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## V. 研究成果の刊行物・別刷

# Natural killer (NK)–dendritic cell interactions generate MHC class II-dressed NK cells that regulate CD4<sup>+</sup> T cells

Masafumi Nakayama<sup>a</sup>, Kazuyoshi Takeda<sup>b</sup>, Mitsuko Kawano<sup>a</sup>, Toshiyuki Takai<sup>c</sup>, Naoto Ishii<sup>d</sup>, and Kouetsu Ogasawara<sup>a,1</sup>

Departments of <sup>a</sup>Immunobiology and <sup>c</sup>Experimental Immunology, Institute of Development, Aging and Cancer, Tohoku University, Sendai 980-8575, Japan; <sup>b</sup>Department of Immunology, Juntendo University School of Medicine, Tokyo 113-8421, Japan; and <sup>d</sup>Department of Microbiology and Immunology, Tohoku University Graduate School of Medicine, Sendai 980-8575, Japan

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**Natural killer (NK) cells contribute to not only innate but also to adaptive immunity by interacting with dendritic cells (DCs) and T cells. All activated human NK cells express HLA-DR and can initiate MHCII-dependent CD4<sup>+</sup> T-cell proliferation; however, the expression of MHCII by mouse NK cells and its functional significance are controversial. In this study, we show that NK–DC interactions result in the emergence of MHCII-positive NK cells. Upon in vitro or in vivo activation, mouse conventional NK cells did not induce MHCII transcripts, but rapidly acquired MHCII protein from DCs. MHCII *H2-Ab1*-deficient NK cells turned I-A<sup>b</sup>-positive when adoptively transferred into wild-type mice or when cultured with WT splenic DCs. NK acquisition of MHCII was mediated by intercellular membrane transfer called “trocytosis,” but not upon DAP10/12- and MHCII-binding NK cell receptor signaling. MHCII-dressed NK cells concurrently acquired costimulatory molecules such as CD80 and CD86 from DCs; however, their expression did not reach functional levels. Therefore, MHCII-dressed NK cells inhibited DC-induced CD4<sup>+</sup> T-cell responses rather than activated CD4<sup>+</sup> T cells by competitive antigen presentation. In a mouse model for delayed-type hypersensitivity, adoptive transfer of MHCII-dressed NK cells attenuated footpad swelling. These results suggest that MHCII-dressed NK cells generated through NK–DC interactions regulate T cell-mediated immune responses.**

Natural killer (NK) cells have long been known to play important roles in innate immunity, but recently their contributions to adaptive immunity have also been reported (1). Following immunization or infection, NK cells migrate to draining lymphoid organs, where they interact with dendritic cells (DCs) and/or T cells (2). By production of cytokines such as IFN- $\gamma$  and TNF- $\alpha$  and direct cell–cell contact, NK cells activate DCs to induce T-cell proliferation and differentiation (3, 4). However, NK cells can also negatively regulate adaptive immune responses (2, 5). For example, in mouse models of autoimmune diseases such as rheumatoid arthritis and experimental autoimmune encephalomyelitis, depletion of NK cells by anti-NK1.1 mAb exacerbates these diseases (5). However, the molecular mechanisms underlying negative immune regulation by NK cells are poorly understood (5). Although conventional mouse NK cells do not express MHCII, subpopulations of activated mouse NK cells have been found to express MHCII (6–9), suggesting that NK cells may directly regulate CD4<sup>+</sup> T-cell responses. Of note, activated human NK cells express HLA-DR and can induce MHCII-dependent CD4<sup>+</sup> T-cell proliferation (6, 10–13).

MHCII molecules are crucial for the presentation of peptides processed from extracellular proteins to CD4<sup>+</sup> T cells, and they shape T-cell receptor repertoire development during T-cell maturation and lineage commitment. The constitutive expression of MHCII is restricted to professional antigen-presenting cells (APCs) such as DCs, macrophages, and B cells (14). In addition to professional APCs, basophils also express MHCII and play a crucial role as APCs (15). MHCII expression is transcrip-

tionally regulated by the class II transactivator (CIITA) in APCs, including basophils (14, 15). However, the regulation of MHCII expression in murine NK cells and the mechanism by which MHCII<sup>+</sup> NK cells are generated remain unclear.

A wide variety of immune cell lineages communicate with each other through direct cell–cell contact or cytokine production to establish appropriate immune responses. Several recent studies have shown that during direct cell–cell interactions, plasma membrane fragments of one cell are transferred to the opposite cell (16, 17). This phenomenon is currently called “trocytosis” (17, 18), and it may generate novel cell populations that result from the interaction between two different types of immune cells. Indeed, Wakim and Bevan have recently reported that DC–DC interactions generate a novel DC subset, called “cross-dressed” DCs, which acquire peptide–MHCII complexes from donor DCs to drive memory CD8<sup>+</sup> T-cell activation (19). Therefore, immune responses can be regulated not only by lineage-committed cell populations but also by cell populations generated independently of transcription, by trocytosis. Whether NK–DC interactions produce novel cell populations in a similar manner has not been explored. Therefore, we investigated whether MHCII-positive NK cells could be generated through interactions between mouse conventional NK cells and splenic DCs, and asked whether the resulting MHCII-positive NK cells could regulate CD4<sup>+</sup> T-cell responses.

## Results

**Activated NK Cells Express MHCII Protein, Although Not the Transcript, in Vivo.** MHCII expression has been observed on human NK cells prepared from mixed lymphocyte cultures or pathogen-infected organs, suggesting that activated NK cells induce MHCII (10, 11). Consistent with these reports, we observed MHCII I-A<sup>b</sup> expression on splenic NK1.1<sup>+</sup> cells in C57BL/6 mice injected i.v. with the double-stranded RNA synthetic analog polyI:C, but not in naive mice (Fig. 1A). These NK1.1<sup>+</sup> cells also expressed NKG2D and DX5 (Fig. 1A), indicating that these cells were conventional NK cells. Expression levels of MHCII on splenic B cells and T cells were not altered by polyI:C administration (Fig. S1). To confirm that I-A<sup>b</sup> NK1.1<sup>+</sup> cells were a single NK cell population and not merely conjugates of NK cells and I-A<sup>b</sup> cells such as DCs, we sort-purified these cells and analyzed them by

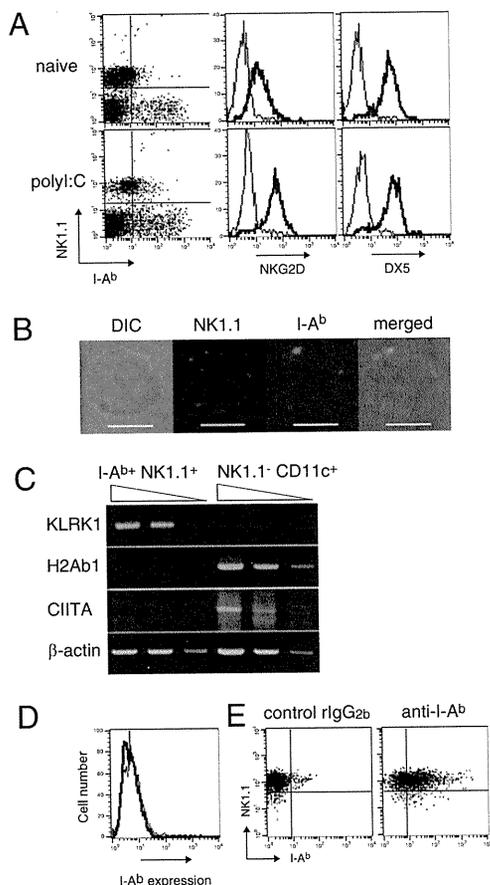
Author contributions: M.N. and K.O. designed research; M.N. performed research; T.T. and N.I. contributed new reagents/analytic tools; K.T., M.K., and K.O. analyzed data; and M.N., K.T., and K.O. wrote the paper.

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<sup>1</sup>To whom correspondence should be addressed. E-mail: ogasawara@idac.tohoku.ac.jp.

