1b) and Tregs have been shown to be highly effective in suppressing GVHD in murine models [32–37], we evaluated whether Tregs were increased in the GVHD setting. Indeed, as early as day 7 after allo-HSCT, recipients that received MDSCs had at least a two-fold higher percentage of Tregs in the splenic H-2K^bCD4⁺ population when compared to those mice that had not received MDSCs (Fig. 1f). This increase in Treg frequency was even more evident in long-term (day 107) surviving mice that had received MDSCs (Fig. S1D). These results suggest that

MDSC-induced Tregs may contribute to the long-term suppression of GVHD. We further evaluated the therapeutic effect of *in vitro*-generated MDSCs. Similar to tumor-bearing-mouse-derived MDSCs, *in vitro*-generated MDSCs also ameliorated GVHD severity and mortality in both MHC-mismatched and MHC-matched miHA-mismatched murine GVHD models (Fig. 2).

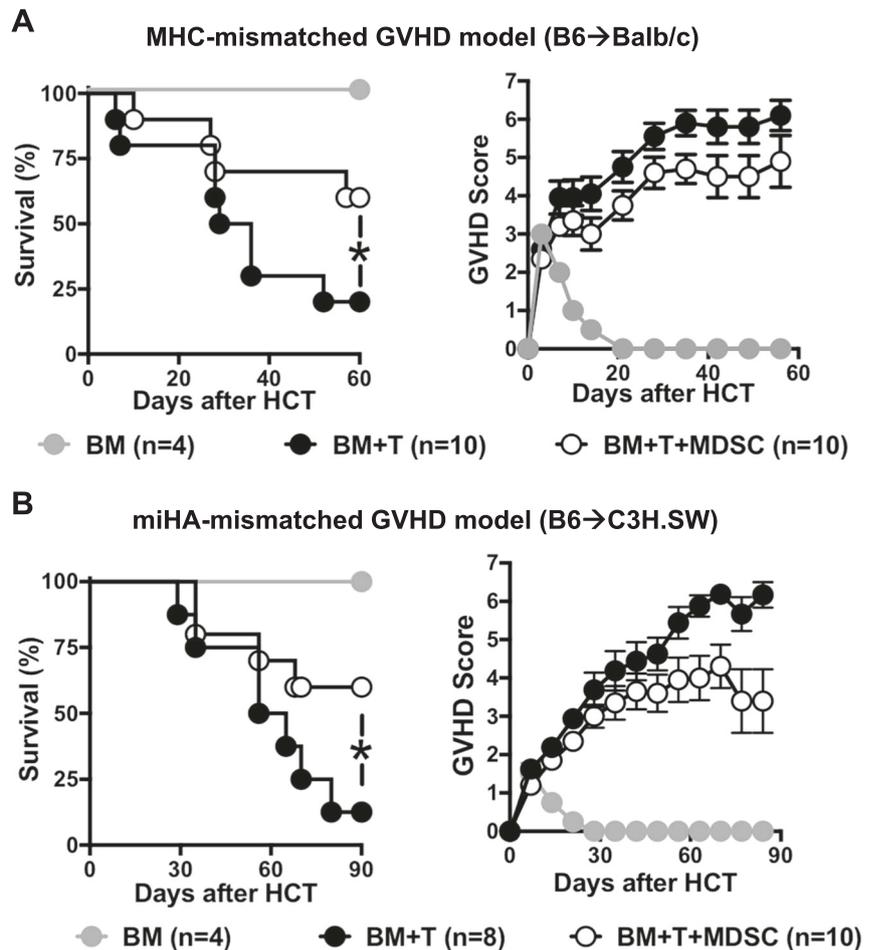
MDSCs preserve allo-reactive T-cell cytotoxicity and GVL activity

The profound suppression of GVHD by MDSCs prompted us to examine whether the GVL activity of donor-derived T cells is maintained in the presence of MDSCs. Lethally irradiated BALB/c mice were co-transplanted with A20 cells (a B-cell lymphoma/leukemia cell line, H-2^d, 1×10^5 /mouse) to induce tumor development in the hosts. Consequently, 100% of recipients that received TCDBM plus A20 died before day 30 from progressive leukemia (Fig. 3a). Mice that had received TCDBM and A20 plus T cells also died between day 20 and day 51 after transplantation, due to GVHD-related damage rather than tumor growth, as exhibited by the development of severe signs of acute GVHD but no tumor formation as evidenced by the absence of tumor nodules in the liver and lymph nodes of 6 mice examined pre- or post-mortem. Strikingly, 67% of tumor-challenged animals that received TCDBM plus T cells supplemented with tumor-bearing-mouse-derived MDSCs survived over 100 days. These long-term surviving mice displayed neither GVHD nor leukemia symptoms, suggesting that, in the absence of GVHD, donor T cells were able to efficiently eradicate tumor cells. These results indicate that MDSCs can actively suppress GVHD while retaining beneficial GVL activity after allo-HSCT.

To explore the underlying mechanisms by which MDSCs maintain GVL activity, we examined the activation of donor-derived T cells and their functional activity. Based on the expression levels for a panel of activation markers, including CD69 (Fig. S2A), CD44, CD62L (Fig. S2B), and CD25 (data not shown), the activation of donor-derived CD4 and CD8 T cells in the spleens or lymph nodes isolated from MDSC-treated recipients at day 7 after allo-HSCT was not significantly altered compared with their counterparts that had not received MDSC treatment. Intracellular staining of re-isolated cells after incubation *in vitro* in the presence or absence of phorbol myristate acetate (PMA) plus ionomycin stimulation also revealed no substantial differences in IFN γ production as evaluated by the percentages of IFN γ -producing CD4⁺ and CD8⁺ cells in the total splenic CD4 or CD8 T cells recovered from the two groups (Fig. 3b).

The granzyme B/perforin pathway represents one of the primary mechanisms mediating direct T-cell cytotoxicity. Although we have not examined the expression of perforin,

Fig. 2 Treatment of in vitro-derived MDSCs alleviates GVHD in multiple murine models. GVHD models and MDSC treatment was described in Methods. Survival curves and GVHD scores for MHC-mismatched GVHD model (a) and miHA-mismatched GVHD model (b) * $P < 0.05$, data are presented as mean \pm S.D



