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Evidence that CD8⁺ Dendritic Cells Enable the Development of γδ T Cells that Modulate Airway Hyperresponsiveness²

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Abstract

Airway hyperresponsiveness (AHR), a hallmark of asthma and several other diseases, can be modulated by $\gamma\delta$ T cells. In mice sensitized and challenged with ovalbumin, AHR depends on allergen-specific $\alpha\beta$ T cells, but $V\gamma\delta1^+\gamma\delta$ T cells spontaneously enhance AHR, whereas $V\gamma4^+\gamma\delta$ T cells after being induced by airway challenge suppress AHR. The activity of these $\gamma\delta$ T cell modulators is allergen-nonspecific, and how they develop is unclear. We now show that CD8 is essential for the development of both the AHR-suppressor and enhancer $\gamma\delta$ T cells although neither type needs to express CD8 itself. Both cell types encounter CD8-expressing non-T cells in the spleen, and their functional development in an otherwise CD8-negative environment can be restored with transferred spleen cell preparations containing CD8⁺ DC, but not CD8⁺ T cells or CD8⁻ DC. Our findings suggest that CD8⁺ DC in the lymphoid tissues enable an early step in the development of $\gamma\delta$ T cells, through direct cell-contact. DC-expressed CD8 might take part in this interaction.

Keywords

Rodent; dendritic cells; T cells; cell differentiation; spleen and lymph nodes

Introduction

Airway hyperresponsiveness (AHR) is a symptom in several potentially life-threatening diseases, including asthma and chronic obstructive pulmonary disease. The pathways leading to AHR are not yet completely mapped. In allergic asthma, antigen-specific T cells are initiators of a cascade of immune-dependent mechanisms that result in the production of the critical cytokine interleukin 13, eosinophilic airway inflammation, allergen-specific IgE antibodies, mast cell activation, goblet cell differentiation and mucus production, and AHR (1,2). The development and role of allergen-specific $\alpha\beta$ T cells have been studied extensively in animal models of allergic airway inflammation and AHR. In the process, it became apparent that innate antigen-nonspecific T cells were involved as well, which are capable of suppressing and enhancing AHR in allergic airway disease and other pathological conditions of the lung (3-7). Both $\alpha\beta$ T cells and $\gamma\delta$ T cells are included among these innate AHR-modulating T cells and are potentially attractive targets for immune intervention (8). However, little is known

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about these cells and their development. We have studied AHR-modulating $\gamma\delta$ T cells *in vivo*. In mice sensitized and challenged with ovalbumin (OVA), which display hypersensitivity when given the cholinergic agonist methacholine (MCh) (9), we found that $\gamma\delta$ T cells can dramatically alter AHR whereas their effect on airway inflammation is comparatively small (5). The $\gamma\delta$ T cells capable of modulating AHR belong to two subsets distinguished by their T cell receptors (TCR): A V γ 1⁺ subset which contains $\gamma\delta$ T cells spontaneously capable of enhancing AHR (10,11) and a V γ 4⁺ subset which contains $\gamma\delta$ T cells that can be induced to suppress AHR (12-15). Neither cell-type requires specific Ag-priming (11,15) and it is unclear how these AHR-modulators develop. However, both are found in spleen and lung (16). Moreover, following airway challenge, selectively depleting either subset in the lung (using aerosolized inhaled anti TCR-antibodies) alters AHR in the predicted fashion, i.e. depletion of V γ 1⁺ cells decreases and depletion of V γ 4⁺ cell increases AHR (8,12), suggesting that the $\gamma\delta$ T cells within the challenged lung are functionally competent.

In the development of antigen-specific T cells, dendritic cells (DC) are critical (17). The role of DC in the development of innate T cells is less well defined. Evidence that $\gamma\delta$ T cells and DC interact has come mainly from studies in vitro. These revealed maturational effects on both the DC and the $\gamma\delta$ T cells (18-24), raising the question of whether such interactions might also be critical for the functional development of yo T cells in vivo. However, DC are heterogeneous and their functions diverse (25-31). The present study began with the finding that AHRsuppression by $\gamma\delta$ T cells could not be demonstrated in mice genetically deficient in CD8 α , and our attempts to determine which cells express the required CD8. Using adoptive cell transfer, we find that the $\gamma\delta$ T cells themselves need not express CD8, but that CD8 is required in the donor mice in which the $\gamma\delta$ T cells develop. Further, the development of $\gamma\delta$ T cells capable of AHR-suppression can be restored in CD8-deficient mice by reconstituting them with cell preparations containing CD8⁺DC, but not with CD8-expressing T cells or CD8⁻ DC. Mice deficient in CD8 α also fail to give rise to $\gamma\delta$ T cells capable of AHR-enhancement and again, their development can be restored by transfer of CD8⁺ DC, suggesting that interactions with $CD8^+ DC$ in lymphoid tissues represent a critical step in the development of $\gamma\delta$ T cells towards functional competence.

Materials and Methods

Animals

C57BL/6, B6.CD8 $\alpha^{-/-}$, B6.TCR- $\beta^{-/-}$, B6.TCR- $\delta^{-/-}$, and B6.TCR- $\beta^{-/-}/\delta^{-/-}$ mice were purchased from The Jackson Laboratory (Bar Harbor, Maine); TCR-V γ 4/6^{-/-} mice deficient in V γ 4⁺ and V γ 6⁺ $\gamma\delta$ T cells (32) were a kind gift from K. Ikuta (Kyoto, Japan). They were backcrossed to the C57BL/6 genetic background and established after 11 backcrossed generations (designated B6.TCR-V γ 4/6^{-/-}). B6.TCRV γ 4/6^{-/-}CD8 $\alpha^{-/-}$ mice were generated starting from the F2 generation of a cross between B6.CD8 $\alpha^{-/-}$ and B6.TCR-V γ 4/6^{-/-} mice. All mice were maintained on an OVA-free diet, and were cared for at National Jewish Medical and Research Center (Denver, Colorado), following guidelines for immune deficient animals when necessary. All experiments were conducted under a protocol approved by the Institutional Animal Care and Use Committee.

Sensitization and airway challenge

Mice received the following treatments: no OVA treatment (NT) or sensitization to OVA by intraperitoneal injection of 20µg of OVA (Grade V; Sigma) emulsified in 2.25mg alum (AlumImject®; Pierce, Rockford, Illinois) in a total volume of 100µl on days 0 and 14 (2ip), or sensitization followed by airway exposure to nebulized OVA (1% in saline), using ultrasonic nebulization (particle size 3–5µm) for 20 mins on days 28, 29 and 30 (2ip3N). Donors of DC-

preparations either received no treatment or only the sensitizing treatment (2ip). Airway responsiveness was assessed 48h after the last nebulized OVA exposure for 2ip3N treated mice.

