

CD8⁺CD122⁺CD49d^{low} regulatory T cells maintain T-cell homeostasis by killing activated T cells via Fas/FasL-mediated cytotoxicity

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The Fas/FasL (CD95/CD178) system is required for immune regulation; however, it is unclear in which cells, when, and where Fas/FasL molecules act in the immune system. We found that CD8⁺CD122⁺ cells, which are mostly composed of memory T cells in comparison with naïve cells in the CD8⁺CD122[−] population, were previously shown to include cells with regulatory activity and could be separated into CD49d^{low} cells and CD49d^{high} cells. We established in vitro and in vivo experimental systems to evaluate the regulatory activity of CD122⁺ cells. Regulatory activity was observed in CD8⁺CD122⁺CD49d^{low} but not in CD8⁺CD122⁺CD49d^{high} cells, indicating that the regulatory cells in the CD8⁺CD122⁺ population could be narrowed down to CD49d^{low} cells. CD8⁺CD122⁺ cells taken from lymphoproliferation (*lpr*) mice were resistant to regulation by normal CD122⁺ Tregs. CD122⁺ Tregs taken from generalized lymphoproliferative disease (*gld*) mice did not regulate wild-type CD8⁺CD122[−] cells, indicating that the regulation by CD122⁺ Tregs is Fas/FasL-dependent. CD122⁺ Tregs taken from IL-10-deficient mice could regulate CD8⁺CD122[−] cells as equally as wild-type CD122⁺ Tregs both in vitro and in vivo. MHC class I-missing T cells were not regulated by CD122⁺ Tregs in vitro. CD122⁺ Tregs also regulated CD4⁺ cells in a Fas/FasL-dependent manner in vitro. These results suggest an essential role of Fas/FasL as a terminal effector of the CD122⁺ Tregs that kill activated T cells to maintain immune homeostasis.

Tregs | Fas/FasL | cytotoxicity T cells | immune homeostasis | central memory phenotype

Fas (CD95) and its ligand FasL (CD178) compose a unique system that is strongly related to programmed cell death (1). FasL has been considered one of the effector molecules involved in the killing of target cells by cytotoxic T lymphocytes (CTLs) (2). When Fas binds to FasL, it induces downstream signal transduction pathways that subsequently activate cell death induction pathways (3, 4). Thus, the Fas/FasL system appears to act as an effector for CTL-mediated killing of virus-infected or cancer cells, similar to the perforin-granzyme system (5, 6). However, because Fas-mutant (*lpr*) and FasL-deficient (*gld*) mice show lymphoproliferative changes, it has been suggested that the Fas/FasL system is important for suppression/regulation of activated effector T cells (7, 8).

Molecular mechanisms after Fas activation have been thoroughly investigated, and the signal transduction pathway has been largely elucidated (9, 10). However, research on the cellular events that use the Fas/FasL system has progressed comparatively slowly. There are only a few relevant reports in this respect, mostly of human CD4⁺Foxp3⁺ Tregs that use the Fas/FasL system for their regulatory activity. [In this article, we use the term “Treg(s)” for any kind of T cells that show immune regulatory activity.] To complicate matters further, some contradictory reports claim that such Fas/FasL-engaging Tregs do not exist (8, 11, 12). No reliable reports suggesting that CD8⁺ T cells use the Fas/FasL system for their

regulatory action are available. Therefore, it is not clear precisely which subset of T cells express FasL or where Fas/FasL-dependent CD8⁺ Tregs, if such cells exist, are located and at which point they function. Thus, the ultimate pathophysiological role of the Fas/FasL system is still unknown.

Regulation of the immune reaction is of pivotal importance for maintaining health in multicellular organisms. In highly developed animals, Tregs, a subset of T lymphocytes especially known as CD4⁺CD25⁺Foxp3⁺ cells, are the main regulators of the immune response (13–16). However, it is not quite clear whether CD8⁺ regulatory T cells exist; there are only few and contradictory reports on their existence, in contrast to the reports on CD4⁺ Tregs. A population of predominantly CD8⁺ suppressor T cells has been described and debated in the 1980s (17, 18). In the 2000s, we found and reported that the cells of central memory phenotype (CD8⁺CD122⁺) also possess the regulatory function, and some other reports regarding CD8⁺ Tregs with some contradictory findings have been published. To date, more than 10 CD8⁺ Treg populations with different markers have been reported (19).