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**Fig. 1.** Activated NK cells express MHCII protein but not the transcript in vivo. (A) NK1.1 and I-A<sup>b</sup> expression on a non-T/B-cell population in spleen from naive or polyI:C (100 μg per mouse)-treated mice was analyzed (Left). Expression of NKG2D (Center) and DX5 (Right) on NK1.1<sup>+</sup> cells from splenocytes was analyzed using isotype control mAbs (thin lines) and specific mAbs (thick lines). (B) Purified I-Ab<sup>+</sup> DX5<sup>+</sup> cells were stained by AF488-anti-I-A<sup>b</sup> mAb and biotinylated anti-NK1.1 mAb, followed by Dylight 594-streptavidin. (Scale bars, 5 μm.) (C) Expression of the indicated transcripts in purified splenic I-Ab<sup>+</sup> NK1.1<sup>+</sup> cells or NK1.1<sup>-</sup> CD11c<sup>+</sup> cells was analyzed by semiquantitative RT-PCR using 10-fold serially diluted cDNA templates. (D) Splenic NK cells were purified and then cultured with IL-2 (1,000 U/ml) for 5 d, and then stained with control rat IgG<sub>2b</sub> (thin line) or anti-I-A<sup>b</sup> mAb (thick line). (E) NK cells cultured as described in D were labeled with CFSE and then transferred into mice. The following day, I-A<sup>b</sup> and NK1.1 expression level on CFSE<sup>+</sup> cells in spleen was analyzed. Similar results were obtained in three (A, D, and E) or two (B and C) independent experiments.

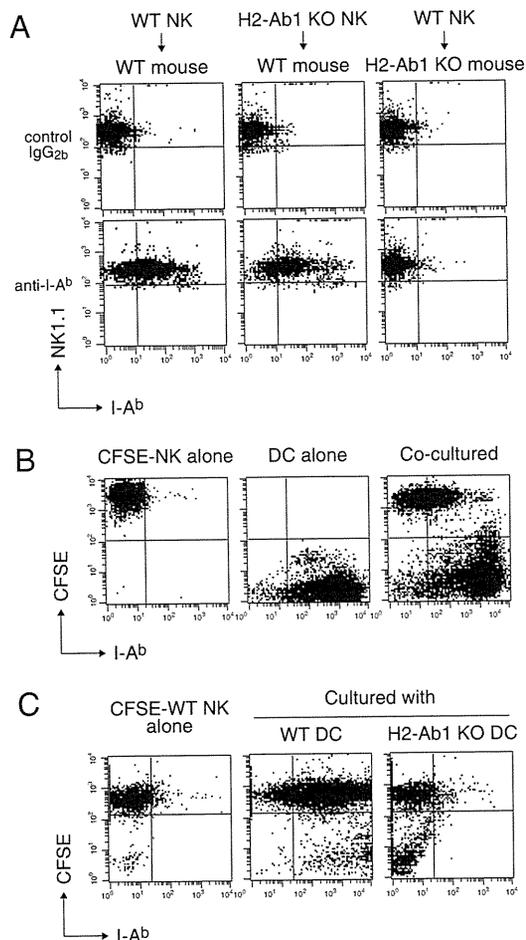
confocal microscopy. Fig. 1B shows that NK1.1<sup>+</sup> cells per se express I-A<sup>b</sup> on their cell surface.

To explore whether I-A<sup>b</sup> expression on NK cells depended on transcriptional regulation, we next performed semiquantitative PCR on mRNA from sort-purified NK cells. Unexpectedly, we observed neither MHCII *H2-Ab1* transcript nor the transactivator *CIITA* transcript in I-A<sup>b</sup><sup>+</sup> NK1.1<sup>+</sup> cells, whereas we did detect the *Klrk1* gene transcript that encodes NKG2D protein in these cells (Fig. 1C). Thus, it appeared that the expression of MHCII protein on NK cells occurred independently of transcriptional regulation in vivo.

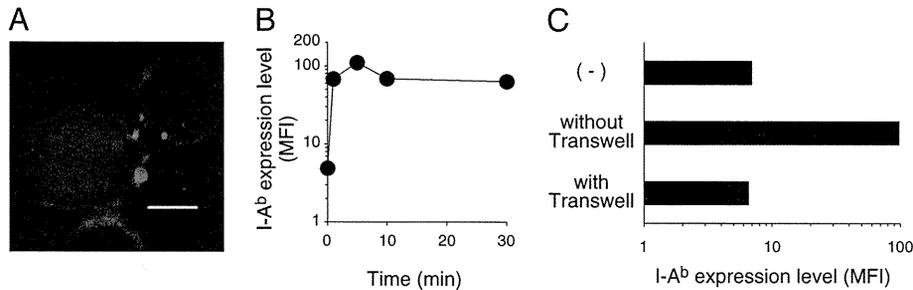
We next addressed whether activated NK cells express MHCII in vitro. We purified conventional NK cells from naïve mouse spleens and cultured them with IL-2 for 5 d. Unexpectedly, these conventional NK cells remained MHCII-negative in vitro (Fig. 1D). Interestingly, when we labeled the cells with 5- (and 6-) carboxyfluorescein diacetate succinimidyl ester (CFSE) and

transferred them into naïve mice, these NK cells turned MHCII-positive in the spleens of recipient mice (Fig. 1E). Nevertheless, we were unable to detect *H2-Ab1* or *CIITA* transcripts in transferred I-A<sup>b</sup><sup>+</sup> NK cells purified from the spleens of recipient mice (Fig. S2). Thus, our findings suggest that activated NK cells become MHCII-positive in vivo, and do so by transcription-independent mechanisms.

**Activated NK Cells Acquire MHCII from DCs Through NK-DC Interaction.** To further elucidate the pathway for MHCII<sup>+</sup> NK cell generation, we next adoptively transferred IL-2-activated MHCII *H2-Ab1* gene-deficient NK cells into wild-type (WT) mice. Surprisingly, we found that MHCII was expressed at high levels on *H2-Ab1*-deficient NK cells in WT mouse spleens within 1 d of transfer (Fig. 2A). In contrast, WT NK cells transferred into *H2-Ab1*-deficient mice remained MHCII-negative (Fig. 2A). Given that activated NK cells interact with DCs in vivo (2), we hypothesized that NK cells acquire MHCII from DCs. To address this possibility, we cocultured CFSE-labeled NK cells and splenic DCs at a 1:1 ratio. We observed that IL-2-activated NK cells turned MHCII-positive within 1 h of coculture (Fig. 2B). In contrast, IL-



**Fig. 2.** Activated NK cells acquire MHCII from DCs. (A) WT or *H2-Ab1*-deficient NK cells prepared as described in Fig. 1E were adoptively transferred into WT or *H2-Ab1*-deficient mice. The following day, NK1.1 and I-A<sup>b</sup> expression level on CFSE<sup>+</sup> cells in spleen was analyzed. (B) CFSE-labeled NK cells prepared as described in Fig. 1D were cocultured with splenic CD11c<sup>+</sup> cells at a 1:1 ratio for 1 h. I-A<sup>b</sup> expression on these cells was analyzed. (C) CFSE-labeled WT NK cells were cocultured with WT or *H2-Ab1*-deficient splenic DCs, and I-A<sup>b</sup> expression on these cells was analyzed as described in B. Similar results were obtained in three independent experiments.



**Fig. 3.** Intercellular MHCII transfer is mediated by trogocytosis. (A) TAMRA-labeled NK cells cocultured with DCs were stained with AF488-anti-I-A<sup>b</sup> mAb and analyzed by confocal microscopy. (Scale bar, 5  $\mu$ m.) (B) CFSE-labeled NK cells were cocultured with DCs as described in Fig. 2B for the indicated periods of time. The mean fluorescence intensity (MFI) of I-A<sup>b</sup> expression on NK cells was analyzed by flow cytometry. (C) CFSE-labeled NK cells were cocultured with DCs for 1 h together (no transwell) or separated by a semipermeable membrane (transwell). The MFI of I-A<sup>b</sup> expression on NK cells was analyzed by flow cytometry. Similar results were obtained in three (A and B) and two (C) independent experiments.

2-activated WT NK cells remained MHCII-negative after coculture with *H2-Ab1*-deficient DCs (Fig. 2C), providing evidence that NK cells acquire MHCII from WT DCs. The acquired MHCII protein remained on NK cells for at least 12 h after removal of DCs (Fig. S3A). Freshly isolated splenic NK cells did not acquire MHCII protein from DCs effectively (Fig. S3B), and thus NK cells gain this ability once activated.