Depletion with monoclonal antibody against TCR

Cell-depletion was achieved by injection into the tail vein of 200 μ g hamster mAb against V γ 4 (UC3) (33), 3 days before the first OVA challenge, and depletion was assessed as described (10). Sham depletion was carried out using hamster Ig (Jackson Laboratories, Bar Harbor, Maine). Throughout this article, we use the nomenclature for murine TCR-V γ genes introduced by Heilig and Tonegawa (34).

Determination of airway responsiveness

Airway responsiveness was assessed as a change in respiratory system resistance after provocation with aerosolized methacholine (MCh) using a method previously described in detail (5). MCh aerosol was administered for 10 s (60 breaths/min, 0.5 ml of tidal volume) in increasing concentrations as indicated in the figures. Maximum values of lung resistance (R_L) and minimum values of dynamic compliance (C_{dyn}) were recorded and expressed as percentage change from baseline after saline aerosol.

Histological analysis of the spleen

For the tissue preparation, sectioning and staining, essentially the same methods were used as described previously for the analysis of $\gamma\delta$ T cells in the lung (16). Briefly, sections of 5–10 mm thickness cut from frozen blocks were mounted on microscopic slides, air dried, dehydrated in acetone, hydrated, blocked with a mixture of normal mouse serum and Avidin, washed, stained with primary and secondary antibodies, and finally mounted with coverslips. Slides were viewed using a Leica (Knowhill, UK) DMRXMA upright fluorescent microscope, and digital images were generated on an Apple MacIntosh computer connected with the microscope, using the SlideBook imaging program (3I Inc., Atlanta, GA). The tissue autofluorescence (green) was used to distinguish overall tissue organization, especially the localization of central arterioles, and the density of the cellular distribution. The higher celldensity of the white pulp around the arterioles was used to determine the extent of the PALS. Cells and cell-contacts were enumerated and compared either across the entire tissue (whole spleen) or limited to the PALS. Antibodies used for histology: anti TCR-δ (mAb GL3), anti Vγ4 (mAb UC3), anti Vγ1 (mAb 2.11), anti-CD8α (mAb 53-6.7, eBioscience), anti-CD11c (mAb HL3, Pharmingen), anti CD103 (mAb 2E7, eBisocience), anti DEC-205 (mAb ATCC# HB 290), plus secondary reagents as described(16).

Cell preparation from spleen and T cell-sorting using the MO FLO

A suspension of splenocytes was prepared by mechanical dispersion. Suspended cells were treated with Gey's red cell lysis solution. Nylon wool non-adherent (NAD) cells were prepared from spleens of OVA-sensitized and challenged B6.TCR- $\beta^{-/-}$ mice 2 days after the last challenge. These preparations contained >75% T cells. Total cell counts were determined using a Coulter counter. NAD cells in PBS/5% FBS were incubated with FITC conjugated anti-Vγ4 mAb (UC3) and PE-conjugated anti-CD8 α (53–6.7, Pharmingen) or PE-conjugated anti-CD8 β (53–5.8, Pharmingen) (20mins, 4°C), and then washed. Cells were next sorted based on their expression of Vγ4 and CD8 (α or β) using a MO FLO cell sorter. Purified Vγ4⁺/CD8⁺ or Vγ4⁺/CD8⁻ cells were washed in PBS and resuspended to 5 × 10⁴ cells/ml PBS, and 1 × 10⁴ cells/mouse injected in 200µl of PBS via the tail vein into OVA sensitized B6.TCR-Vγ4/6^{-/-} mice within 1 hour prior to the first airway challenge.

Adoptive transfer of yo T cells

Nylon wool nonadherent (NAD) cells were prepared from spleens of OVA-sensitized and challenged B6.TCR- $\beta^{-/-}$ or B6.CD8 $\alpha^{-/-}$ mice 2 days after the last challenge. These cells contained >75% T cells. Total cell counts were determined using a Coulter counter. NAD cells in PBS/5% FBS were incubated with biotinylated anti-V γ 4 mAb (UC3) or anti-V γ 1 mAb (2.11) (35) (20 mins, 4°C) then washed and incubated with streptavidin-conjugated magnetic beads (streptavidin microbeads; Miltenyi Biotec, Bergisch Gladbach, Germany) for 15 mins at 4°C and passed twice through magnetic columns to purify V γ 4⁺ or V γ 1⁺ cells. This produced a cell population containing >80% V γ 4⁺ or V γ 1⁺ viable cells as determined by two-color staining with anti-TCR- δ and anti-V γ 4 or anti-V γ 1 mAbs. These splenic V γ 4⁺ or V γ 1⁺ cells were washed in PBS and resuspended to 5 × 10⁴ cells/ml PBS and 1 × 10⁴ cells/mouse were injected in 200µl of PBS via the tail vein into OVA sensitized B6.TCR-V γ 4/6^{-/-} or B6.TCR- $\delta^{-/-}$ mice, within 1 hour prior to the first airway challenge.

Phenotype of splenocytes and enriched DC from B6.TCR- $\beta/\delta^{-/-}$ mice

Red cell depleted splenocytes from naïve or 2ip treated B6.TCR- $\beta/\delta^{-/-}$ mice (or magnetically enriched DC) were incubated with APC conjugated anti-CD8 α mAb (53–6.7, eBioscience), FITC conjugated anti-CD11c mAb (HL3, Pharmingen), biotinylated anti CD45R (B220, eBioscience, clone RA3–6B2), biotinylated anti CD3 ϵ mAb (KT3) and PE conjugated anti-MHC II (M5/114.15.2, Pharmingen) in PBS/5% FBS (20min, 4°C), then washed and analyzed on a FACSCAN.