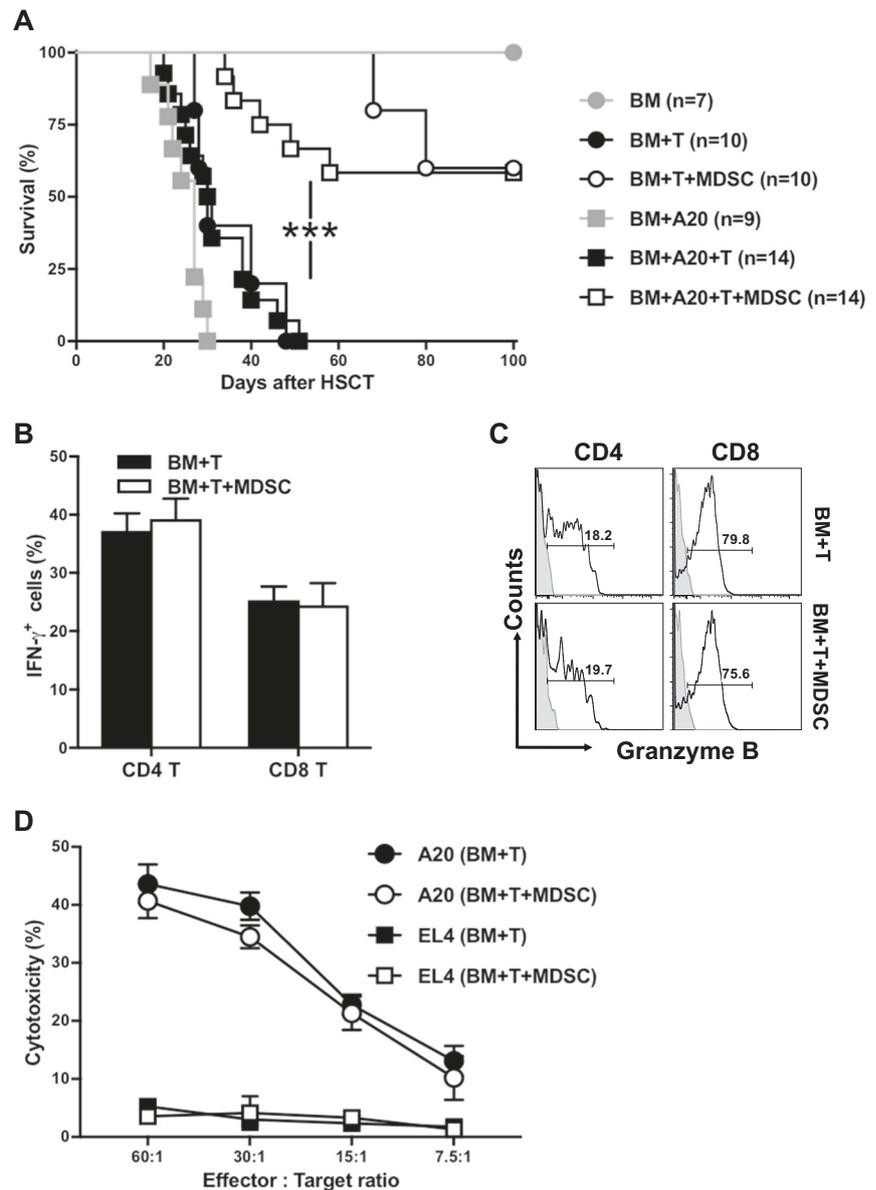
no apparent changes in granzyme B expression were detected in the donor-derived T lymphocyte compartment (Fig. 3c), suggesting that the cytotoxic capabilities of donor-derived CD8 T cells were unaffected by MDSCs. To verify this, we conducted cytolytic assays using splenic T cells isolated on day 14 after allo-HSCT. As shown in Fig. 3d, while CD8 T cells derived from mice that had or had not received MDSCs showed minimal killing against syngeneic tumor targets EL4 (H-2^b), they both exhibited similar cytotoxicity profiles against allogeneic target A20 cells (H-2^d). Indeed, cytotoxicity against allogeneic tumor cells was preserved among T cells isolated from MDSC-treated long-term survivors (Fig. S2C, D). These data indicate that adoptively transferred MDSCs do not alter the activation or functional activity of donor-derived T cells, which may explain in part why the GVL activity of allo-reactive T cells is retained in the presence of MDSCs.

Upregulation of NKG2D during allogeneic T-cell responses in the presence of MDSCs

Since treatment with MDSCs limited only the expansion, not the activation or effector function of donor T cells, we

next determined the cellular composition of MLR in the presence or absence of tumor-bearing-mouse-derived MDSCs. We found that the ratio of responder (H2K^{b+}) CD8-to-CD4 T cells in the MLR absence of MDSCs was significantly lower than that in the MLR in the presence of MDSCs (Fig. S3A), suggesting that CD4 T cells are more susceptible to suppression by MDSCs. We further confirmed this notion by assessing the proliferation of responder T cells labeled with carboxyfluorescein succinimidyl ester (CFSE). There was a 66% decrease in proliferating CD4 T cells versus a 48% decrease in proliferating CD8 T cells in MLR in the presence of MDSCs when compared to MLR without MDSCs (Fig. S3B). We further analyzed the phenotypes of T cells in MLR. The addition of MDSCs to MLR led to significant increases in the percentages of CD44⁺CD62L⁻ CD4 T cells and CD44⁺CD62L⁺ CD8 T cells (Fig. 4a and Fig. S3C). Strikingly, NKG2D, a crucial co-stimulatory molecule that is involved in anti-tumor immunity of CD8 T cells was also significantly upregulated when MDSCs were added to the MLR culture (Fig. 4a and Fig. S3C). We further investigated NKG2D⁺ T cells in murine GVHD/GVL models. Consistent with the in vitro MLR results, the

Fig. 3 Treatment of tumor-bearing-mouse-derived MDSCs preserves GVL activity. The GVHD/GVL model with Balb/c recipients and MDSC treatment was described in Methods. **a** Survival curve. *** $P < 0.001$, BM + A20 + T + MDSC versus BM + A20 + T. Data shown are combined from three independent experiments. **b** The frequency of donor-derived IFN γ -producing CD4 and CD8 splenic T cells. **c** Granzyme B expression in donor-derived splenic T-cell subsets (BM + T and BM + T + MDSC). Gray line indicates Isotype. **d** Cytolytic activity of donor CD8 T cells isolated from mice that had received allo-HSCT. Splenic T cells isolated on day 14 after allo-HSCT were co-cultured with tumor cells, A20 (H-2K^d) or EL4 (H-2K^b) for 4 hours at indicated ratios. The extent of cytotoxicity was determined by LDH release. Data are presented as mean \pm S. D., $n = 3$