Intercellular protein transfer between immune cells is mediated by several pathways, including membrane nanotubes (transient long-distance connections), trogocytosis (a rapid, cell contact-dependent transfer of membrane fragment), and exosomes (secreted membrane nanovesicles of  $\leq 100$  nm) (16–18, 20). By confocal microscopy, we could not observe nanotubes between NK cells and DCs, but we found I-A<sup>b</sup>-containing membrane fragments ( $\sim 1$   $\mu$ m) from DCs on NK cells (Fig. 3A). Although DCs have been reported to secrete MHCII-containing exosomes (20), we did not detect exosomes in culture supernatants until 12 h after incubation (Fig. S4). In contrast, MHCII-containing membrane fragment transfer occurred within minutes (Fig. 3B). Furthermore, MHCII acquisition was completely abrogated by culturing NK cells and DCs in transwell plates (Fig. 3C), indicating that acquisition was dependent upon cell–cell contact. These results are consistent with the transfer of MHCII being mediated through trogocytosis rather than exosome secretion.

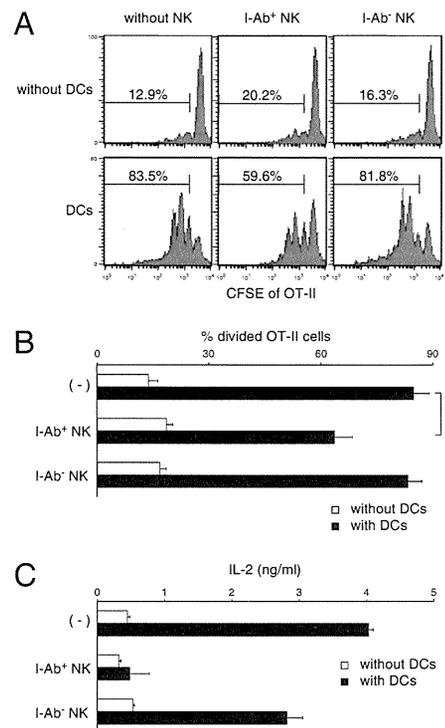
Trogocytosis is generally thought to be triggered by receptor signaling, and it has been reported that NK cells acquire MHC I and MHC I-related chain B from target cells by using NK cell receptors (NKR) (21–23). Therefore, we conducted further studies to determine whether NKR signaling is required for MHCII acquisition. However, DAP10/12-deficient NK cells as well as WT NK cells acquired MHCII from DCs (Fig. S5A). WT NK cells also acquired MHCII from  $\beta 2$ -microglobulin-deficient DCs that lacked cell surface expression of MHC I, indicating that MHC I-binding NKR are not involved in this process (Fig. S5A). Consistent with these results, MHCII acquisition was not inhibited by an inhibitor of Syk family kinase, PI3K, or Src kinase (Fig. S5B). Neither did we observe the involvement of NK effector molecules such as perforin, IFN- $\gamma$ , FasL, or TRAIL (24) in this process (Fig. S5). Interestingly, pretreatment of NK cells or DCs with cytochalasin D, an inhibitor of actin polymerization, or sodium azide, which depletes intracellular ATP, substantially inhibited MHCII transfer (Fig. S5B). Fixation of either NK cells or DCs with paraformaldehyde completely inhibited MHCII transfer (Fig. S5B). Collectively, these results suggest that activated NK cells acquire MHCII-containing DC membranes through a process that is dependent on plasma membrane and actin cytoskeleton interactions, but not NKR signaling or NK effector molecules.

#### MHCII-Dressed NK Cells Negatively Regulate CD4<sup>+</sup> T-Cell Proliferation.

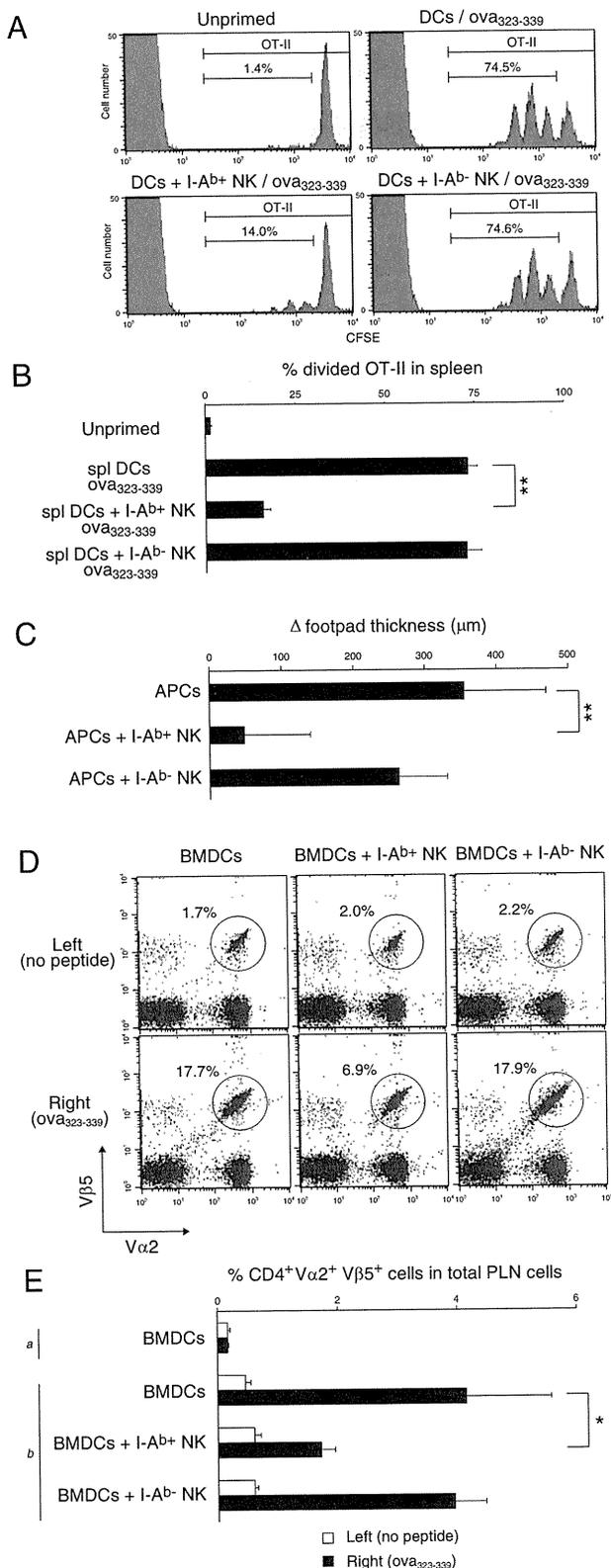
To explore whether MHCII-dressed NK cells could act as APCs

to stimulate naïve T cells, we examined whether NK cells could also acquire costimulatory molecules from DCs by an in vitro coculture assay. Activated NK cells acquired CD80 and CD86, but the levels of these molecules on NK cells were minimal (Fig. S6).

We next examined whether MHCII-dressed NK cells could influence CD4<sup>+</sup> T-cell responses. We performed an antigen-presentation study using OT-II CD4<sup>+</sup> T cells specific for OVA<sub>323–339</sub>/I-A<sup>b</sup>. After coculture with WT or *H2-Ab1*-deficient



**Fig. 4.** MHCII<sup>+</sup> NK cells suppress CD4<sup>+</sup> T-cell proliferation induced by DCs. (A) NK cells were purified by cell sorting after coculture with WT DCs (I-A<sup>b</sup>+ NK) or *H2-Ab1*-deficient DCs (I-A<sup>b</sup>- NK). CFSE-labeled OT-II CD4<sup>+</sup> T cells were cocultured with purified NK cells at a 1:10 ratio (NK:T), DCs at a 1:10 ratio (DC:T), or NK cells and DCs at a 1:1:10 ratio (NK:DC:T) in the presence of OVA<sub>323–329</sub> peptides (10 ng/mL) for 3 d. The CFSE intensity of OT-II CD4<sup>+</sup> cells was then analyzed. The percentages of divided OT-II cells are shown as means  $\pm$  SD of triplicates in B (\*\**P* < 0.01 compared with DCs alone). (C) Production of IL-2 in the culture supernatants at 48 h after coculture was measured by ELISA. Similar results were obtained in three independent experiments.