Adoptive transfer of accessory non-T cells

Cells were isolated from the spleen of untreated or 2ip treated B6.TCR- $\beta^{-/-}/\delta^{-/-}$ and were treated with Gey's red cell lysis solution. To reconstitute with whole splenocytes, one "spleen equivalent" (~ 5×10^7 cells) was then transferred per 2ip treated B6.CD8 $\alpha^{-/-}$ mouse. A small sample of these cells was analysed for CD8 expression by flow cytometry (mAb 53-6.7-PE, Pharmingen). Approx. 9% were found to be CD8⁺. Reconstituted B6.CD8 $\alpha^{-/-}$ mice were then nebulized as described earlier, and used as donors of $V\gamma 4^+$ or $V\gamma 1^+$ cells. To reconstitute with CD8-enriched or CD8-depleted splenocytes, the red cell depleted splenocytes from 2ip treated B6.TCR- $\beta^{-/-}/\delta^{-/-}$ mice, or B6.TCR- $\delta^{-/-}$ mice (for CD8⁺ T cell adoptive transfer), were incubated with biotinylated anti-CD8 mAb 53-6.7 in PBS/5% FBS (20 min, 4°C), then washed and incubated with streptavidin-conjugated magnetic beads (streptavidin microbeads; Miltenyi Biotec, Bergisch Gladbach, Germany) for 15 mins at 4°C and passed twice over MS magnetic columns to purify CD8⁺ cells. This produced a CD8-enriched fraction containing approx. 45% $CD8^+$ cells as determined by flow cytometry, and a CD8-depleted fraction containing < 1% CD8⁺ cells. Approx. 1×10^{6} viable enriched cells were transferred to 2ip-treated B6.CD8 $\alpha^{-/-}$ recipients. The B6.CD8 $\alpha^{-/-}$ mouse was then nebulized as described earlier and these mice were used as donors of $V\gamma 4^+$ cells. Finally, to reconstitute with more highly enriched CD8⁺ DC, a CD8⁺ purification kit (Miltenyi Biotec, Bergisch Gladbach, Germany) was used as per the manufacturer's recommendations. Briefly, freshly isolated red cell depleted splenocytes derived from the B6.TCR- $\beta^{-/-}/\delta^{-/-}$ mice were incubated with biotinylated antibody cocktail (anti-CD90, anti-CD45R and anti-CD49b) in PBS/5% FBS (10min, on ice), then incubated with anti-biotin microbeads (15min, on ice), then washed and passed over a LD magnetic column to enrich DCs. The enriched DCs were incubated with anti-CD8 α (Ly-2) microbeads (30mins, on ice) then washed and passed twice over a MS magnetic column retaining the CD8⁺ DCs. In the worst case (with 2ip-treated mice), this still produced a preparation containing 60–70% viable CD8 α^+ DC, based on the dual criterion of MHC class II and CD11c expression. The concentration of CD8⁺ DC in this preparation might actually be higher due to variable MHC class II expression. 1.3×10^5 of these enriched cells containing approx. 10^5 CD8⁺DC were transferred to 2ip-treated B6.CD8 $\alpha^{-/-}$ recipients, which were then nebulized as described above and used as donors of V γ 4⁺ cells. Donors of V γ 1⁺ cells (nonsensitized B6.CD8 $\alpha^{-/-}$ mice that received enriched CD8⁺ DC from non-sensitized B6.TCR- $\beta^{-/-}/\delta^{-/-}$ mice - these could be obtained at a higher purity than those from 2ip-treated mice) remained non-challenged.

Statistical analysis

Data are presented as means +/- SEM. The unpaired T test was used for two group comparisons and ANOVA for analysis of differences in three or more groups. Pairwise comparisons were performed using the Tukey-Kramer honest significant difference test. Statistically significant levels are indicated as follows: 1 symbol (*, #, +) = p < 0.05; 2 symbols = p < 0.01; 3 symbols = p < 0.001.

Results

AHR-suppressive $\gamma\delta$ T cells depend on the CD8-molecule but need not themselves express CD8

We have previously found that $V\gamma 4^+\gamma\delta$ T cells suppress AHR in several different mouse strains including C57BL/6 (B6) (12-15). Here, we have used B6 mice to take advantage of available mutations on this background. As predicted by earlier studies (36), by comparison with wildtype mice (Fig.1a), mice deficient in CD8 (B6.CD8 $\alpha^{-/-}$) (Fig.1b) exhibited much-reduced AHR after being sensitized and challenged with OVA. Unexpectedly, however, i.v. injection of a cell-depleting anti TCR-Vy4 antibody (mAb UC3), which increases AHR in wild-type B6 mice (Fig.1a), failed to increase AHR in the CD8-deficient mice (Fig. 1b). This difference suggested a role for CD8 in the suppression of AHR by $\gamma\delta$ T cells but it remained unclear when the molecule might be needed and which cells must express it. To address these questions, we chose a cell-transfer approach. We compared mice adoptively transferred with purified $V\gamma 4^+$ $\gamma\delta$ T cells derived from the spleens of OVA-sensitized and challenged wild-type B6 (Fig. 1c) or B6.CD8 $\alpha^{-/-}$ donors (Fig. 1d), using recipients deficient in V γ 4⁺ cells but expressing normal levels of CD8 (B6.TCR-V γ 4/6^{-/-}). These B6.TCR-V γ 4/6^{-/-} mice develop strong AHR following OVA-sensitization and challenge (13). Following cell transfer, the B6.TCR- $V\gamma 4/6^{-/-}$ mice that received $V\gamma 4^+$ cells from wild-type-mice exhibited reduced AHR (Fig. 1c), but those that received V γ 4⁺ cells from B6.CD8 $\alpha^{-/-}$ mice did not (Fig. 1d). In a similar comparative experiment transferring $\gamma\delta$ T cells derived from the lung, again the wild-typederived cells suppressed AHR whereas the cells from CD8-deficient mice failed to do so (not shown). These results indicated that CD8-expression in the cell-transfer recipients is not sufficient to support the AHR-modulating function of the $\gamma\delta$ T cells. CD8-expression by the donor cells is required, either by the $\gamma\delta$ T cells themselves or by other cells in the donors that might determine the development of the $\gamma\delta$ T cells. Whether in addition CD8 might play a role during yo T cell-mediated AHR-suppression in the cell transfer-recipients was not pursued because the CD8-deficient recipient-mice bred for this purpose (B6.TCR-V γ 4/6^{-/-}CD8 α ^{-/-}) failed to develop AHR (not shown).

We previously found that some of the $V\gamma4^+\gamma\delta$ T cells in spleen and lung express CD8, including CD8 $\alpha\beta$ and CD8 $\alpha\alpha$ (12). To test whether the AHR-suppressive $\gamma\delta$ T cells must themselves express CD8 $\alpha\beta$ or CD8 $\alpha\alpha$, we separated purified $V\gamma4^+$ cells from the spleen of sensitized and challenged donors into CD8 β -positive or –negative, or into CD8 α -positive or –negative fractions, and tested each fraction separately in the cell-transfer model (Fig. 2a, b). All fractions suppressed AHR, indicating that CD8 expression by the $\gamma\delta$ T cells is not required for the suppressive function. This was confirmed by our subsequent experiments with $\gamma\delta$ T cells derived from B6.CD8 $\alpha^{-/-}$ mice (see below). The finding that $\gamma\delta$ T cells need not express CD8 implied that the required CD8 must be expressed by another cell-type. Furthermore, since CD8

is required in the donor, the "other" CD8-expressing cell-type might be involved in the development of the AHR-suppressive $\gamma\delta$ T cells.