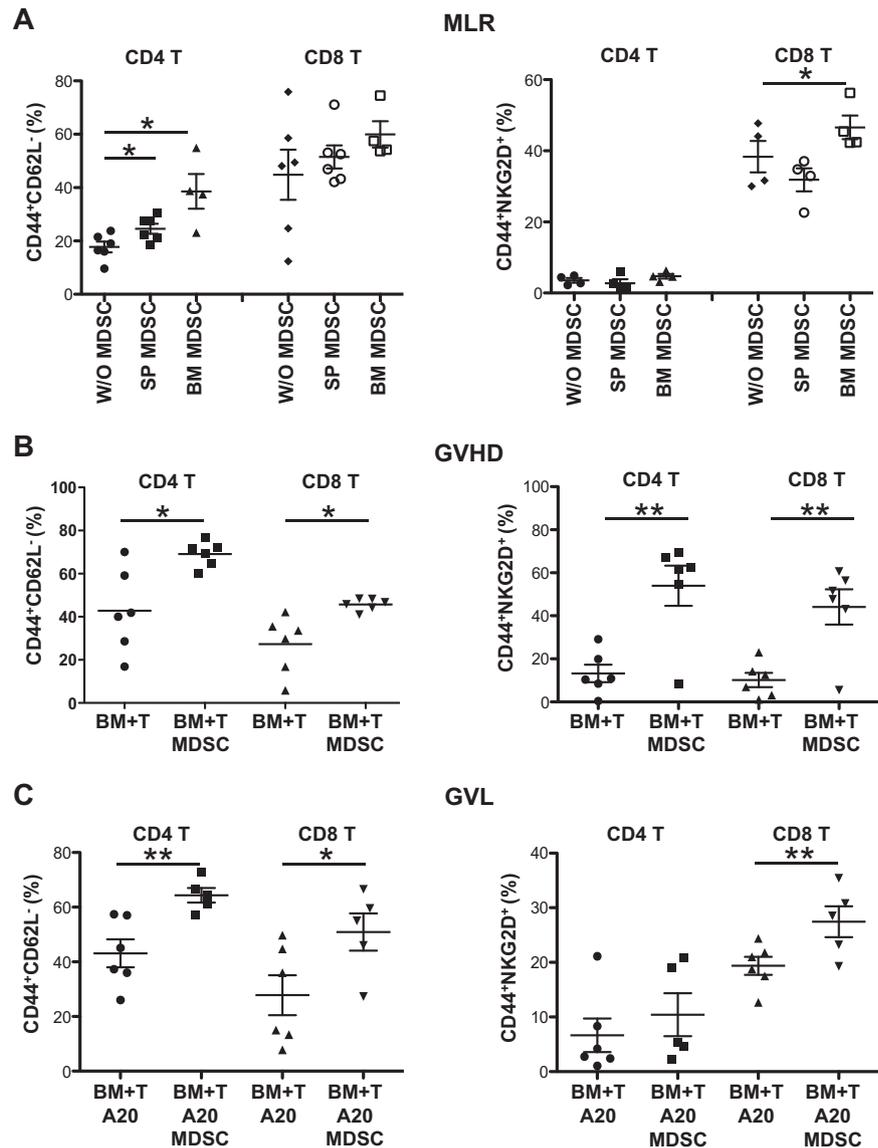
frequency of NKG2D⁺ CD8 T cells was significantly increased in mice that had received tumor-bearing-mouse-derived MDSCs treatment at day 30 after allo-HSCT (Fig. 4b, c). Interestingly, MDSC-treated mice also showed increase of NKG2D expression on CD4 T cells in the GVHD model, indicating that NKG2D⁺ CD4 T cells may also be involved in preventing and/or inhibiting GVHD pathogenesis after MDSCs treatment. Moreover, we observed a further increase in CD44⁺NKG2D⁺ and CD44⁺CD62L⁻ T cells on day 100 after allo-HSCT (Fig. S3D). The results suggest that NKG2D⁺ CD8 T cells may play a pivotal role in GVL activity after MDSC treatment. Since NKG2D⁺ T cells were upregulated, we further evaluated whether MDSCs express NKG2D ligands. Indeed, MDSCs from tumor-bearing mice expressed a significant level of

NKG2D ligands, e.g. RAE-1 and a certain amount of MULT-1 as detected by flow cytometric analysis (Fig. S4).

The cytolytic activity of MDSC-conditioned allo-reactive T cells depends on NKG2D and CD3

To determine the role of NKG2D in cytotoxicity against tumors mediated by MDSC-conditioned allo-reactive T cells, we performed cytotoxicity assays in the presence of CD3 or NKG2D blocking antibodies. T cells purified from 5-day MLR cultures containing tumor-bearing-mouse-derived MDSCs exhibited stronger cytotoxicity against allogeneic A20 cells and Yac-1 cells than control MLR cultures without MDSCs (Fig. 5a). A significantly lower level of cytotoxicity against syngeneic EL4 cells was also

Fig. 4 Tumor-bearing-mouse-derived MDSCs induce NKG2D⁺ memory T cells. Flow cytometric analysis of CD44⁺CD62L⁻ and CD44⁺NKG2D⁺ splenic T-cell populations in 5-days MLR (a), GVHD (b), and GVL models (c) at day 30 after allo-HSCT and MDSC treatment. SP MDSC and BM MDSC represent MDSCs isolated from spleen and bone marrow of tumor-bearing mice, respectively. * $P < 0.05$, ** $P < 0.01$, $n = 6$. The data are from three independent experiments



detected in T cells purified from MLR cultured in the presence or absence of MDSCs (data not shown). The cytotoxicity of T cells purified from control or MDSC-containing MLR was significantly inhibited in the presence of NKG2D or CD3 blocking antibodies. Importantly, purified T cells from MDSC-treated mice exhibited strong cytotoxic activity against allogeneic A20 cells and YAC-1 that was partially blocked by CD3 and NKG2D blocking antibodies (Fig. 5a, b), indicating that both NKG2D and CD3 are required for the cytotoxicity of donor T cells.

Since NKG2D was upregulated on CD8 T cells in the presence of MDSCs, we further examined the role of the NKG2D ligand RAE-1 in T-cell-mediated cytolytic activity. We first assessed the expression of RAE-1 on tumor cells (Fig. S5). Indeed, not only did A20 tumor cells normally express high levels of RAE-1, but irradiation also increased

RAE-1 expression (Fig. S5A). However, results from cytotoxicity assays with RAE-1 blockade (Fig. S5B) suggested that RAE-1 may not be the only ligands involved in cytotoxicity mediated by NKG2D⁺ T cells. We further evaluated the role of NKG2D in cytotoxicity of MDSC-conditioned T cells against RAE-1 low-expressing non-hematopoietic tumor cells. As expected, T cells from MLR culture or mice treated with tumor-bearing-mouse-derived MDSC exhibited cytotoxicity against RAE-1 low-expressing tumor cells (Fig. S5C, S5D and Fig. 5c). However, NKG2D blockade did not affect the cytotoxicity against these tumor cells (Fig. 5c), suggesting that diverse mechanisms are involved in cytotoxicity of MDSC-conditioned T cells against RAE-1 low-expressing non-hematopoietic malignancies and RAE-1 high-expressing hematopoietic malignancies.