**Fig. 5.** MHCII<sup>+</sup> NK cells suppress CD4<sup>+</sup> T-cell responses in vivo. (A) B6 mice ( $n = 3$ ) adoptively transferred with CFSE-labeled OT-II CD4<sup>+</sup> T cells were i.v. injected with OVA<sub>323-339</sub>-loaded DCs or a mixture of these DCs and NK cells precultured with DCs together (I-A<sup>b+</sup> NK cells) or separated (I-A<sup>b-</sup> NK cells) as described in Fig. 3C. Two days later, CFSE intensity of CD4<sup>+</sup> Vα2<sup>+</sup> OT-II cells in spleen was analyzed. The percentage of divided OT-II cells is shown in B. B6 mice [ $n = 4$  (C) or  $n = 3$  (D and E)] adoptively transferred with activated OT-II CD4<sup>+</sup> T cells were s.c. injected with OVA<sub>323-339</sub>-loaded APCs or a mixture of

DCs, I-A<sup>b</sup>-dressed (I-A<sup>b+</sup>) or I-A<sup>b</sup>-negative (I-A<sup>b-</sup>) NK cells were sort-purified. Purified I-A<sup>b+</sup> NK cells did not induce naive OT-II CD4<sup>+</sup> T-cell proliferation (Fig. 4A and B), possibly due to insufficient expression of costimulatory molecules (Fig. S6). Interestingly, purified I-A<sup>b+</sup> NK cells suppressed OT-II CD4<sup>+</sup> T-cell proliferation induced by DCs, whereas I-A<sup>b-</sup> NK cells cocultured with *H2-Ab1*-deficient DCs did not (Fig. 4A and B). Moreover, I-A<sup>b+</sup> NK cells suppressed IL-2 production from OT-II CD4<sup>+</sup> T cells more effectively than I-A<sup>b-</sup> NK cells (Fig. 4C). These results indicate that MHCII-dressed NK cells regulate CD4<sup>+</sup> T-cell responses to DCs via antigen presentation on MHCII without sufficient costimulation.

Furthermore, we addressed whether I-A<sup>b+</sup> NK cells could suppress CD4<sup>+</sup> T-cell responses in vivo. Interestingly, I-A<sup>b+</sup> NK cells significantly suppressed OT-II CD4<sup>+</sup> T-cell proliferation induced by DCs loaded with OVA<sub>323-339</sub> peptides in spleen, although I-A<sup>b-</sup> NK cells did not affect the proliferation (Fig. 5A and B). In a mouse model for delayed-type hypersensitivity (DTH) where activated OT-II CD4<sup>+</sup> T cells were i.v. transferred followed by s.c. injection with OVA<sub>323-339</sub>-loaded APCs, I-A<sup>b+</sup> NK cells attenuated footpad swelling (Fig. 5C) and reduced OT-II CD4<sup>+</sup> T-cell accumulation in the draining popliteal lymph nodes (PLNs) (Fig. 5D and E). Taken together, these results suggest that MHCII-dressed NK cells suppress CD4<sup>+</sup> T-cell responses in vivo.

## Discussion

In this study, we provide evidence that conventional murine NK cells do not transcriptionally induce MHCII but instead rapidly acquire MHCII protein from DCs through NK-DC interactions. These MHCII-dressed NK cells suppress CD4<sup>+</sup> T-cell responses to DCs by presenting antigen-MHCII complexes without sufficient costimulation, which might induce anergy in CD4<sup>+</sup> T cells. In addition, adoptive transfer of MHCII-dressed NK cells attenuated dermal DTH. Therefore, our findings may provide a mechanistic explanation for the negative immune regulation of T-cell immunity by NK cells.

Several recent studies have identified NK/DC hybrid-phenotype cells, which have functional properties characteristic of both NK cells and DCs (11, 25–28). IFN-producing killer DCs (IKDCs), also called B220<sup>+</sup> NK1.1<sup>+</sup> DCs, were identified as a novel DC subset harboring killer activity (27, 28). On the contrary, more recent studies have proposed that these killer DCs are functionally and developmentally activated NK cells (7–9). It remains unclear whether the MHCII-dressed NK cells we describe here are identical to the NK/DC hybrid-phenotype cells described in previous studies (25–28), although at least IKDCs and MHCII-dressed NK cells have similar antigenic phenotypes: CD11c<sup>+</sup> B220<sup>+</sup> MHCII<sup>+</sup> NKG2D<sup>+</sup> CD86<sup>dull</sup> Gr1<sup>-</sup> (Fig. S7). Interestingly, in a mouse model of type 1 diabetes, CD11c<sup>+</sup> DX5<sup>+</sup> cells, which are functionally and phenotypically similar to MHCII-dressed NK cells, were found to negatively regulate pathogenic T-cell activation (25). Of note, we observed that a small population of DCs cocultured with IL-2-activated NK cells became Ly49G2-positive (Fig. S8), suggesting that DCs

these APCs and I-A<sup>b+</sup> NK or I-A<sup>b-</sup> NK cells into the right footpads. The left footpads were injected with the same cell population without OVA<sub>323-339</sub> as controls. The following day, Δfootpad thickness was calculated by subtracting the left hind footpad thickness from the right (C). Three days after the s.c. injection, the percentage of Vα2<sup>+</sup> Vβ5<sup>+</sup> cells in total CD4<sup>+</sup> T cells in PLNs was analyzed (D) and the percentage of CD4<sup>+</sup> Vα2<sup>+</sup> Vβ5<sup>+</sup> cells in total PLN cells is calculated in E. (a) Naive mice. (b) OT-II CD4<sup>+</sup> T-cell transferred mice. \* $P < 0.05$ , \*\* $P < 0.01$  compared with APCs alone. Similar results were obtained in two (A and B) or three (C and D) independent experiments. BMDCs, bone marrow-derived DCs.

could also acquire NK cell surface proteins. Although this observation might also account for the generation of NK/DC hybrid-phenotype cells described in previous studies (25, 26), further studies will be necessary to characterize these hybrid-phenotype cells.

In addition, we observed that activated murine NK cells acquired MHCII from B cells in coculture experiments. However, the acquisition level of MHCII on these NK cells was lower than that on NK cells cocultured with DCs (Fig. S9), suggesting that activated NK cells preferentially acquired MHCII from DCs.

In contrast to mouse NK cells, after activation, all human NK cells synthesize HLA-DR as well as costimulatory molecules including CD80, CD86, and OX40 ligand (6, 11, 12). Unlike in humans, activation of mouse NK cells apparently does not induce the endogenous expression of MHCII.

Given that many cell types store a large excess of membrane on their cell surface (16), intercellular membrane transfer might occur frequently during immune cell-cell interactions. Recently, DCs have been reported to acquire peptide-MHCI complexes from distinct donor DCs and subsequently drive memory CD8<sup>+</sup> T-cell activation (19). T cells have also been reported to acquire CD80 and CD86 proteins from DCs by CTLA-4, thereby downmodulating the delivery of costimulatory signals (29). Our findings show that activated NK cells can acquire MHCII from DCs and regulate T-cell immune responses *in vitro* and *in vivo*. Taken together, it is possible that immune cells acquire additional functions and/or alter their intrinsic functions through intercellular transfer of immune regulatory molecules such as MHCII in lymphoid organs. Such newly generated cell populations could play important roles in the regulation of immune responses through their effects on other cell types.

## Materials and Methods

Further details are available in *SI Materials and Methods*.

**Mice.** C57BL/6 mice were obtained from CLEA Japan. MHCII *H2-Ab1*-deficient mice (30) were kindly provided by D. Mathis (Harvard Medical School, Boston, MA). OT-II transgenic/*rag-1* knockout mice were obtained from Taconic. These mice were maintained under specific pathogen-free conditions, and used according to the guidelines of the Institutional Animal Care and Use Committee established at Tohoku University.