Encounters between $\gamma\delta$ T cells and CD8⁺ non-T cells in the spleen

Because $\gamma\delta$ T cells prepared from the spleen are capable of modulating AHR, we examined their distribution and cell-contacts here. In C57BL/6 mice (Fig. 3a), and in B6.TCR- $\beta^{-/-}$ mice (Fig. 3b), both of which give rise to functionally normal AHR-regulatory $\gamma\delta$ T cells (11,13), $\gamma\delta$ T cells were mainly present in the peri-arteriolar lymphoid sheath (PALS), although some were also detected outside the PALS. The $\gamma\delta$ T cells were often in close proximity/contact with DEC-205⁺ or CD8⁺ cells (Fig. 3a and 3b). The CD8⁺ cells are not $\alpha\beta$ T cells because they are present in the spleen of B6.TCR- $\beta^{-/-}$ mice. Both V γ 4⁺ and V γ 1⁺ cells were found in such interactions (Fig.3c, d), with little change following sensitization and challenge (Fig. 4). The frequency of CD8⁺ non-T cells in contact with $\gamma\delta$ T cells was higher inside the PALS (50–60%) than in the spleen in general (30–40%) (Fig. 4). However, the high frequency of such contacts might be a peculiarity of B6.TCR- $\beta^{-/-}$ mice. In accordance with the phenotype of "conventional" CD8⁺ DC (31), the CD8⁺ non-T cells in the PALS of the B6.TCR- $\beta^{-/-}$ mice co-expressed CD11c (Fig.3e), and many co-expressed CD103 and DEC-205 (Fig.3h, i). Collectively, these findings suggest that $\gamma\delta$ T cells, including those types implicated in the modulation of AHR, encounter CD8⁺ DC in the spleen.

Enriched CD8⁺ DC enable the development of AHR-suppressive $\gamma\delta$ T cells in mice lacking CD8

Because CD8-negative $\gamma\delta$ T cells were capable of suppressing AHR (Fig. 2), and because of our previous finding that the AHR-suppressors can develop in mice lacking all $\alpha\beta$ T cells (5, 12,13), we concluded that CD8⁺ non-T cells must satisfy the CD8-requirement. Furthermore, the histological data of this study suggested CD8⁺ DC as candidate CD8⁺ non-T cells. Therefore, we tested whether B6.CD8 $\alpha^{-/-}$ mice might be able to generate V γ 4⁺ AHRsuppressors if reconstituted with CD8⁺ DC. To avoid all CD8⁺ T cells, we chose B6.TCR- $\beta^{-/-}/\delta^{-/-}$ mice as a source of the CD8⁺ accessory cells. Furthermore, because the AHRsuppressive $V\gamma 4^+ \gamma \delta T$ cells from the spleen require prior sensitization and challenge (15), we OVA-sensitized (2ip) both the donors of the CD8⁺ accessory cells and the CD8 α ⁻ recipients, prior to i.v. transfer of the accessory cells. The reconstituted B6.CD8 $\alpha^{-/-}$ mice were then immediately challenged with OVA (3N) so that all cells in these mice were exposed to the full protocol of sensitization and challenge (2ip3N). $V\gamma 4^+ \gamma \delta T$ cells of these mice were subsequently tested for their ability to suppress AHR in B6.TCR-V γ 4/6^{-/-} recipients, using the same method as described in Fig.1. Indeed, these $V\gamma 4^+$ cells suppressed AHR, indicating that genetically CD8-deficient $V\gamma 4^+\gamma\delta$ T cells still retain the potential of developing into AHRsuppressors, and that B6.TCR- $\beta^{-/-}/\delta^{-/-}$ mice (CD8wt) contain non-T cells capable of restoring their development (Fig. 5a). To determine if this capability was contained within the CD8⁺ non-T cells themselves, we compared CD8 α^+ and CD8 α^- splenocytes from B6.TCR- $\beta^{-/-}$ $\delta^{-/-}$ mice for their ability to restore $\gamma\delta$ T cell-development. Only the CD8 α^+ fraction enabled the development of the AHR-suppressors, suggesting that the reconstituting cells themselves must express CD8 (Fig. 5b). This experiment also ruled out the possibility that the mere process of cell-transfer induced the AHR-suppressors. Next, to better enrich splenic CD8 α^+ DC from the B6.TCR- $\beta^{-/-}/\delta^{-/-}$ mice, we used a purification kit for CD8 α^+ DC, which depletes CD90 (Thy-1.2), CD45R (B220) and CD49b-positive cells, followed by positive magnetic beadselection of CD8 α^+ cells. This cell-fraction contained at a minimum 60% CD8⁺ DC (MHC class II⁺, CD11c⁺, CD8 α^+). These enriched cells, when injected i.v. (10⁵ CD8 α^+ DC/ inoculum), were sufficient to support the development of AHR-suppressive V $\gamma 4^+ \gamma \delta$ T cells in B6.CD8 $\alpha^{-/-}$ mice (Fig. 5c). In contrast, neither CD8 α^{-} splenocytes nor enriched CD8 α^{-} DC (Fig. 5c) nor splenic CD8⁺ $\alpha\beta$ T cells (from B6.TCR- $\delta^{-/-}$ donors) had any reconstituting activity (Fig. 5d), albeit provided in larger numbers. These data suggest that CD8⁺ DC function

as accessory cells in the development of the V γ 4⁺ AHR-suppressor $\gamma\delta$ T cells whereas other DC-types or T cells do not.