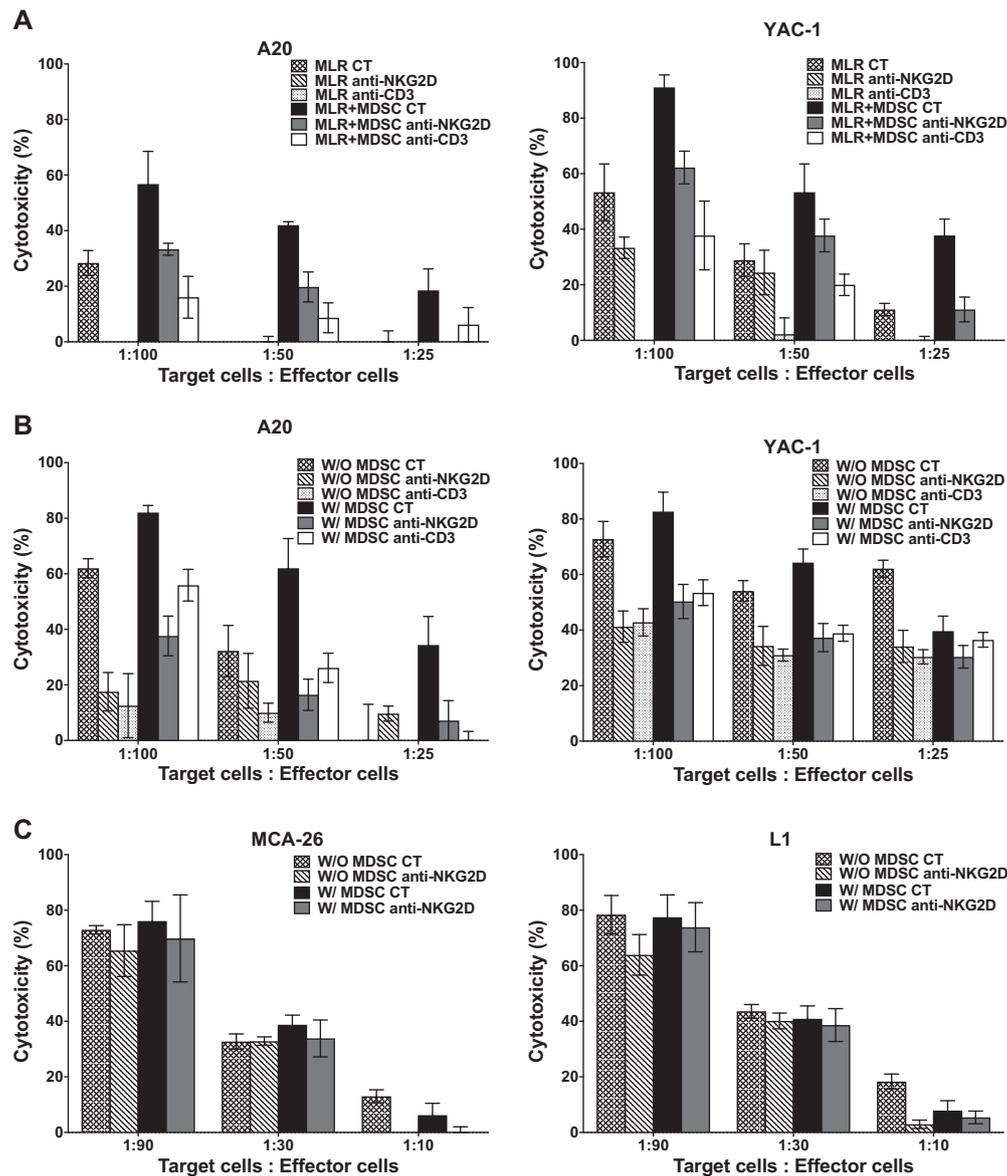


Fig. 5 NKG2D and CD3 are essential for MDSC-conditioned T-mediated tumor lysis. **a** Cytotoxicity of T cells from MLR in the presence or absence of tumor-bearing-mouse-derived MDSCs. T cells were purified from MLR and their cytotoxicity against allogeneic A20 cells and Yac-1 cells was determined at various effector: target ratios in the presence or absence of CD3 or NKG2D blocking antibodies. **b** Cytotoxicity of donor T cells purified from Balb/c mice at day 14 after allo-HSCT with or without treatment of tumor-bearing-mouse-

derived MDSC. The cytotoxicity against allogeneic A20 cells and Yac-1 cells at various effector: target ratios was assessed in the presence of CD3 or NKG2D blocking antibodies or isotype control. **c** Cytotoxicity of donor T cells against RAE-1 low-expression tumor cells, L1 and MCA-26, in the presence of NKG2D blocking antibodies. Splenic donor T cells were purified from Balb/c mice at day 14 after allo-HSCT with or without treatment of tumor-bearing-mouse-derived MDSCs. Data are presented as mean \pm S.D., $n = 3$

NKG2D-expressing T cells are essential for maintaining GVL activity after MDSCs treatment

We next determined whether the NKG2D expression is required for GVL activity after MDSC treatment. As shown in Fig. 6a, tumor-bearing-mouse-derived MDSC treatment resulted in significantly prolonged survival of A20-bearing mice that had received allo-HSCT. In mice where the NKG2D⁺ cell population was depleted using anti-NKG2D

antibody (BM + T + A20 + MDSC + anti-NKG2D), GVL activity was substantially inhibited compared to mice treated with the corresponding control IgG (BM + T + A20 + MDSC + IgG) ($P = 0.046$) (Fig. 6a). Furthermore, we used NKG2D knockout T cells (KOT) and bone marrow cells (KOBM) as donor cells to clarify whether donor-derived NKG2D⁺ T cells were required for the GVL activity. MDSC treatment significantly increased survival in mice receiving wild-type donor cells (BM + T + A20 + MDSC)

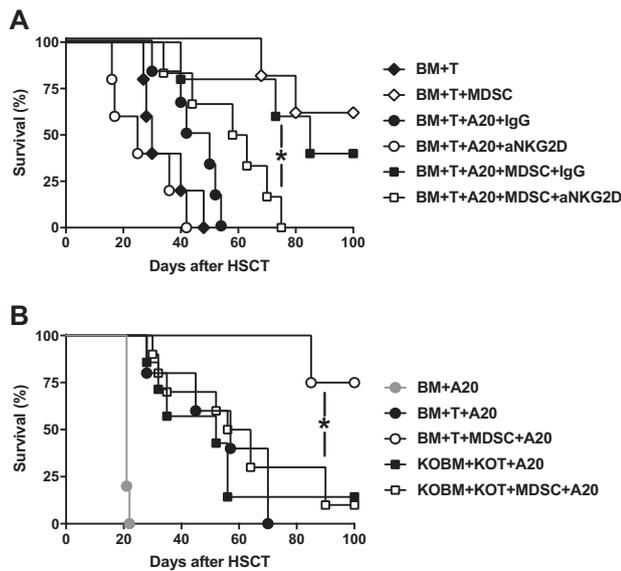


Fig. 6 NKG2D is required for maintaining GVL activity in MDSCs-treated mice after allo-HSCT. **a** NKG2D blocking antibodies abrogate GVL effect in allo-HSCT mice treated with MDSC. Lethally irradiated BALB/c mice were grafted with C57BL6 TCDBM and T cells, tumor-bearing-mouse-derived MDSC, and A20. MDSC recipients were given MDSCs on days 0, 4, and 10 after transplantation. For depletion of NKG2D population, mice were continuously injected with anti-NKG2D blocking antibodies (100 $\mu\text{g}/\text{mouse}/\text{day}$) for 10 days starting on day 14 post-transplantation. $*P < 0.05$, $n = 5$ to 6 per group. **b** NKG2D-deficient T cells fail to mediate GVL effect in allo-HSCT mice treated with MDSCs. BM and T cells were isolated from NKG2D-deficient or wild-type C57BL6 mice then transplanted into lethally irradiated BALB/c mice that were injected with A20. MDSC recipients received tumor-bearing-mouse-derived MDSCs on days 0, 4, and 10 post-transplantation. Significantly prolonged survival is observed in the BM/T/A20/MDSC versus KOBM/KOT/A20/MDSC group. $*P < 0.05$, $n = 5$ to 10 per group