**RT-PCR.** Total RNA was purified from cells using Sepasol (Nacalai). Complementary DNAs were synthesized from total RNAs by using oligo(dT) primer (Invitrogen). PCR was performed by using 10-fold serially diluted cDNA templates, AmpliTaq poly (Applied Biosystems), and primers listed in *SI Materials and Methods*.

**NK-DC Interaction.** Mouse splenic NK cells and DCs were prepared as described previously (31, 32). CFSE (0.5  $\mu$ M)-labeled IL-2 (1,000 U/mL)-activated NK cells ( $5 \times 10^5$  per well) and splenic DCs ( $5 \times 10^5$  per well) were cocultured in a 96-well flat-bottom plate for the indicated periods at 37 °C. Then cells were stained with APC-labeled anti-I-A<sup>b</sup> mAb (BioLegend) and analyzed on a FACSCanto II (BD Biosciences).

**Confocal Microscopy.** Cells were stained with 5- (and 6-) carboxyethylrhodamine succinimidyl ester (TAMRA; Invitrogen) or the following mAbs: AF488-anti-I-A<sup>b</sup>, biotinylated anti-NK1.1 mAbs, and streptavidin-DyLight 594 (BioLegend). Then these cells were analyzed on a Carl Zeiss LSM510 confocal laser-scanning microscope equipped with a 63 $\times$  objective lens as described previously (33).

**In Vitro Antigen Presentation Assay.** The antigen presentation assay was performed as described previously (32) with some modifications. Bone marrow-derived DCs ( $5 \times 10^3$  per well) and/or sort-purified NK cells ( $5 \times 10^3$  per well) were cocultured with CFSE (10  $\mu$ M)-labeled OT-II CD4<sup>+</sup> T cells ( $5 \times 10^4$  per well) in a 96-well U-bottom plate for 3 d in the presence of OVA<sub>323-339</sub> peptides (10 ng/mL; Abgent). CFSE fluorescence intensity of OT-II CD4<sup>+</sup> T cells was analyzed by flow cytometry. Production of IL-2 in the culture supernatant at 48 h postaddition of OT-II CD4<sup>+</sup> T cells was measured by ELISA (eBioscience).

**In Vivo OT-II Proliferation Assay.** CFSE-labeled OT-II CD4<sup>+</sup> T cells ( $3 \times 10^6$  per mouse) were adoptively transferred into B6 mice. The following day, mice were *i.v.* injected with OVA<sub>323-339</sub> (1  $\mu$ g/mL)-loaded splenic DCs ( $3 \times 10^6$  per mouse) or a mixture of these DCs ( $3 \times 10^6$  per mouse) and NK cells ( $3 \times 10^6$  per mouse). Two days later, mice were killed, and CFSE dilution of CD4<sup>+</sup> V $\alpha$ 2<sup>+</sup> splenocytes was analyzed by flow cytometry.

**Dermal DTH.** B6 mice adoptively transferred with activated OT-II CD4<sup>+</sup> T cells ( $6 \times 10^6$  per mouse) were *s.c.* injected with 50  $\mu$ L of OVA<sub>323-339</sub>-loaded APCs ( $2 \times 10^7$  per mouse) or a mixture of these APCs ( $2 \times 10^7$  per mouse) and NK cells ( $1 \times 10^7$  per mouse) into the right footpad. The left footpads were injected with the same cell population without OVA<sub>323-339</sub> as controls. Footpad thickness and infiltration of OT-II cells into PLNs were analyzed.

**ACKNOWLEDGMENTS.** We thank Hiromi Yoshida for cell sorting; Shota Endo and Hisaya Akiba for advice on OT-II experiments; and Chika Takahashi, Misato Tsugita, and Kazusa Ishizaki for technical assistance. This work was supported by Grants-in-Aid for Scientific Research from the Japanese Ministry of Education, Culture, Sports, Science and Technology (to M.N., K.T., and K.O.); by Grants-in-Aid for Scientific Research from the Ministry of Health, Labour and Welfare of Japan H22-meneki-ippan-004 (to K.O.), and H22-meneki-ippan-005 (to M.N.); by a grant from the Takeda Science Foundation (to M.N.); and by a grant from the Mochia Memorial Foundation for Medical and Pharmaceutical Research (to M.N.).

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# Supporting Information

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## SI Materials and Methods

**Mice.** C57BL/6 mice were obtained from CLEA Japan. MHCII *H2-Ab1*-deficient mice (1) were kindly provided by D. Mathis (Harvard Medical School, Boston, MA). C57BL/6 mice deficient in *perforin*, *IFN- $\gamma$* , or  *$\beta$ 2-microglobulin* were obtained from The Jackson Laboratory. OT-II transgenic/*rag-1* knockout mice were obtained from Taconic. DAP12 knockout mice and DAP10/12 double knockout mice were generated as described previously (2).

**Adoptive Natural Killer Cell Transfer Assay.** Natural killer (NK) cells were prepared from WT B6 mice or *H2-Ab1* knockout mice as described previously (3). In brief, spleen cells were incubated with anti-CD4 and anti-CD8 mAbs, followed by magnetic beads coated with goat anti-mouse Ig and goat anti-rat Ig Abs (Advanced Magnetics). After removing CD4, CD8, and surface Ig-positive cells by magnetic cell sorting, NK cells were stained with phycoerythrin (PE)-labeled DX5 mAb (BioLegend) and purified using anti-PE MACS beads (Miltenyi Biotec). Purified NK cells were cultured with rhIL-2 (1,000 U/mL; Wako) for 5 d. These activated NK cells ( $5 \times 10^6$  per mouse) were labeled with 5- (and 6-) carboxyfluorescein diacetate succinimidyl ester (CFSE) (2  $\mu$ M; Invitrogen) and transferred i.v. into WT B6 mice or *H2-Ab1* knockout mice. The following day, mice were killed, and splenocytes were stained with PE-anti-NK1.1 and antigen-presenting cell (APC)-anti-I-A<sup>b</sup> mAbs (BioLegend). Expression of I-A<sup>b</sup> and NK1.1 on CFSE<sup>+</sup> cells was analyzed on a FACSCanto II (BD Biosciences).

**RT-PCR.** Splenic NK1.1<sup>+</sup> cells were purified from polyI:C (100  $\mu$ g per mouse; InvivoGen)-treated mice or naive mice as described above. Splenic CD11c<sup>+</sup> dendritic cells (DCs) were purified as described previously (4). In brief, spleens were digested with 400 U/mL collagenase (Wako) in the presence of 5 mM EDTA and separated into low- and high-density fractions on an Opti-Prep gradient (Axis-Shield). Low-density cells were purified using anti-CD11c MACS beads (Miltenyi Biotec). Then I-Ab<sup>+</sup> NK1.1<sup>+</sup> cells or NK1.1<sup>-</sup> CD11c<sup>+</sup> cells were further purified on an Epics ALTRA cell sorter (Beckman Coulter) (>98% I-Ab<sup>+</sup> NK1.1<sup>+</sup> cells; >96% NK1.1<sup>-</sup> CD11c<sup>+</sup> cells). Total RNA was purified from these purified cells using Sepasol (Nacalai). Complementary DNAs were synthesized from total RNAs by using oligo(dT) primer (Invitrogen). PCR was performed by using 10- or 100-fold serially diluted cDNA templates and AmpliTaq poly (Applied Biosystems). Primers used in this study were as follows: Klrk1-F, 5'-ATG GCA TTG ATT CGT GAT CGA AAG-3'; Klrk1-R, 5'-TTA CAC CGC CCT TTT CAT GCA GAT-3'; H2-Ab1-F, 5'-CGC ATA CGA TAT GTG ACC AGA TAC-3'; H2-Ab1-R, 5'-CAA CAT CAT TTT GCT CCA GGC AGA CTC-3'; class II transactivator (Ciita)-F, 5'-ATG CGC TGC CTG GTT CCT GGC CCT TCT-3'; Ciita-R, 5'-TCC CTG GGG CAG AGT GGT GAA GAT-3'; Actb-F, 5'-ATG GAT GAC GAT ATC GCT GCG CTG-3'; Actb-R, 5'-GTA GCC ACG CTC GGT CAG GAT CTT-3'.