AHR-enhancing $\gamma\delta$ T cells also depend on CD8, and enriched CD8⁺ DC enable their development

 $V\gamma 1^+ \gamma \delta$ T cells enhance AHR, as shown by cell-depletion and transfer experiments (10). In fact, development of AHR in young mice depends of these cells (10,11) whereas AHR in older mice can develop in their absence while remaining subject to the influence of AHR-suppressive $\gamma \delta$ T cells (5,14). In contrast to the AHR-suppressors, the AHR-enhancing $\gamma \delta$ T cells develop spontaneously, without need for sensitization or challenge of the cell-donors (11). However, we found that B6.CD8 $\alpha^{-/-}$ mice were also unable to produce these AHR-enhancers (Fig. 6a). Therefore, we tested whether reconstituting these mice with CD8⁺ non-T cells would also restore development of the AHR-enhancing $\gamma \delta$ T cells, in the absence of any sensitization or challenge. Indeed, we found that reconstituting B6.CD8 $\alpha^{-/-}$ mice with the CD8 α^+ splenocyte fraction of B6.TCR- $\beta^{-/-}/\delta^{-/-}$ mice (Fig. 6b), or with small numbers of enriched CD8 α^+ DC (Fig. 6c), restored their ability to produce the AHR-enhancers, whereas the CD8 α^- splenocyte fraction had no effect (Fig. 6b). Because the restoration of the AHR-enhancers occurred without sensitizing the source of the DC, and without sensitizing or challenging the donors of the $\gamma \delta$ T cells, this result suggested that interactions between CD8⁺ DC and $\gamma \delta$ T cells are an integral part of normal and spontaneous $\gamma \delta$ T cell-development.

Are the encounters between $\gamma\delta$ T cells and CD8⁺ DC in the spleen functionally relevant?

In B6.CD8 $\alpha^{-/-}$ mice, $\gamma\delta$ T cells are still present in the PALS (Fig. 3f), while in T cell-deficient mice (B6.TCR- $\beta^{-/-}/\delta^{-/-}$), CD8⁺ non-T cells still populate the general vicinity of the central arterioles (Fig. 3g). Although there likely are quantitative differences in cell distributions by comparison with wt mice, this indicates that neither cell type absolutely requires the other to accumulate at this site.

To test whether the transferred CD8⁺ DCs that restore the development of AHR-regulatory $\gamma\delta$ T cells in B6.CD8 $\alpha^{-/-}$ mice indeed appear in the spleen and engage $\gamma\delta$ T cells, we transferred purified CD8⁺ non-T cells from B6.TCR- $\beta^{-/-}/\delta^{-/-}$ spleen in larger numbers (1×10⁶) to facilitate their detection *in situ*. Five days later, when the regulatory $\gamma\delta$ T cells are normally harvested, the CD8⁺ non-T cells were indeed detectable in the PALS of the B6.CD8 $\alpha^{-/-}$ mice, proximal to and frequently in direct contact with the endogenous $\gamma\delta$ T cells (Fig. 3k). Thus, the transferred CD8⁺ non-T cells (in the B6.CD8 $\alpha^{-/-}$ mice) accumulated and encountered $\gamma\delta$ T cells in the same splenic location as the endogenous CD8⁺ non-T cells in the B6.TCR- $\beta^{-/-}$ mice. Because these transferred cells also enable splenic $\gamma\delta$ T cell function (Figs. 5, 6), their interactions with $\gamma\delta$ T cells in the spleen are likely to be critical for the functional development of the AHR-regulators.

Finally, because development of the AHR-suppressive $\gamma\delta$ T cells in the spleen depends on donor sensitization with OVA, we also examined the histological distribution of i.p. injected OVA/alum (Fig. 31). We found OVA/alum in the spleen, in aggregates or contained within CD8⁻ phagocytes, but not within CD8⁺ non-T cells or $\gamma\delta$ T cells. However, both the CD8⁺ non-T cells and the $\gamma\delta$ T cells were seen in proximity/contact with the OVA/alum in the spleen (Fig. 31).

Discussion

Because innate T cells can become powerful modulators of AHR, it is important to understand their development. We have examined the development of two $\gamma\delta$ T cell-types in mice, capable of either suppressing or enhancing allergen-induced AHR. These cells express V $\gamma4$ and V $\gamma1$,

respectively, and they are present in the lung and in secondary lymphoid tissues (10,16). The AHR-suppressive $V\gamma 4^+$ cells are induced by allergen-sensitization and challenge (15) whereas the AHR-enhancing V γ 1⁺ cells develop spontaneously (11). Here, we report three observations related to the development of these two cell-types, namely that the process requires CD8, that the $\gamma\delta$ T cells encounter CD8⁺ non-T cells which appear to be DC in the splenic PALS, and that enriched CD8⁺ DC from the spleen can restore the development of these two $\gamma\delta$ T cell types when transferred into genetically CD8-deficient mice. Crosstalk between $\gamma\delta$ T cells and DC occurs in vitro (20-22,24), but little is known about their cellular interactions in the spleen and other lymphoid tissues. In mice lacking $\alpha\beta$ T cells (B6.TCR- $\beta^{-/-}$), we found many of the $\gamma\delta$ T cells within the splenic PALS in proximity or contact with what appear to be CD8⁺DC based on morphology, phenotype and tissue location. In wt mice (C57BL/6), there are fewer $\gamma\delta$ T cells. These still accumulate in the PALS (Fig. 3a), but due to their smaller numbers and perhaps also because of competition with the $\alpha\beta$ T cells (37,38), one might expect to find overall fewer contacts with DC. However, in mice lacking CD8 which contain $\alpha\beta$ T cells, the $\gamma\delta$ T cells were found in proximity of what appeared to be endogenous DEC-205⁺DC in the PALS, and they were not prevented from contacts with transferred splenic CD8⁺ non-T cells which probably are CD8⁺DC (Fig.3k). Contacts with endogenous splenic CD8⁺ non-T cells seem to occur continuously as there was little change in frequency following OVA-sensitization/ challenge, and both $V\gamma 1^+$ cells and $V\gamma 4^+$ cells take part. Likewise, enriched CD8⁺ DC can restore the development of both $V\gamma 1^+$ AHR-enhancers and $V\gamma 4^+$ AHR-suppressors. Our preliminary data (not shown) suggest that the transfer of CD8⁺ DC drives the $\gamma\delta$ T cells into cell cycle (C.L.R. et al., unpublished). However, the CD8⁺ DC appear to exert their effect at an early stage in the development of the AHR-modulating $\gamma\delta$ T cells, and these interactions alone seem insufficient to explain the functional differentiation of the AHR-enhancers and suppressors. Notably, development of the suppressors requires airway challenge, and thus may depend on a lung-derived signal (11,15).