compared to mice that had not received MDSC treatment (BM + T + A20) ($P = 0.0065$). However, MDSC treatment lost its protective effect in mice that received NKG2D-deficient donor cells (KOBM + KOT + MDSC + A20) compared to mice that received wild-type donor cells ($P = 0.0332$) (Fig. 6b). These results suggest that NKG2D expression on T cells is crucial for preserving GVL activity after MDSCs treatment.

Discussion

The data presented here strongly suggest that MDSC treatment might represent a cell-based tolerogenic therapy that is capable of preventing lethal GVHD while retaining sufficient GVL activity to eradicate leukemia cells. Co-transplantation with MDSCs resulted in suppressed development of lethal acute GVHD but did not jeopardize the GVL activity of allo-reactive T cells or the long-term

immunity of treated mice, making MDSC administration an attractive approach to GVHD prophylaxis.

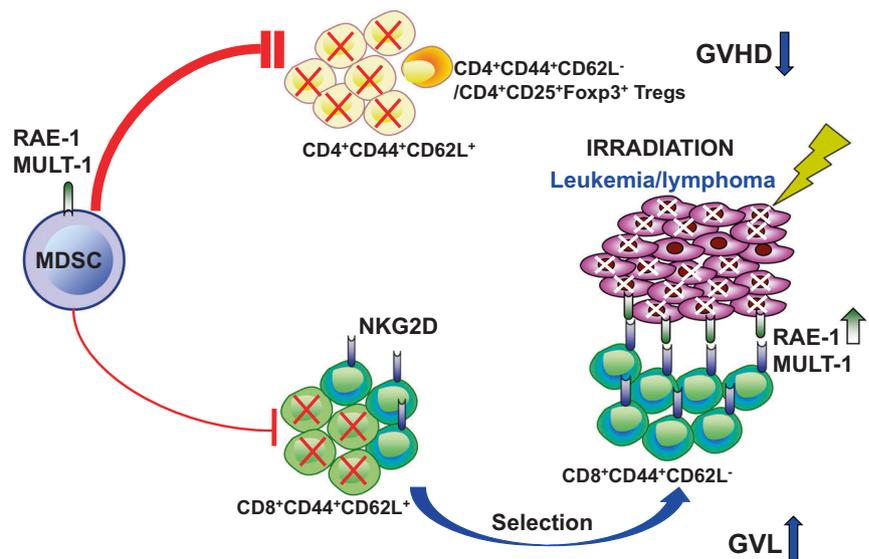
In a recent report, in vitro-generated MDSCs have been shown to prevent GVHD by inducing T helper 2 (Th2) cells while preserving the cytotoxic activity of donor T cells against H-2K mismatched thymoma (JM6) cells [38]. In contrast to our study, Tregs were not increased in MDSC-treated group. This may be attributed to the functional differences between MDSCs isolated from tumor-bearing mice and in vitro-generated MDSCs in the presence of granulocyte-macrophage colony-stimulating factor (GM-CSF) and G-CSF.

In this study, we have shown that MDSCs exhibit a potent immunosuppressive activity, effectively suppressing allo-specific immune responses and, remarkably, augmenting regulatory T cells ($\text{CD4}^+\text{CD25}^+\text{Foxp3}^+$). We also observed that MDSC treatment favored the development of a central and effector memory T cell and NKG2D^+ T-cell populations in vitro and in vivo, which may be partially attributed to the expression of NKG2D ligands on MDSC (Fig. S4) and on tumor cells (Fig. S5). Mechanistically, both CD3 and NKG2D played a role in T-cell-mediated cytolytic activity against tumor cells in vitro. To assessing the contribution of NKG2D to GVL activities by using NKG2D depleting Ab and NKG2D-deficient mice as the source of bone marrow and T cells, we demonstrated that donor-derived NKG2D positive T cells play an important role in the GVL effect observed in allo-HSCT mice treated with MDSC.

NKG2D^+ CD8 T cells have been shown to mediate GVHD and GVL responses [39]. Since NKG2D ligands were transiently induced on normal tissues of allo-HSCT mice, transient NKG2D blockade from days 0 to 6 after transplantation was able to attenuate GVHD while preserving GVL effect of NKG2D^+ CD8 T cells. In our GVHD/GVL model, the acute allo-responses were suppressed by MDSCs, thereby preventing the attack on normal cells that transiently expressed NKG2D ligands by NKG2D^+ CD8 T cells, which depends on both CD3 and NKG2D (Fig. 5). At later stages of GVHD, NKG2D induction in normal tissues of recipient mice and the suppressive activities of MDSC waned thereby allowing donor NKG2D^+ CD8 T cells to regain GVL function and eradicate allogeneic tumor cells without negatively affecting normal tissues in recipient mice.

Based on our findings in the present study, we propose the model in Fig. 7 that MDSC can prevent GVHD by suppressing the early expansion of donor-derived T cells and inducing Tregs. Furthermore, MDSC can augment GVL activity by favoring the development and activation of donor-derived NKG2D^+ CD8 T cells while retaining their effector function, such as tumor cytolytic activity as well as $\text{IFN}\gamma$ production. In conclusion, we have demonstrated that

Fig. 7 Proposed mechanisms by which MDSCs differentially regulate GVHD and GVL effect. NKG2D ligand-expressing MDSCs favor Tregs expansion that prevent GVHD, while preserving GVL activity by selection expansion of NKG2D⁺ memory CD8 T cells. RAE-1 Retinoic acid early inducible 1, MULT-1 mouse UL16-binding protein-like Transcript 1, NKG2D Natural killer group 2, member D; Tregs regulatory T cells



NKG2D expression on T cells can be induced by MDSCs during allogeneic immune responses and its expression is essential for maintaining GVL activity after allo-HSCT. Our findings may provide new insight into understanding the mechanism of how MDSCs control GVHD while preserving GVL activity.

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Compliance with ethical standards

Conflict of interest The authors declare that they have no conflict of interest.

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Supplementary Materials for

The mechanistic study behind suppression of GVHD while retaining GVL activities by myeloid-derived suppressor cells.

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This PDF file includes:

Figure S1. MDSCs prevent tissue damage and inhibit allo-reactive T cell proliferation

Figure S2. MDSCs preserve cytotoxicity of allo-reactive T cells.