**ELISA for DC-Derived Exosomes.** Anti-mouse CD9 mAb (10  $\mu$ g/mL; BioLegend) was precoated on a 96-well EIA/RIA plate (Costar). After blocking with 5% FCS/PBS at room temperature for 1 h, the culture supernatants were added to the wells and incubated at room temperature for 2 h. After washing with PBS containing 0.05% Tween 20, exosomes were detected by biotinylated anti-I-A<sup>b</sup> mAb (BioLegend) and streptavidin-HRP (eBioscience).

**NK-DC Interaction.** CFSE (0.5  $\mu$ M)-labeled IL-2 (1,000 U/mL)-activated NK cells ( $5 \times 10^5$  per well) and splenic DCs ( $5 \times 10^5$  per well) were cocultured in a 96-well flat-bottom plate for the indicated periods at 37 °C. For the transwell assay, NK cells ( $3 \times 10^6$  per well) were cultured in the upper chamber of a 12-well plate and separated from DCs ( $3 \times 10^6$  per well) by a 0.4  $\mu$ m-pore semipermeable membrane (Greiner Bio-One). As controls, NK cells and DCs ( $3 \times 10^6$  per well) were cultured together in a 12-well plate. For the inhibition assay, NK cells or DCs were pretreated for 1 h with the following reagents: piceatannol (100  $\mu$ M), LY294002 (100  $\mu$ M), cytochalasin D (20  $\mu$ M), PP2 (50  $\mu$ M), paraformaldehyde (4%; Wako), polyI (10  $\mu$ g/mL), dextran sulfate (10  $\mu$ g/mL; Sigma), RGDS peptides (10 mM; Peptide Institute), NaN<sub>3</sub> (50 mM; Nacalai), and mAbs against Fc $\gamma$ RIIb/III (2.4G2, 10  $\mu$ g/mL; BD Biosciences), FasL (K10, 10  $\mu$ g/mL), or TRAIL (N2B2, 10  $\mu$ g/mL; BioLegend). After coculture for the indicated times, cells pretreated with 2.4G2 mAb were stained with the following mAbs: APC-labeled anti-I-A<sup>b</sup>, anti-CD80, anti-CD86, or anti-CD11c (BioLegend). In some assays, NK cells were then purified by cell sorting and analyzed via flow cytometry.

**In Vitro Antigen Presentation Assay.** The antigen presentation assay was performed as described previously (4) with some modifications. Bone marrow-derived DCs (BMDCs) were generated by rmGM-CSF (40 ng/mL; PeproTech). CFSE-labeled NK cells ( $1 \times 10^7$  per well) and BMDCs ( $5 \times 10^6$  per well) were cocultured in a 6-well plate for 1 h. Then these cells were stained with APC-anti-I-A<sup>b</sup> mAb, followed by sorting into I-A<sup>b+</sup> and I-A<sup>b-</sup> NK subsets on an Epics ALTRA cell sorter (Beckman Coulter) (>98% I-A<sup>b+</sup> CFSE<sup>+</sup>; >99% I-A<sup>b-</sup> CFSE<sup>+</sup>). BMDCs ( $5 \times 10^3$  per well), and/or sorted NK cells ( $5 \times 10^3$  per well) were cocultured with CFSE (10  $\mu$ M)-labeled OT-II CD4<sup>+</sup> T cells ( $5 \times 10^4$  per well) in a 96-well U-bottom plate for 3 d in the presence of OVA<sub>323-339</sub> peptides (10 ng/mL; Abgent). CFSE fluorescence intensity of OT-II CD4<sup>+</sup> T cells was analyzed by flow cytometry. Production of IL-2 in the culture supernatant at 48 h post-addition of OT-II CD4<sup>+</sup> T cells was measured by ELISA (eBioscience). For statistical evaluation, an unpaired Student's *t* test, two-tailed, was used. *P* values less than 0.05 were considered significant.

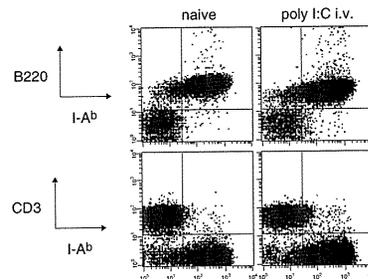
**In Vivo OT-II CD4<sup>+</sup> Cell Proliferation Assay.** OT-II CD4<sup>+</sup> cell proliferation in spleen was analyzed as described previously (4) with some modifications. CFSE-labeled naive OT-II CD4<sup>+</sup> T cells ( $3 \times 10^6$  per mouse) were adoptively transferred into B6 mice. The next day, mice were i.v. injected with OVA<sub>323-339</sub> (1  $\mu$ g/mL)-loaded splenic DCs ( $3 \times 10^6$  per mouse) or a mixture of these DCs ( $3 \times 10^6$  per mouse) and NK cells cocultured with these DCs together (I-A<sup>b+</sup> NK cells;  $3 \times 10^6$  per mouse) or separately (I-A<sup>b-</sup> NK cells;  $3 \times 10^6$  per mouse). Two days later, mice were killed, and CFSE dilution of CD4<sup>+</sup> V $\alpha$ 2<sup>+</sup> splenocytes was analyzed by flow cytometry. For statistical evaluation, an unpaired Student's *t* test, two-tailed, was used. *P* values less than 0.05 were considered significant.

**Dermal Delayed-Type Hypersensitivity.** A dermal delayed-type hypersensitivity study was performed as described previously (5) with some modifications. OT-II CD4<sup>+</sup> T cells were stimulated with OVA<sub>323-339</sub> (100 ng/mL) for 3 d, and then cells were maintained in IL-2 (10 U/mL)-containing medium without OVA<sub>323-339</sub> for another 3 d. These cells ( $6 \times 10^6$  per mouse) were transferred i.v. into B6 mice. The next day, recipient mice

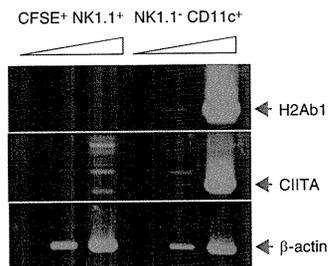
were s.c. injected with 50  $\mu$ L of OVA<sub>323–339</sub>-loaded APCs (collagenase-digested whole splenocytes or BMDCs;  $2 \times 10^7$  each per mouse) or a mixture of these APCs ( $2 \times 10^7$  per mouse) and NK cells cocultured with these DCs together (I-A<sup>b+</sup> NK cells;  $1 \times 10^7$  per mouse) or separately (I-A<sup>b-</sup> NK cells;  $1 \times 10^7$  per mouse) into the right footpad. The left footpads were injected with the same cell population without OVA<sub>323–339</sub> as controls.

The following day,  $\Delta$ footpad thickness was calculated by subtracting the left hind footpad thickness from the right. Three days after the s.c. injection, the OT-II CD4<sup>+</sup> V $\alpha$ 2<sup>+</sup> V $\beta$ 5<sup>+</sup> cell population migrating into popliteal lymph nodes was measured by flow cytometry. For statistical evaluation, an unpaired Student's *t* test, two-tailed, was used. *P* values less than 0.05 were considered significant.