Our data suggest a strict requirement for CD8 in the development of the AHR-regulatory $\gamma\delta$ T cells although these cells themselves need not express CD8. A role for CD8 "in trans" in $\gamma\delta$ T cell development has not been reported (39). Since transferred enriched CD8⁺ DC can restore development of the AHR-modulators in the absence of other CD8⁺ cells in the recipients, it seems likely that the DC-expressed CD8 satisfies the requirement for CD8 although other more complex scenarios cannot be excluded. A direct mechanism would be interesting because, although CD8⁺ DC in the lymphoid tissues have been characterized as a phenotypically and functionally distinct population (40), no functional significance has been assigned to the CD8 they express. In fact, in one study comparing the largely overlapping populations of CD8⁺ and DEC-205⁺ DC for their ability to support $\alpha\beta$ T cell-responses, DC-expressed CD8 appeared to be of no consequence as $CD8^+$ DC could be replaced with DEC-205⁺ DC without change (41). Our findings contrast with this study, inasmuch as endogenous DEC- 205^+ cells, which are plentiful in the spleen of CD8⁻ mice and co-localize here with $\gamma\delta$ T cells (Fig. 3j), failed to support the development of the AHR-modulating $\gamma\delta$ T cells. Thus, DC-expressed CD8 α might have a function after all, in connection with the development of $\gamma\delta$ T cells, and perhaps as an activating receptor (42, 43).

CD8⁺ DC, which reside in the lymphoid tissues(28,29), have been found to receive Ag and activating signals from phagocytes that shuttle from peripheral tissues such as skin and lung to the secondary lymphoid organs (44,45), to excel at Ag-cross-presentation, and to prime naïve $\alpha\beta$ T cells to become Ag-specific CTLs and type 1 helper (Th1) cells (46-48). To our knowledge, they have not previously been connected with $\gamma\delta$ T cells. However, a dependence on CD8⁺ DC aligns the $\gamma\delta$ T cells with the above-mentioned types of $\alpha\beta$ T cells (CD8⁺ and Th1), and sets them apart from others (Th2), and from NK cells, neither of which depend on CD8⁺ DC although they require DC-priming (25,49). We have not yet determined which factors might be involved in the DC- $\gamma\delta$ T interaction. Candidates include TNF- α and IL-15,

both of which are produced by CD8⁺ DC, and TNF- α , because the function and development of AHR-regulatory $\gamma\delta$ T cells depends on it (50,51), and IL-15, because it supports homeostatic expansion of $\gamma\delta$ T cells (37). Incidentally, IL-15 is "trans"-presented by DC, requires DC-lymphocyte contact, and also is essential in the "priming" of NK cells (49).

Connecting the observations reported in this study, we propose that developing $\gamma\delta$ T cells must interact with CD8⁺DC in the lymphoid tissues as an early step on their way to functional competence (e.g. towards becoming AHR-modulators). Because this interaction appears to involve direct cell-contact, because CD8 is absolutely required for $\gamma\delta$ T cell-development, and because development can take place when the only available CD8 appears to be expressed by the DC, we envisage that the DC-expressed CD8-molecules might engage ligands on the $\gamma\delta$ T cells they contact. Known ligands for CD8 are MHC class I or class I-like molecules (12,52). We have not yet examined AHR-enhancement, but we found that AHR-suppression by V γ 4⁺ $\gamma\delta$ T cells indeed depends on MHC class I expression (Cook et al., unpublished). Thus, our data are consistent with the scenario outlined, but fall short of proving it. However, in another study with naturally occurring Treg whose functional activation requires both CD8 and MHC I, a similar cell-cell interaction was proposed (53).

The interaction between $\gamma\delta$ T cells and CD8⁺DC documented here likely represents only one of several steps in the development of the $\gamma\delta$ T cells. As both AHR-suppressive and AHR-enhancing types depend on CD8⁺DC, their functional differences might be determined elsewhere, earlier or later in their development (54). Additional accessory cells such as the CD8⁻ phagocytes shown in Fig. 31, which engulf OVA/alum particles and appear following sensitization in the splenic PALS near the CD8⁺DC and the $\gamma\delta$ T cells, are perhaps involved as well (44,45). These cells or cells from the challenged lung might mediate the functional differentiation of the splenic AHR-suppressors, which require induction by OVA-sensitization and challenge (15).

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Figure 1.

AHR in mice after systemic sensitization and airway challenge with OVA (2ip3N protocol). (**a-d**) Lung resistance (R_L) and dynamic compliance (C_{dyn}) as percent changes from saline controls in relation to increasing doses of aerosolized MCh. Treatments: none (NT), intraperitoneal sensitization and airway challenge (2ip3N), 2ip3N plus hamster IgG i.v., 2ip3N plus anti Vγ4 mAb UC3 i.v., 2ip3N plus purified Vγ4⁺ cells from 2ip3N-treated donors (wt or CD8 $\alpha^{-/-}$) i.v. (**a**) Normal C57BL/6 mice after 2ip3N or 2ip3N plus depletion of Vγ4⁺ cells with mAb UC3 (* non-treated compared with 2ip3N, # 2ip3N compared with 2ip3N plus UC3); (**b**) B6.CD8 $\alpha^{-/-}$ mice after 2ip3N plus depletion of Vγ4⁺ cells with mAb UC3 (* non-treated compared with 2ip3N); (**c**) B6.TCR-Vγ4/6^{-/-} mice after 2ip3N or 2ip3N plus Vγ4⁺ cells from B6.TCR- $\beta^{-/-}$ (CD8 α^{wt}) donors (* non-treated compared with 2ip3N; there was no significant difference between non-treated and 2ip3N plus WT Vγ4 cells); (**d**) B6.TCR-Vγ4/6^{-/-} mice after 2ip3N or 2ip3N plus Vγ4⁺ cells from B6.CD8 $\alpha^{-/-}$ donors (there was no significant difference between 2ip3N and 2ip3N plus CD8^{-/-} Vγ4 cells at any dose of MCh).



Figure 2.