Figure S3. MDSCs inhibit allogenic T cell proliferation while up-regulate NKG2D expression.

Figure S4. MDSCs express NKG2D ligands.

Figure S5. NKG2D ligand RAE-1 expression on tumor cells.

Table S1. Antibodies for flow cytometry and functional assays.

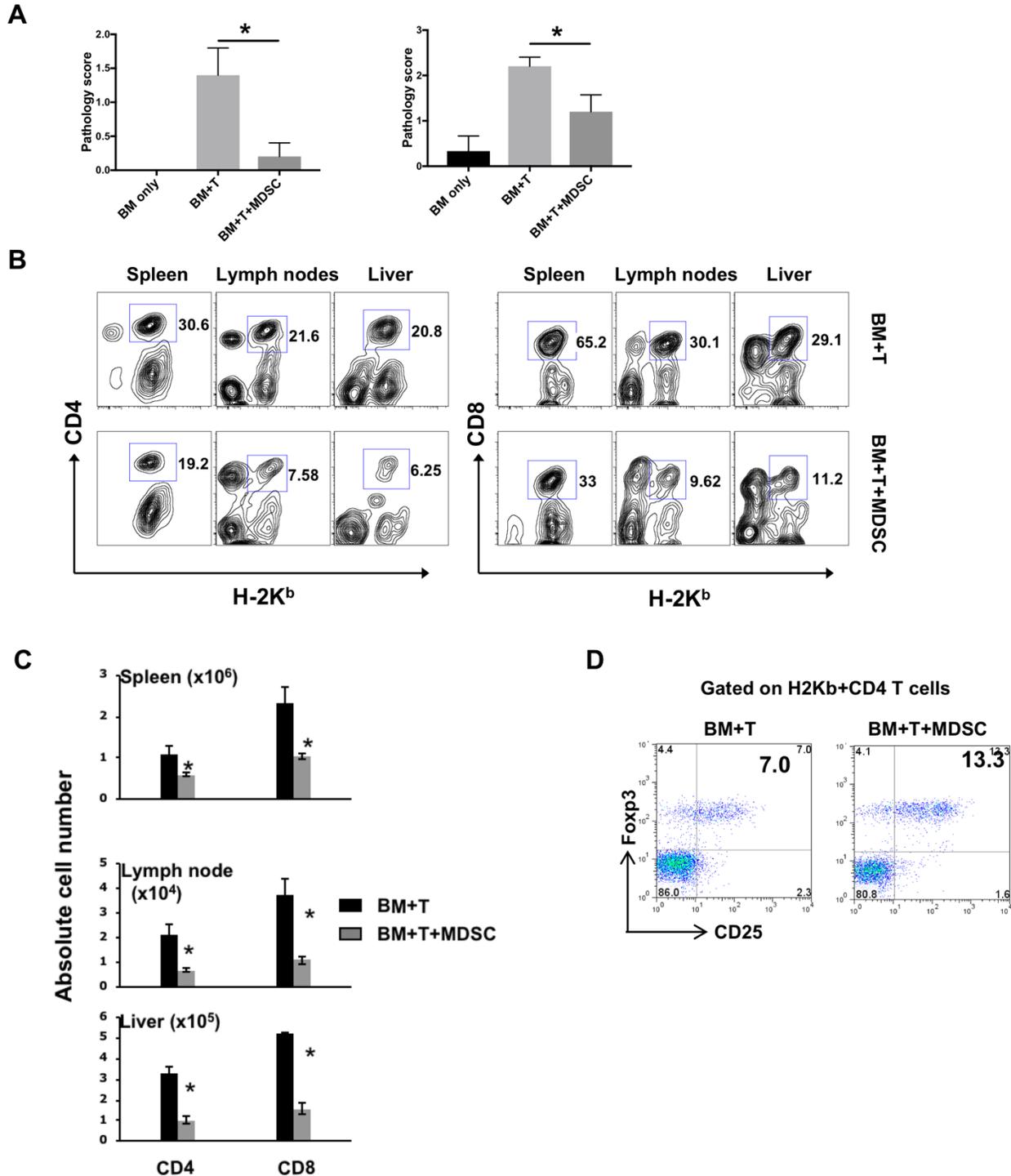


Figure S1. MDSCs prevent tissue damage and inhibit allo-reactive T cell proliferation. The Balb/c GVHD model and tumor-bearing-mouse-derived MDSC treatment are described in Methods. (A) Histopathological analyses of GVHD target tissues (skin, left panel; liver, right panel). Tissues were collected at day 21-30 after allo-HSCT, data are presented as mean \pm S.D., $n = 4-5$ per group, * $P < 0.05$. (B) Representative flow plots of donor H2K^b+CD4 and CD8 cell populations in the spleen, lymph node and liver tissues of mice. (C) Absolute CD4 and CD8 T cell numbers in spleen, lymph node and liver day 14 post-treatment. Data are presented as mean \pm S.D. ($n = 3$). * $P < 0.05$. (D) Flow analysis of Splenic Treg population at day 107 after allo-HSCT.