One of the best characterized CD8⁺ Treg populations is the CD8⁺CD122⁺ Treg (122⁺ Treg) population. To confirm the existence of CD8⁺ Tregs, markers that may be possible to distinguish Tregs from other T cells were examined. We previously found that CD49d can separate CD8⁺CD122⁺ cells into at least two subsets (CD49d^{low} and CD49d^{high}) (20). To judge which cells have a stronger regulatory activity, we prepared two experimental systems:

Significance

Our study demonstrated that CD8⁺CD122⁺CD49d^{low} regulatory T cells induced apoptosis in target T cells depending on death receptor Fas (CD95)–FasL (CD178) interaction. This in vitro phenomenon reflected the regulatory activity of these cells in vivo, which was also reproducible in the in vivo experimental system of adoptive T-cell transfer. By using natural mutant mice of Fas or FasL and the gene-targeted mice of IL-10, we confirmed that the regulatory activity of CD8⁺CD122⁺CD49d^{low} cells was mediated by Fas–FasL but not by IL-10. In addition, the necessity of MHC class I molecules on the target T cells was also confirmed by analysis using CD8⁺ T cells that lack β2-microglobulin and consequently class I MHC.

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confirmed that CD8⁺ Tregs are encompassed in the CD49d^{low} subset. Thus, in the following text, CD49d^{low} cells are described as “CD8⁺ Tregs” to reduce the text volume.

CD8⁺CD122⁺CD49d^{low} Tregs Cannot Eliminate or Suppress Fas-Mutated CD8⁺CD122⁺ Cells from *lpr* Mice. We tested the regulatory activity of CD8⁺ Tregs under Fas-related conditions. CD8⁺ Tregs, taken from wild-type mice, were cocultured with CD8⁺CD122⁺ cells, taken from Fas-mutant *lpr* mice, and the percentage of target CD8⁺CD122⁺ cells among total live cells was determined at various time points. The ratio of CD8⁺CD122⁺ cells from *lpr* mice to wild-type CD8⁺ Tregs decreased as coculture continued (Fig. 2*A* and *B*). However, the rate and extent of decrease were significantly higher when CD8⁺CD122⁺ cells and CD8⁺ Tregs, both taken from wild-type mice, were cocultured (Fig. 2*A* and *B*). These changes in the ratio of cell number were also observed in the data of absolute cell numbers (Fig. 2*C*). The proliferation rate of CD8⁺CD122⁺ cells from *lpr* mice was

not different from that of CD8⁺CD122⁺ cells from wild-type mice (Fig. 2*D*). In addition, we examined the proliferation of cocultured cells by the dilution of prelabeled CFSE and further confirmed that any types of cells used in this study show a similar pattern of CFSE peaks (Fig. 2*D*). This result confirmed that the difference observed between CD8⁺CD122⁺ cells from *lpr* mice and those from wild-type mice in the coculture experiment was not due to a difference in proliferation rate but due to resistance to apoptosis of the *lpr* mice-derived CD8⁺CD122⁺ cells. In the in vivo experiment, RAG-2-deficient mice that had received CD8⁺CD122⁺ cells taken from *lpr* mice died, similar to those injected with CD8⁺CD122⁺ cells taken from wild-type mice. Mice that had received a mixture of CD8⁺CD122⁺ cells from *lpr* mice and CD8⁺ Tregs obtained from wild-type mice died, similar to those that had received CD8⁺CD122⁺ cells from *lpr* mice alone (Fig. 2*E*). Together, these findings indicated that Fas-mutant CD8⁺CD122⁺ cells are resistant to regulation by CD8⁺ Tregs, which have a marked effect on wild-type CD8⁺CD122⁺ cells and prolong the survival of CD8⁺CD122⁺ cell-recipient mice.

FasL-Mutant CD8⁺ Tregs Cannot Eliminate or Suppress Conventional CD8⁺CD122⁺ T Cells. First, the expression of FasL in CD8⁺ Tregs was confirmed by real-time PCR (Fig. 3*A*). Furthermore, the existence of the FasL product on the cell surface was confirmed by flow cytometry (Fig. 3*B*). This result indicated that FasL is a candidate mediator of killing of Fas-expressing target cells.