AHR in $\gamma\delta$ T cell-deficient recipients reconstituted with CD8-selected V $\gamma4^+$ cells, after systemic sensitization and airway challenge with OVA (2ip3N protocol). (**a**,**b**) Lung resistance (R_L) and dynamic compliance (C_{dyn}) as percent changes from saline controls in relation to increasing doses of aerosolized MCh. Recipients: B6.TCR-V $\gamma4/6^{-/-}$ mice; donors: 2ip3N-treated B6.TCR- $\beta^{-/-}$ or C57BL/6 mice. Treatments: none (NT), systemic sensitization and airway challenge (2ip3N), 2ip3N plus selected purified V $\gamma4^+$ /CD8 β^+ or $^-$ or V $\gamma4^+$ / CD8 α^+ or $^-$ cells from the donors. (**a**) Recipients untreated or after 2ip3N or 2ip3N plus selected V $\gamma4^+$ /CD8 β^+ or $^-$ cells from B6.TCR- $\beta^{-/-}$ donors (* non-treated compared with 2ip3N); (**b**) Recipients untreated or after 2ip3N or 2ip3N plus selected V $\gamma4^+$ /CD8 α^+ or $^-$ cells from C57BL/6 donors (* non-treated compared with 2ip3N); (**b**) Recipients untreated or after 2ip3N or 2ip3N plus Selected V $\gamma4^+$ /CD8 α^+ or $^-$ cells from C57BL/6 donors (* non-treated compared with 2ip3N); (**b**) Recipients untreated or after 2ip3N or 2ip3N plus Selected V $\gamma4^+$ /CD8 α^+ or $^-$ cells from C57BL/6 donors (* non-treated compared with 2ip3N) plus CD8 α^+ V $\gamma4$ cells, + 2ip3N compared with 2ip3N plus CD8 α^- V $\gamma4$ cells).



Figure 3.

Localization of $\gamma\delta$ T cells, CD8 α^+ DCs and CD8 α^- OVA/alum-containing phagocytes in the spleen (**a-l**) Frozen sections of spleen stained with mAbs, tissue autofluorescence (green), antibody staining (red or blue), coincidence of antibody staining (pink/purple), (**a**) C57BL/6 spleen, stained with anti TCR- δ mAb (red) and anti DEC-205 mAb (blue); (**b**) B6.TCR- $\beta^{-/-}$ spleen, stained with anti TCR- δ mAb (red) plus anti CD8 α mAb (blue); (**c**) B6.TCR- $\beta^{-/-}$ spleen, stained with anti TCR- δ mAb (red) plus anti CD8 α mAb (blue), or with (**d**) anti TCR- δ v γ 1 mAb (red) plus anti CD8 α mAb (blue); (**e**) B6.TCR- $\beta^{-/-}$ spleen, stained with anti CD11c mAb (red) plus anti CD8 α mAb (blue), cells co-expressing these markers appear violet; (**f**) B6.CD8 $\alpha^{-/-}$ spleen, stained with anti TCR- δ mAb (red); (**g**) B6.TCR- $\beta^{-/-}\delta^{-/-}$ spleen, stained with anti CD12 α mAb (blue), cells co-expressing these markers appear violet; (**f**)

with anti CD8 α mAb; (h) B6.TCR- $\beta^{-/-}$ spleen, stained with an antibody against CD103 (red) plus anti CD8 α mAb (blue), cells co-expressing these markers appear violet; (i) B6.TCR- $\beta^{-/-}$ spleen, stained with an antibody against DEC-205 (red) plus anti CD8 α mAb 53–6/7 (blue), cells co-expressing these markers appear violet; (j) B6.CD8 $\alpha^{-/-}$ spleen, stained with anti TCR- δ mAb (red) plus anti DEC-205 mAb (blue); (k) 2ip-treated B6.CD8 $\alpha^{-/-}$ spleen after transfer of CD8⁺ cells from the spleen of a 2ip-treated B6.TCR- $\beta^{-/-}\delta^{-/-}$ mouse, stained with anti TCR- δ mAb (red) plus anti CD8 α mAb (blue), enlarged detail showing two endogenous $\gamma\delta$ T cells in contact with transferred CD8⁺ DC; (l) same tissue and staining as in (k), additional detail: OVA/alum appears as bright white/yellow fluorescence in CD8 α^{-} , TCR- δ^{-} phagocytes, which are found in close proximity to both a $\gamma\delta$ T cell and a CD8⁺ DC.

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Figure 4.

Distribution of contacts between $\gamma\delta$ T cells and CD8⁺ DCs in the spleen (**a**-**d**) Frozen sections of spleen stained with mAbs were evaluated for relative frequencies of $\gamma\delta$ T cell-DC contacts (**a**) B6.TCR- $\beta^{-/-}$ spleen non-treated and 2ip3N, fractions of TCR- δ^+ , V γ 1⁺ and V γ 4⁺ cells in contact with CD8⁺ non-T cells; (**b**) B6.TCR- $\beta^{-/-}$ spleen non-treated and 2ip3N, comparison of PALS and "outside the PALS", fractions of TCR- δ^+ cells in contact with CD8⁺ non-T cells. (**c**) B6.TCR- $\beta^{-/-}$ spleen non-treated and 2ip3N, fractions of CD8⁺ non-T cells in contact with TCR- δ^+ , V γ 1⁺ and V γ 4⁺ cells; (**d**) B6.TCR- $\beta^{-/-}$ spleen non-treated and 2ip3N, comparison of PALS and "outside the PALS", fractions of CD8⁺ non-T cells in contact with TCR- δ^+ , V γ 1⁺ and V γ 4⁺ cells; (**d**) B6.TCR- $\beta^{-/-}$ spleen non-treated and 2ip3N, comparison of PALS and "outside the PALS", fractions of CD8⁺ non-T cells in contact with TCR- δ^+ cells.

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Figure 5.

AHR in $\gamma\delta$ T cell-deficient recipients reconstituted with genetically CD8-deficient V $\gamma4^+$ cells from "CD8-restored donors", after intra-peritoneal sensitization and airway challenge with OVA (2ip3N protocol). (**a-d**) Lung resistance (R_L) and dynamic compliance (C_{dyn}) as percent changes from saline controls in relation to increasing doses of aerosolized MCh. Recipients: B6.TCR-V $\gamma4/6^{-/-}$ mice; donors: B6.CD8 $\alpha^{-/-}$ mice. Treatments: none (NT), systemic sensitization and airway challenge (2ip3N), 2ip3N plus V $\gamma4^+$ cells from 2ip3N-treated donors. Donors: The donors of the V $\gamma4^+$ cells either remained non-restored or were restored with splenocytes from 2ip-treated B6.TCR- $\beta^{-/-}/\delta^{-/-}$ mice, including the following fractions: (**a**) whole spleen, (**b**) magnetic bead-selected CD8 α^+ or CD8 α^- splenocytes, or (**c**) DC-enriched, magnetic bead-selected CD8 α^+ T cells from 2ip-treated B6.TCR- $\delta^{-/-}$ mice.