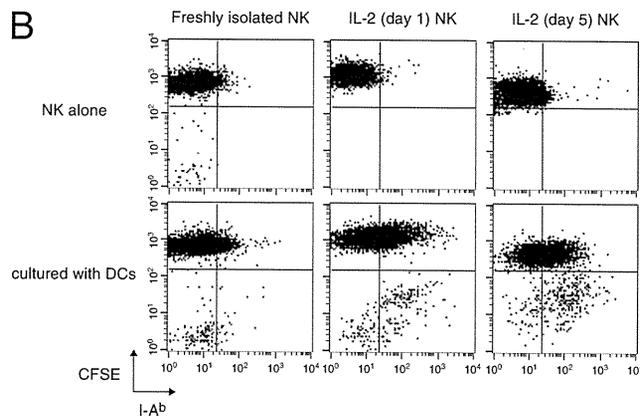
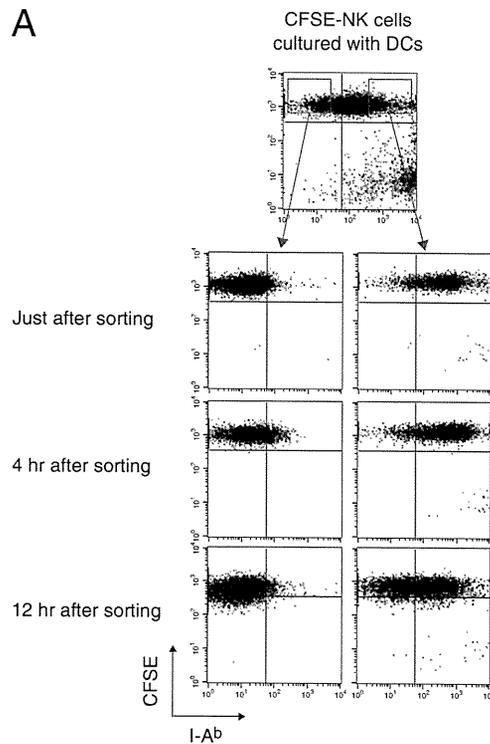
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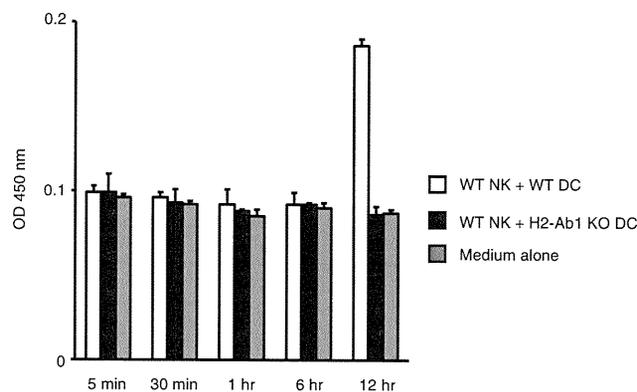
**Fig. S1.** Expression of MHCII on B and T cells. C57BL/6 mice were i.v. injected with 100  $\mu$ g of polyI:C. The following day, collagenase-digested splenocytes from these mice or naive C57BL/6 mice were stained with anti-CD3, anti-B220, and anti-I-A<sup>b</sup> mAbs. Similar results were obtained in three independent experiments.



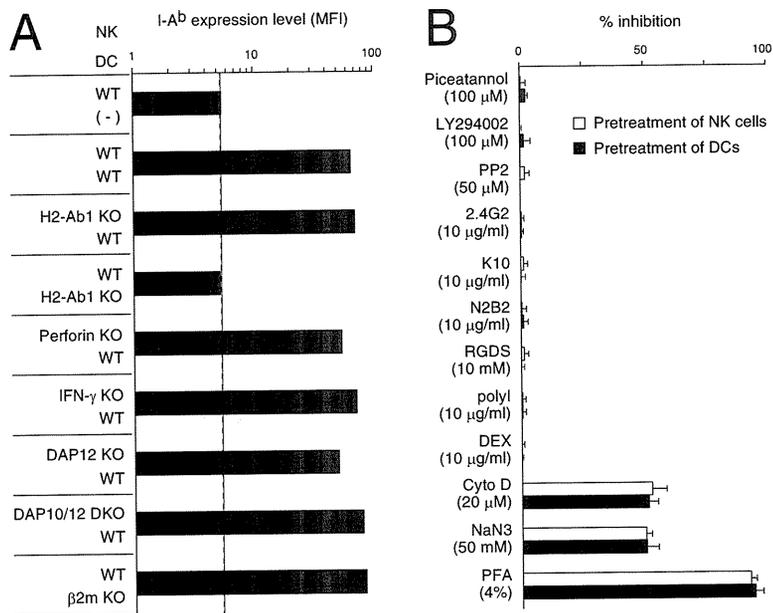
**Fig. S2.** Expression of H2-Ab1 and CIITA transcripts in purified CFSE<sup>+</sup> I-A<sup>b</sup><sup>+</sup> NK1.1<sup>+</sup> cells or NK1.1<sup>-</sup> CD11c<sup>+</sup> cells. IL-2-activated CFSE-labeled NK cells ( $5 \times 10^6$  per mouse) were transferred into mice. The following day, CFSE<sup>+</sup> NK1.1<sup>+</sup> cells in spleen were purified by cell sorting. CD11c<sup>+</sup> cells were purified as described in *S1 Materials and Methods*. Expression of the indicated transcripts in purified cells was analyzed by semiquantitative RT-PCR using 100-fold serially diluted cDNA templates. Similar results were obtained in two independent experiments.



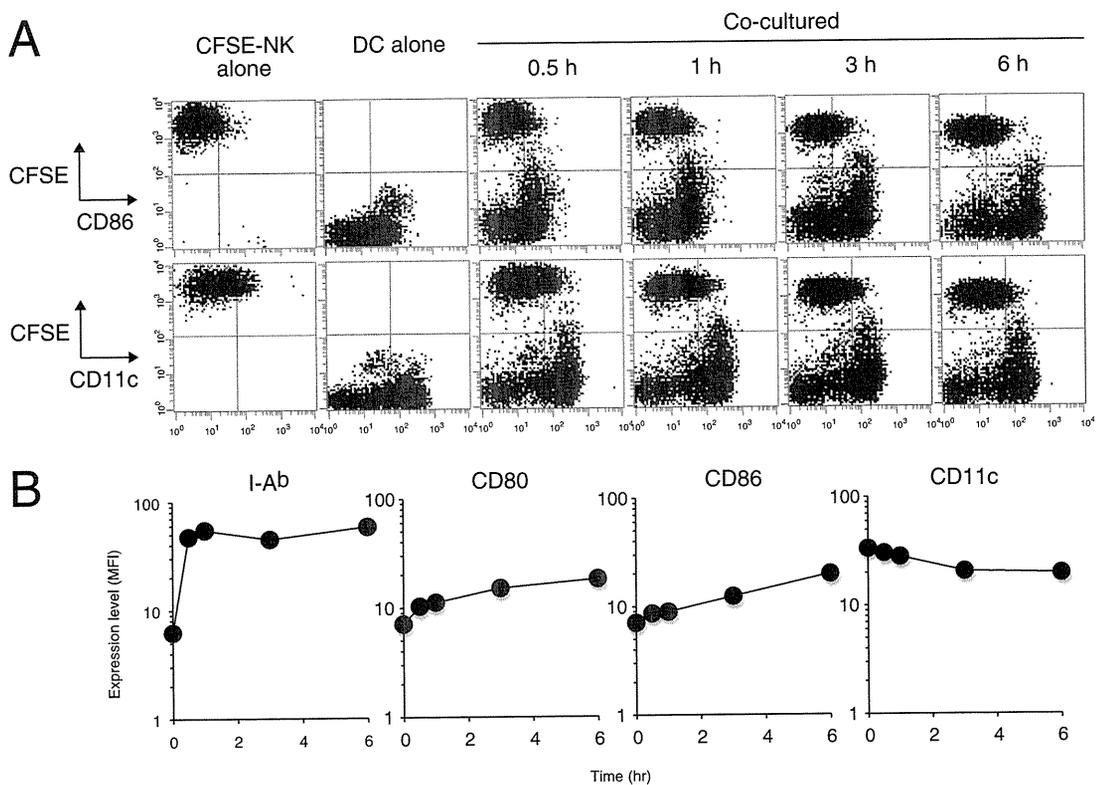
**Fig. S3.** Activated NK cells acquire MHCII. (A) Stability of MHCII on NK cells. I-Ab<sup>-</sup> and I-Ab<sup>+</sup> NK cells were purified by cell sorting after coculture with WT DCs. After the indicated times, cells were stained with anti-I-A<sup>b</sup> mAb. Similar results were obtained in three independent experiments. (B) Freshly isolated NK cells do not acquire MHCII. Freshly isolated splenic NK cells or IL-2 (1,000 U/mL)-activated NK cells were labeled with CFSE and then cocultured with splenic CD11c<sup>+</sup> cells at a 1:1 ratio for 1 h. I-A<sup>b</sup> expression on NK1.1<sup>+</sup> cells was analyzed. Similar results were obtained in two independent experiments.