Next, we examined the activity of FasL-mutant CD8⁺ Tregs. CD8⁺ Tregs taken from *gld* mice were cocultured with CD8⁺CD122⁺ cells taken from wild-type mice, and the percentage of target cells (CD8⁺CD122⁺) among CD8⁺ Tregs was determined at various time points. The percentage of wild-type CD8⁺CD122⁺ cells among CD8⁺ Tregs from *gld* mice decreased over time; however, the rate and extent of decrease were significantly lower than when both types of cells were taken from wild-type mice (Fig. 3*C* and *D*). The difference was also observed in the absolute cell number (Fig. 3*C*). The proliferation rate of CD8⁺ Tregs taken from *gld* mice was not different from that of wild-type CD8⁺ Tregs (Fig. 3*E*). This result confirmed that an insufficient decrease of cells cocultured with CD8⁺ Tregs taken from *gld* mice, compared with those cocultured with wild-type CD8⁺ Tregs, was not due to a relative decrease in the proliferation rate of CD8⁺ Tregs taken from *gld* mice but rather due to insufficient elimination of CD8⁺CD122⁺ cells by FasL-mutated CD8⁺ Tregs.

In the in vivo experiment, RAG-2-deficient mice that had received a mixture of wild-type CD8⁺CD122⁺ cells and CD8⁺ Tregs from *gld* mice died at the same rate as those that had received wild-type CD8⁺CD122⁺ cells alone (Fig. 3*F*). Thus, CD8⁺ Tregs taken from FasL-mutant *gld* mice demonstrated no ability to control CD8⁺CD122⁺ cells and could not prolong the survival of mice that received these cells.

CD8⁺ Tregs Induce Apoptosis in Target-Activated CD8⁺CD122⁺ T Cells.

To confirm the cytotoxic mechanisms underlying the regulatory action of CD8⁺ Tregs, we examined cell death during coculture by staining the cells with annexin V and 7-amino actinomycin D (7-AAD) (Fig. 4*A* and *B*). Although we could not detect annexin V⁺7-AAD⁺ cells, we observed annexin V⁺7-AAD⁺ cells. CD8⁺CD122⁺ cells cocultured with CD8⁺ Tregs showed a higher percentage of 7-AAD⁺ cells than those cocultured with CD49d^{high} cells, indicating that CD8⁺ Tregs induced apoptosis in CD8⁺CD122⁺ T cells. CD8⁺CD122⁺ cells taken from *lpr* mice did not show an increase in 7-AAD-stained cells after coculture with CD8⁺ Tregs, indicating that Fas-mutant CD8⁺CD122⁺ cells are resistant to apoptosis induced by CD8⁺ Tregs (Fig. 4*B*).

CD8⁺CD122⁺CD49d^{low} T Cells Induce Apoptosis via Caspase 8 Pathway.

Judging from the experimental results both in vitro and in vivo, it is strongly suspicious that the regulatory mechanism performed by