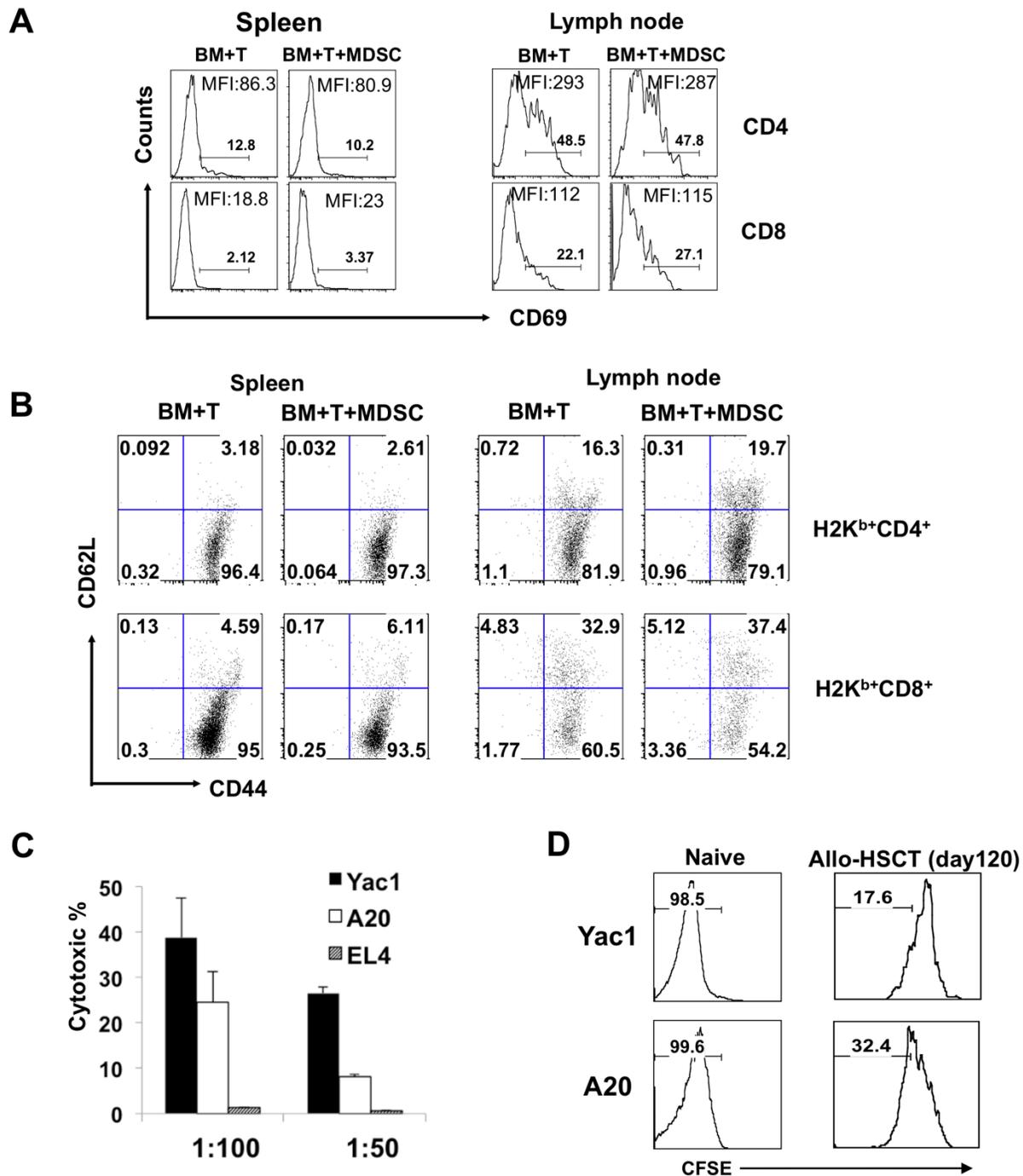


Figure S2. MDSCs preserve cytotoxicity of allo-reactive T cells. The Balb/c GVHD model and tumor-bearing-mouse-derived MDSC treatment are described in Methods. **(A)** Representative flow plots of CD69 expression on donor T cells at day 7 after allo-HSCT. **(B)** Representative flow plots of CD62L and CD44 expression on donor T cells at day 7 after allo-HSCT. **(C)** Cytotoxicity of T cells isolated from two MDSCs-treated long-term (100 days after allo-HSCT) surviving mice against allogeneic A20, Yac-1 and syngeneic EL4 cells. **(D)** Tumor rechallenge in MDSCs-treated long-term (120 days after allo-HSCT) surviving mice. CFSE-labeled Yac1 or A20 were injected into MDSC-treated mice and naïve BALB/C mice. After 40 hrs, spleens were harvested and the proliferation of CFSE-labeled tumor cells was checked by flow cytometry.

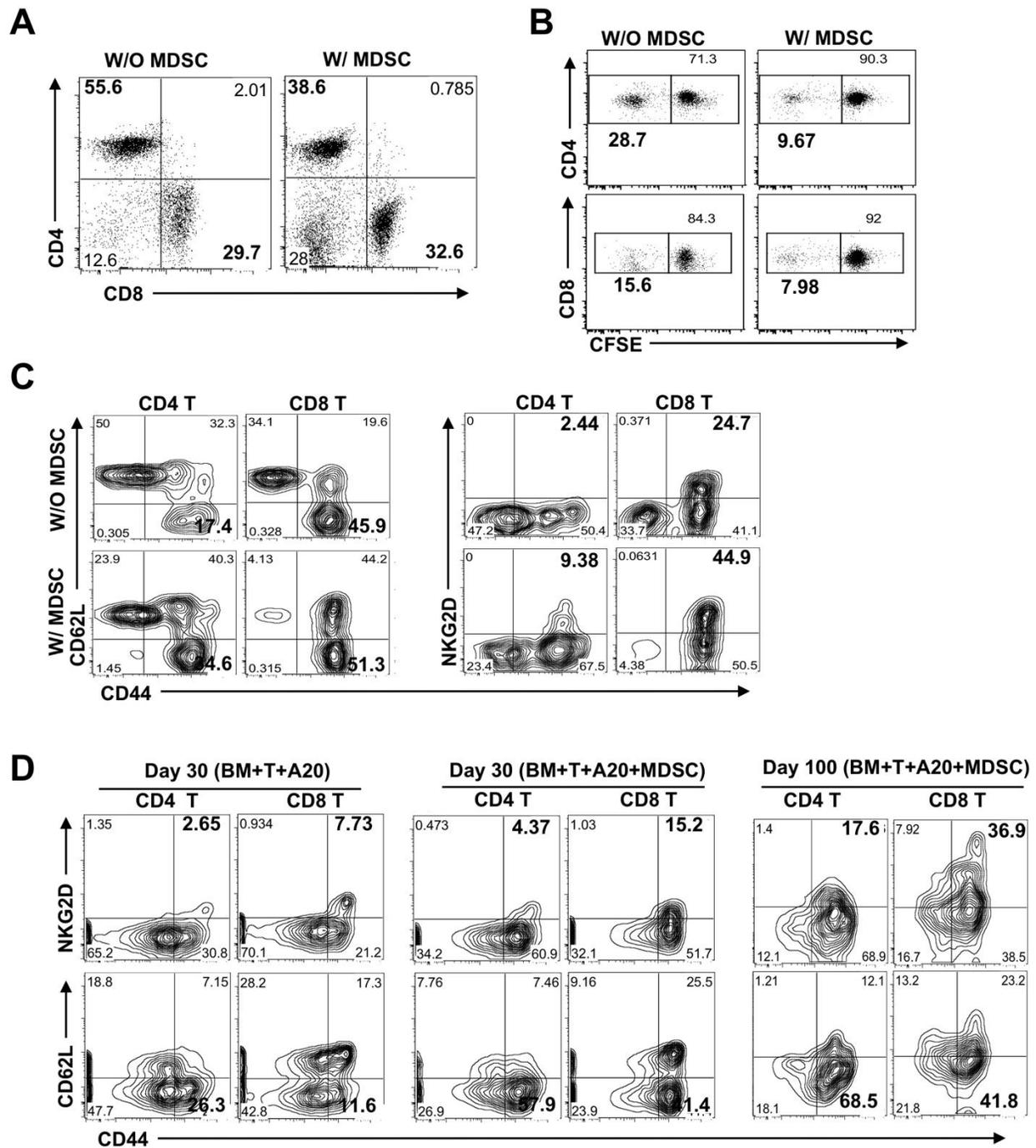


Figure S3. MDSCs inhibit allogenic T cell proliferation while up-regulate NKG2D expression.