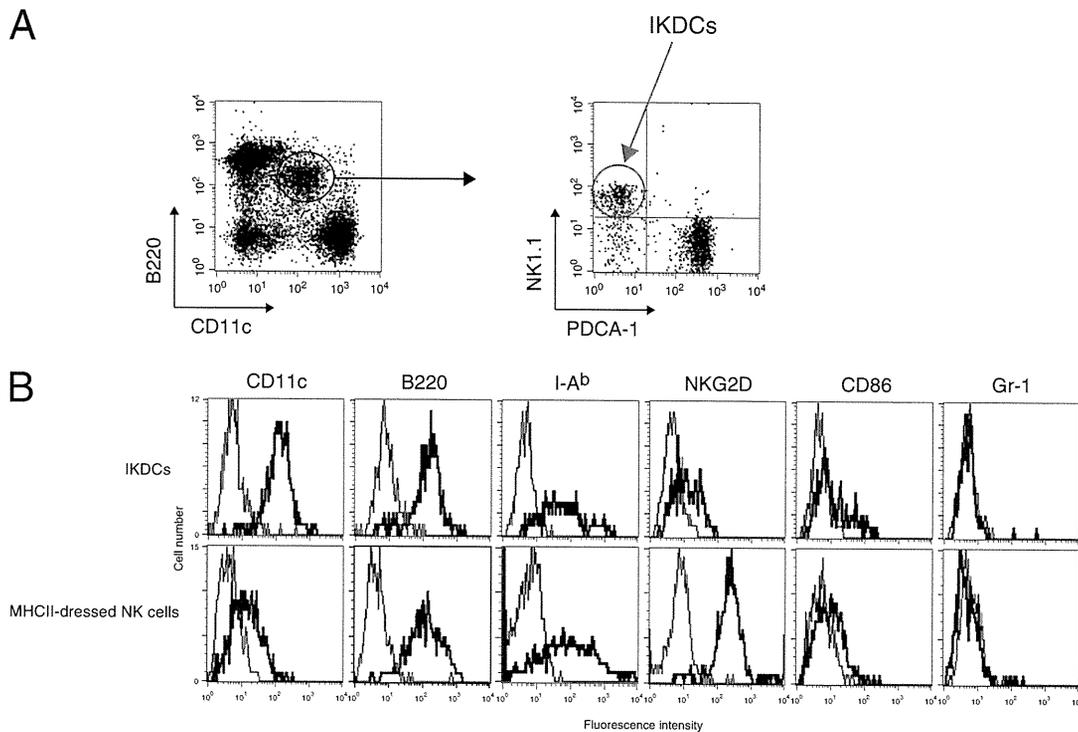
**Fig. S4.** Exosomes in NK/DC coculture supernatants. IL-2-activated WT NK cells were cocultured with WT or H2-Ab1 knockout splenic DCs at a 1:1 ratio for the indicated times. Exosomes in culture supernatants were detected by sandwich ELISA. Similar results were obtained in two independent experiments.



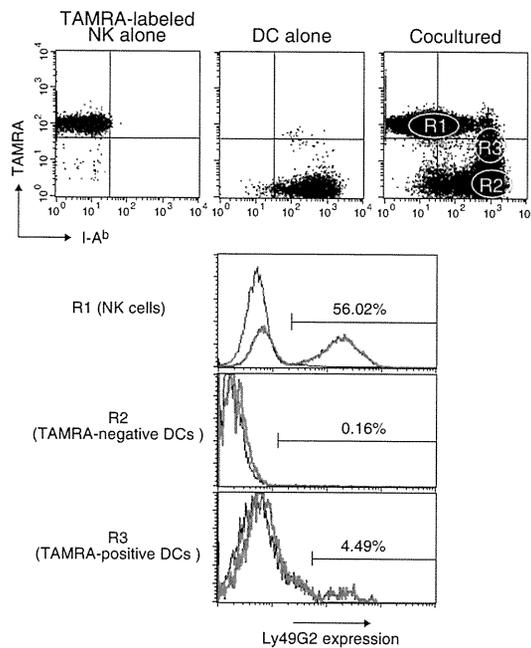
**Fig. 55.** Intercellular MHCII transfer is NK cell receptor-independent. (A) CFSE-labeled NK cells and splenic DCs from the indicated mice were cocultured for 30 min. Surface expression of I-A<sup>b</sup> on NK cells was analyzed by flow cytometry, and the mean fluorescence intensity (MFI) is indicated. (B) NK cells (white columns) or DCs (black columns) were pretreated with the following inhibitors 1 h before coculture: piceatannol (a Syk family kinase inhibitor), LY294002 (a specific inhibitor of PI3K), PP2 (a Src kinase inhibitor), 2.4G2 (an anti-FcγRIIb/III–blocking mAb), K10 (an anti-FasL–blocking mAb), N2B2 (an anti-TRAIL–blocking mAb), RGDS (an integrin-antagonizing peptide), polyI (polyinosinic acid; a negatively charged reagent), DEX (dextran sulfate; a negatively charged reagent), cytochalasin D (an inhibitor of actin polymerization), NaN<sub>3</sub> (an ATP-depleting reagent), and PFA (paraformaldehyde; a cross-linking reagent). The expression level of I-A<sup>b</sup> on NK cells was analyzed by flow cytometry. Percent inhibition was calculated by comparing the MFI in the presence of an inhibitor with the MFI in the absence of inhibitor. Data represent means + SD of triplicates. Similar results were obtained in three independent experiments.



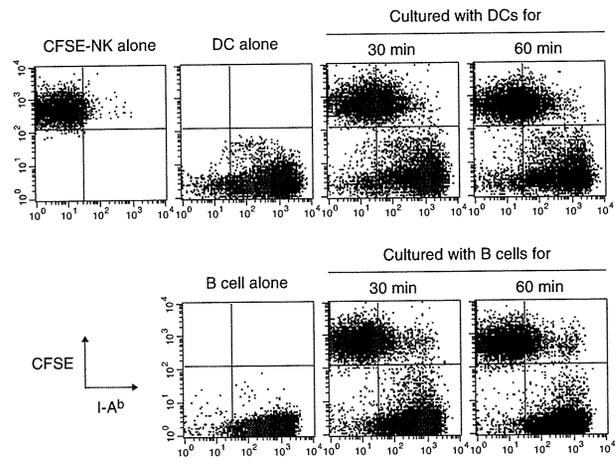
**Fig. 56.** Kinetic analysis of intercellular transfer of MHCII, CD80, CD86, and CD11c. (A) CFSE-labeled NK cells were cocultured with DCs at a 1:1 ratio. After the indicated periods, cells were stained with anti-CD86 and anti-CD11c mAbs. (B) The MFI of I-A<sup>b</sup>, CD80, CD86, and CD11c expression on NK cells cocultured with DCs was analyzed by flow cytometry. Data represent the mean of triplicates. Similar results were obtained in two independent experiments.



**Fig. S7.** Antigenic phenotypes of IFN-producing killer DCs (IKDCs) and MHCII-dressed NK cells. (A) Collagenase-digested low-density spleen cells were stained with anti-B220, anti-CD11c, anti-NK1.1, and anti-PDCA-1 mAbs. The putative IKDC population is indicated by a red arrow. (B) Putative IKDCs and MHCII-dressed NK cells were stained with the indicated mAb (thick histograms) or respective isotype control mAb (thin histograms). Similar results were obtained in three independent experiments.



**Fig. S8.** DC acquisition of Ly49G2 from NK cells. TAMRA [5- (and 6-) carboxytetramethylrhodamine succinimidyl ester]-labeled activated NK cells were co-cultured with splenic DCs at a 1:1 ratio for 1 h. Then cells were stained with anti-I-A<sup>b</sup> and anti-Ly49G2 mAbs or control rat IgG<sub>2a</sub>. In histograms, black and red lines indicate control rIgG<sub>2a</sub> and anti-Ly49G2 mAb staining, respectively. The percentage of Ly49G2-positive cells in R1, R2, and R3 is indicated. Similar results were obtained in two independent experiments.



**Fig. S9.** NK acquisition of MHCII from B cells. CFSE-labeled activated NK cells were cocultured with splenic DCs or B cells at a 1:1 ratio for the indicated periods of time. Then cells were stained with anti-I-A<sup>b</sup> mAb. Similar results were obtained in two independent experiments.