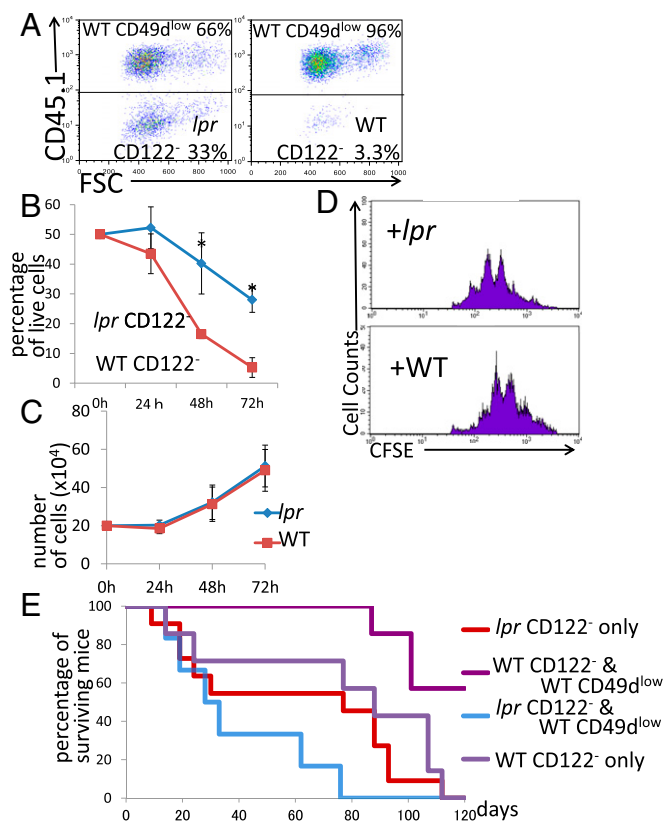


Fig. 2. (A and B) Analysis of regulatory activity in coculture of CD49d^{low} cells (CD45.1) and CD8⁺CD122⁺ cells (CD45.2) derived from *lpr* mice and wild-type mice. (A) Representative results after 72 h of coculture obtained from three independent experiments are shown. (Left) CD122⁺ cells (*lpr*) cocultured with CD49d^{low} cells. (Right) CD122⁺ cells (WT) cocultured with CD49d^{low} cells. (B) The percentages of CD8⁺CD122⁺ cells among total live cells are shown. *Significantly different ($P < 0.05$, Student's t test). (C) Absolute numbers of CD122⁺ T cells derived from WT mice (red line) and *lpr* mice (blue line) in simple culture were determined. At all time points, the values obtained for *lpr* mice-derived cells and WT mice-derived cells were not significantly different ($P > 0.05$, Student's t test). (D) CD8⁺CD122⁺ T cells (WT or *lpr*) labeled with CFSE were cocultured for 48 h and analyzed. The dilution patterns of CFSE are shown as representative results of three independent experiments. (E) Analysis of adoptive T-cell transfer of CD122⁺ T cells from WT or *lpr* mice mixed with CD49d^{low} T cells of WT mice into RAG-2-deficient mice. Addition of WT mice-derived CD49d^{low} T cells to *lpr* mice-derived CD122⁺ cells (blue line) did not affect survival; survival of these mice was not significantly different from that of mice receiving *lpr* mice-derived CD122⁺ cells alone (red line; log-rank test, $P = 0.0878$; Wilcoxon test, $P = 0.2537$).

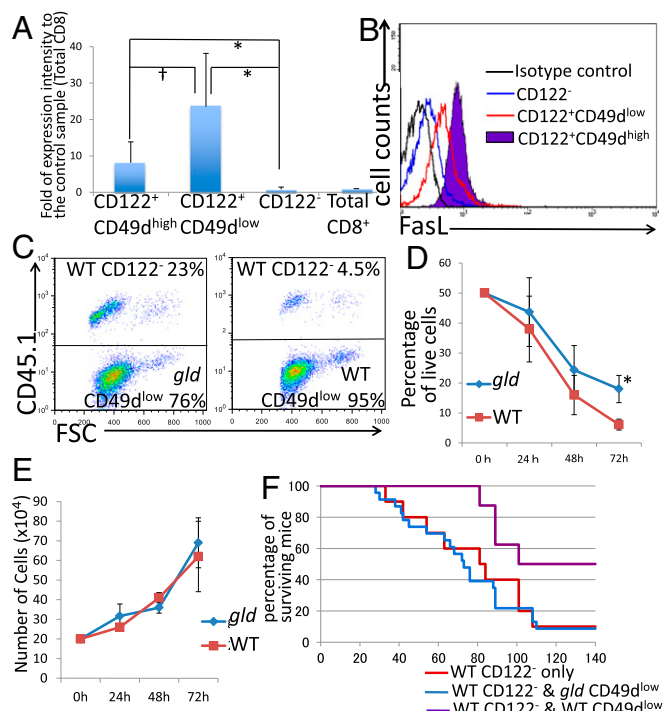


Fig. 3. (A) Analysis of RNA expression level of FasL in each population of CD8⁺ T cells by real-time PCR. For each sample, FasL mRNA levels were normalized to β -actin and are expressed as fold change relative to the control sample (1.00). *Not significantly different ($P > 0.05$, Student's t test); **significantly different ($P < 0.01$, Student's t test). (B) Analysis of FasL expression of flow cytometry. (C and D) Analysis of coculture of CD8⁺CD122⁻CD45.1⁺ T cells and CD49d^{low} from *gld* or WT mice. (C) Representative results obtained from three independent experiments. (D) *Significantly different ($P < 0.05$, Student's t test). (E) Cell growth analysis of CD49d^{low} T cells from *gld* mice and WT mice. At all time points, the results obtained from two different mice were not significantly different ($P > 0.05$, Student's t test). (F) Analysis of adoptive T-cell transfer of CD122⁻ T cells (WT) mixed with CD49d^{low} T cells (WT or *gld*) into RAG-2-deficient mice. Addition of *gld* mice-derived CD49d^{low} T cells to CD122⁻ cells (blue line) had no significant effect on survival compared with transfer of CD122⁻ cells alone (red line; log-rank test, $P = 0.7608$; Wilcoxon test, $P = 0.6859$).

the CD49d^{low} cells is elimination of cells by using the Fas/FasL system. To prove the involvement of the Fas/FasL system in our T-cell coculture assay, we introduced the assay system of PhiPhiLux and CaspaLux. It is well known that caspases are the key molecule of apoptosis, and caspase 8 is especially directly linked to the Fas-derived signal. When caspase 8 is active, CaspaLux-8, a substrate for caspase 8, is digested and becomes a fluorescent substance that can be detected by flow cytometry. As a result, more caspase 8-active cells were observed in the cells cocultured with CD49d^{low} cells (Fig. 4 C and D), indicating that Fas and its downstream caspase, caspase 8, was activated during the process of apoptosis induction.