(A) In vitro MLR assay of C57BL/6 splenocytes co-cultured with irradiated BALB/c splenocytes (1:1 ratio) in the presence of tumor-bearing-mouse-derived MDSC (splenocytes : MDSCs at 1:0.5 ratio). After 4 days co-culture, donor CD4 and CD8 T cell populations (gated on H2Kb⁺) were analyzed. (B) T cell proliferation in the MLR in the presence or absence of MDSCs as assessed by CFSE dilution. (C) Representative flow plots of CD62L, CD44 and NKG2D expression on T cells present in the MLR, and (D) on donor splenic T cells from mice received allo-HSCT with or without tumor-bearing-mouse-derived MDSC treatment.

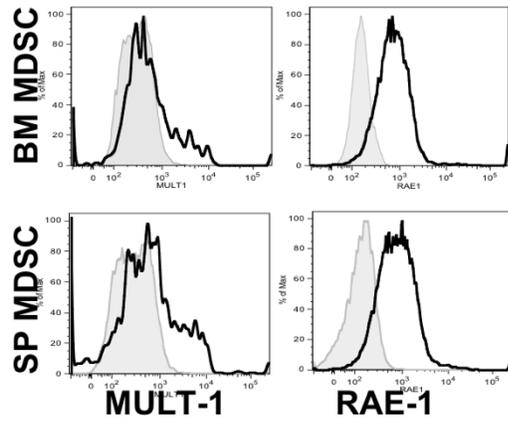


Figure S4. MDSCs express NKG2D ligands.

Flow analysis of expression of NKG2D ligands, RAE-1 and MULT-1 on CD115⁺ MDSCs in spleen and bone marrow from tumor-bearing mice.

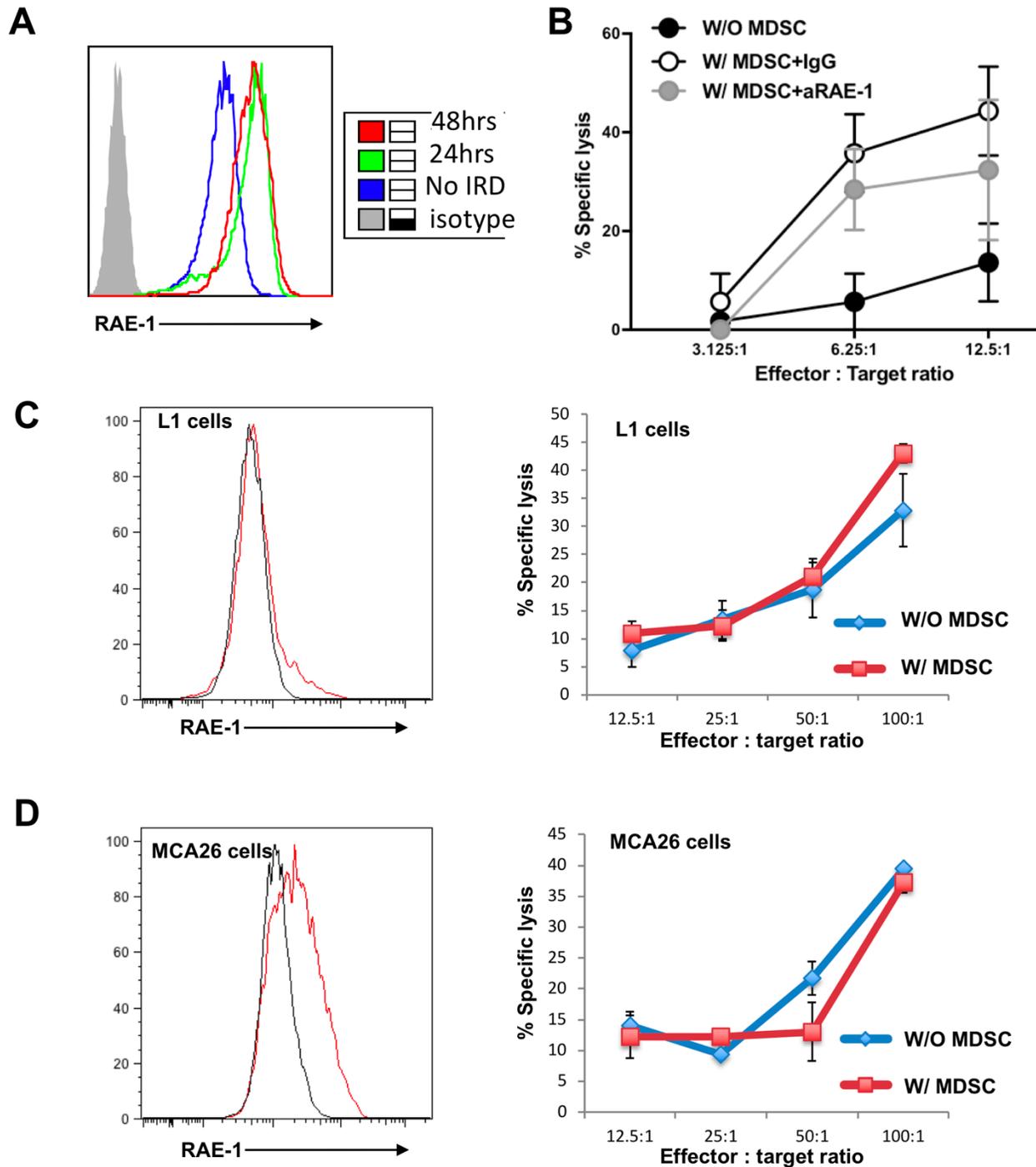


Figure S5. NKG2D ligand RAE-1 expression on tumor cells.

(A) Expression of RAE-1 on non-irradiated and irradiated (30Gy) A20 cells. Cells were analyzed at 24 and 48 hours after irradiation. (B) Cytotoxicity against A20 in presence or absence of anti-RAE-1 antibody. T cells were isolated from *in vitro* MLR in presence or absence of tumor-bearing-mouse-derived MDSCs. Data are presented as mean \pm S.D. ($n = 3$). (C and D) Cytotoxicity against RAE-1 low-expressing tumor cells, L1 and MCA-26. T cells were isolated from *in vitro* MLR in presence or absence of tumor-bearing-mouse-derived MDSCs. Data are presented as mean \pm S.D. ($n = 3$).

Table S1. Antibodies for flow cytometry and functional assays

Target	Cat. #	Company
CD4	563933	BDbiosciences
NKG2D	130212	Biolegend
NKG2D	115710	Biolegend
CD3	100314	Biolegend
CD115	135506	Biolegend
H2Kb	116506	Biolegend
CD90.2	12-0902-82	eBioscience
CD8	47-0081-82	eBioscience
CD44	12-0441-82	eBioscience
CD62L	45-0621-82	eBioscience
IFN- γ	17-7311-82	eBioscience
FoxP3	12-5773-82	eBioscience
CD25	17-0251-82	eBioscience
Granzyme B	12-8898-82	eBioscience
RAE-1	FAB17582P	R&D systems
RAE-1	AF1136	R&D systems
MULT-1	FAB2588R	R&D systems