CD8⁺ Tregs Are Increased in Fas/FasL-Mutant Mice but Not in IL-10-Deficient Mice. We demonstrated that CD8⁺ Tregs, but not CD49d^{high} cells, display regulatory activity that can eliminate cocultured CD8⁺CD122⁻ cells in vitro and prolong survival of CD8⁺CD122⁻ T-cell-recipient RAG-deficient mice in vivo. The cause of death in CD8⁺CD122⁻ T-cell-recipient RAG-deficient mice was not evident but might involve failure of multiple organs, including bone marrow, lung, liver, etc. Additionally, we found that lack of Fas/FasL function affects the regulatory activity of CD8⁺ Tregs both in vitro and in vivo. Therefore, we expected some changes in CD8⁺ Tregs of Fas/FasL-mutant mice. The absolute number of CD8⁺ Tregs and their percentage on total CD8⁺

cells were significantly increased in *lpr* and *gld* mice but not in IL-10-deficient mice, compared with wild-type mice (Fig. 4 E and F). This finding indicates the existence of a negative feedback mechanism that controls the activity of CD8⁺ Tregs.

CD8⁺ Tregs Cannot Eliminate or Suppress MHC Class I-Deficient CD8⁺CD122⁻ Cells. Next, we asked whether recognition of target cells by CD8⁺ Tregs depends on T-cell receptor (TCR)–MHC class I interaction. To address this question, we used β 2-microglobulin-deficient instead of MHC class I-deficient mice; β 2-microglobulin-deficient mice have very few CD8⁺ cells because of a lack of positive selection of CD8⁺ T cells. This problem can be addressed by generating wild-type mice with bone marrow reconstituted with MHC class I-deficient bone marrow. When CD8⁺CD122⁻ cells, taken from the bone marrow-reconstituted mice, were cocultured with CD8⁺ Tregs (Fig. 5 A and B). The rate and extent of reduction were lower than when they were cocultured with wild-type CD8⁺CD122⁻ cells (Fig. 5 A and B). The proliferation of CD8⁺CD122⁻ cells from mice in which the bone marrow had been reconstituted with that of β 2-microglobulin-deficient mice was not significantly different from that of CD8⁺CD122⁻ cells

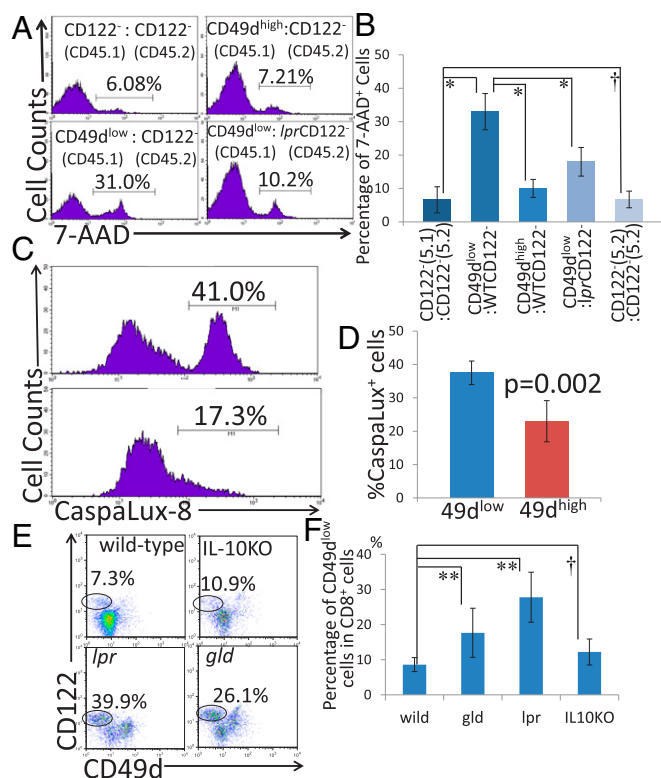


Fig. 4. (A) Result of 7-AAD analysis after 48 h of coculture. Percentages of 7-AAD-positive cells among total cultured cells are shown in the histogram. (B) Percentages of 7-AAD-positive cells in coculture. *Significantly different ($P < 0.01$, Student's t test); †not significantly different ($P > 0.05$, Student's t test). (C and D) Analysis of caspase 8 activity in CD122⁻ (CD45.1⁺) cells using CaspaLux8 after coculture with CD49d^{low} or CD49d^{high} (CD45.1⁻) cells. (E) Representative results obtained from three independent experiments. (F) Percentages of cells positive for CaspaLux8 in CD122⁻ CD45.1⁺ cells are shown. The data are averages of three independent experiments. (G) Spleen cells taken from WT, IL-10-deficient, *lpr*, and *gld* mice were analyzed by flow cytometry. Representative results obtained from three independent experiments for each mouse strain are shown. (H) Percentages of CD49d^{low} cells in total CD8⁺ cells in each group are shown. Statistically significant differences between the WT and each mutant strain were determined. **Significantly different ($P < 0.01$, Student's t test); †not significantly different ($P > 0.05$, Student's t test).

cytokine could be the effector molecule of these CD8⁺ Tregs. Although both the mechanism of recognition of target cells and the effector molecule of regulation by CD8⁺ Tregs remain to be elucidated, this study yielded some important clues: Activation of Fas via interaction with FasL is largely associated with the killing of T cells that are no longer needed and rather harmful for survival. Another important molecular interaction is that of MHC class I and TCR. β 2-microglobulin-deficient cells are neither regulated in the previous study (33) nor in the present study, suggesting the importance of specific recognition between “the regulatory cells” and “the regulated cells.” Another important molecule involved in the action of CD8⁺ Tregs might be Qa-1, a nonclassical MHC class I molecule (MHC-Ib), as H. Cantor and his colleague found and reported a CD8⁺ T-cell subset that showed a regulatory activity under a Qa-1-restricted manner (30). In their recent publications, however, the Qa-1-restricted CD8⁺ Tregs they proposed were apparently shown to be CD122⁺CD44⁺ (34, 35). The Qa-1-restricted CD8⁺ Treg is overlapped with or involved in the CD8⁺CD122⁺ Treg reported by H. Suzuki and his colleagues (22). Two CD8⁺ Tregs that were formerly thought to be distinct cell populations turned out to be the same one, further confirming the existence of CD8⁺ regulatory T cells, regardless of whether they are the naturally arising type or the induced type. Although the relation between CD4⁺ Tregs and CD8⁺ Tregs has not been clarified in detail, both Tregs are indispensable to maintain T-cell homeostasis, and their cooperative work has been suggested in the prevention and cure of an autoimmune disease (31).

This study forms a basis for continued research of immune suppression by CD8⁺CD122⁺ Tregs. Elucidation of the effector mechanisms of CD8⁺ and CD4⁺ Tregs may pave the way to improve immune regulation by Tregs in clinical practice.

Materials and Methods

Mice. Animal care was performed according to the guidelines of Nagoya University (Nagoya, Japan). Experimental protocols were approved by the Ethics Committee of the Nagoya University Graduate School of Medicine (Nos. 27015, 27017, and 27365).

Adoptive T-Cell Transfer to RAG-2^{-/-} Mice. Sorted CD8⁺CD122⁺ T cells (4×10^5 per recipient mouse) simply or mixed with CD8⁺CD122⁺ T cells (49d^{low} or CD49^{high}) (1×10^5 per recipient mouse) were i.v. transferred into 8–10-wk-old RAG-2-deficient mice that had received a sublethal dose of irradiation (5.5 Gy). The survival of the T-cell-recipient mice was followed up after adoptive T-cell transfer, and the results were analyzed using Kaplan–Meier survival curves.

Activation of CD8⁺ T Cells. CD8⁺ T cells in any desired population were collected from murine spleen and lymph node cells by using a FACS SORP Aria II cell sorter (BD Biosciences) and cultured in an anti-mCD3 antibody-coated 96-well plate for 24 h (1.0×10^5 cells per well).

T-Cell Culture. Activated T cells as described above were either cocultured or cultured alone in RPMI-1640 media containing 10% (vol/vol) FBS and 20 ng/mL of interleukin-2 (PeproTech Inc.) in a round-bottom 96-well plate for 72 h (0.5×10^5 cells per well).

For other information, see *SI Materials and Methods*.

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