Recipients: (a) untreated or after 2ip3N plus V γ 4⁺ cells from non-restored donors or donors that were restored with whole spleen (* 2ip3N plus V γ 4 cells compared with 2ip3N plus V γ 4 cells from donor restored with whole spleen, # non-treated compared with 2ip3N plus V γ 4 cells from non-restored donor); (b) untreated or after 2ip3N plus V γ 4⁺ cells from 2ip3N-treated donors restored with CD8-enriched or –depleted splenocytes (* non-treated compared with 2ip3N plus V γ 4 cells from donors reconstituted with CD8⁻ splenocytes, # 2ip3N plus V γ 4 cells from donors reconstituted with CD8⁻ splenocytes compared with 2ip3N plus V γ 4 cells from donor reconstituted with CD8⁺ splenocytes); (c) untreated or after 2ip3N plus V γ 4 cells from donor reconstituted with CD8⁺ splenocytes); (c) untreated or after 2ip3N plus V γ 4 cells from donor reconstituted with CD8⁺ splenocytes); (c) untreated or after 2ip3N plus V γ 4 cells from donor reconstituted with CD8⁺ splenocytes); (c) untreated or after 2ip3N plus V γ 4 cells from donor reconstituted with CD8⁺ splenocytes); (c) untreated or after 2ip3N plus V γ 4 cells from donor reconstituted with CD8⁺ splenocytes); (c) untreated or after 2ip3N plus V γ 4 cells from donor reconstituted with CD8⁺ splenocytes); (c) untreated or after 2ip3N plus V γ 4 cells from donor reconstituted with CD8⁺ splenocytes); (c) untreated or after 2ip3N plus V γ 4 cells from donor reconstituted with CD8⁺ splenocytes); (c) untreated or after 2ip3N plus V γ 4⁺ cells from donor reconstituted with CD8⁺ splenocytes); (c) untreated or after 2ip3N plus V γ 4 cells from donor splenocytes); (c) untreated or after 2ip3N plus V γ 4⁺ cells from donor splenocytes); (c) untreated or after 2ip3N plus V γ 4⁺ cells from donor splenocytes); (c) untreated or after 2ip3N plus V γ 4⁺ cells from donor splenocytes); (c) untreated or after 2ip3N plus V γ 4⁺ cells from donor splenocytes); (c) untreated or after 2ip3N plus V γ 4⁺ cells from donor splenocytes); (c) untreated or after

donors restored with CD8⁺ or CD8⁻ splenic DCs or with non-DCs (mostly CD8⁻) (* nontreated compared with 2ip3N plus V γ 4 cells from donor restored with non-DCs or # with CD8⁻ DCs, \$ 2ip3N plus V γ 4 cells from donor restored with CD8⁺ DCs compared with 2ip3N plus V γ 4 cells from donor restored with non-DCs or # with CD8⁻ DCs; (**d**) untreated or after 2ip3N or 2ip3N plus V γ 4⁺ cells from donors restored with CD8 α ⁺ T cells (* non-treated compared with 2ip3N, # non-treated compared with 2ip3N plus V γ 4 cells from donors reconstituted with CD8 α ⁺ T cells, + there was no significant difference in R_L between 2ip3N and 2ip3N plus V γ 4 cells from donors reconstituted with CD8 α ⁺ T cells, with exception of the response to 12.5 mg MCh).

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Figure 6.

AHR in $\gamma\delta$ T cell-deficient recipients reconstituted with V γ 1⁺ cells from normal or CD8deficient mice, or with CD8-deficient V γ 1⁺ cells from "CD8-restored donors", after systemic sensitization and airway challenge with OVA (2ip3N protocol). (**a-c**) Lung resistance (R_L) and dynamic compliance (C_{dyn}) as percent changes from saline controls in relation to increasing doses of aerosolized MCh. Recipients: B6.TCR- $\delta^{-/-}$ mice; donors: C57BL/6 or B6.CD8 $\alpha^{-/-}$ mice. Treatment of recipients: none (NT), intra-peritoneal sensitization and airway challenge (2ip3N), 2ip3N plus selected purified V γ 1⁺ cells from non-treated donors, 2ip3N plus V γ 1⁺ cells from non-treated B6.CD8 $\alpha^{-/-}$ mice restored with with magnetic bead-selected CD8 α^+ or CD8 α^- splenocytes from 2ip-treated B6.TCR- $\beta^{-/-}/\delta^{-/-}$ mice, or 2ip3N plus V γ 1⁺ cells from non-treated B6.CD8 $\alpha^{-/-}$ mice restored with DC-enriched, magnetic bead-selected CD8 α^+ or CD8 α^- DCs from 2ip-treated B6.TCR- $\beta^{-/-}/\delta^{-/-}$ mice.

Recipients: (a) after 2ip3N or 2ip3N plus V γ 1⁺ cells from C57BL/6 or from B6.CD8 $\alpha^{-/-}$ donors (* 2ip3N compared with 2ip3N plus V γ 1⁺ cells from C57BL/6 donors, # 2ip3N plus V γ 1⁺ cells from C57BL/6 donors (* 2ip3N plus V γ 1⁺ cells from C57BL/6 donors); (b)

after 2ip3N or 2ip3N plus V γ 1⁺ cells from B6.CD8 $\alpha^{-/-}$ donors restored with CD8⁺ or ⁻ spleen cells from non-treated B6.TCR- $\beta^{-/-}/\delta^{-/-}$ mice (* 2ip3N compared with 2ip3N plus V γ 1⁺ cells from B6.CD8 $\alpha^{-/-}$ donors restored with CD8⁺ spleen cells, # 2ip3N plus V γ 1⁺ cells from B6.CD8 $\alpha^{-/-}$ donors restored with CD8⁺ spleen cells compared with 2ip3N plus V γ 1⁺ cells from B6.CD8 $\alpha^{-/-}$ donors restored with CD8⁺ spleen cells); (c) after 2ip3N or 2ip3N plus V γ 1⁺ cells from non-treated B6.CD8 $\alpha^{-/-}$ donors restored with CD8⁻ spleen cells); (c) after 2ip3N or 2ip3N plus V γ 1⁺ cells from non-treated B6.CD8 $\alpha^{-/-}$ donors restored with CD8⁺ spleen cells); (c) after 2ip3N or 2ip3N plus V γ 1⁺ cells from non-treated B6.CD8 $\alpha^{-/-}$ donors restored with CD8⁺ bcs, # 2ip3N compared with 2ip3N plus V γ 1⁺ cells from B6.CD8 $\alpha^{-/-}$ donors restored with CD8⁺ DCs, # 2ip3N plus V γ 1⁺ cells from B6.CD8 $\alpha^{-/-}$ donors restored with 2ip3N plus V γ 1⁺ cells from B6.CD8 $\alpha^{-/-}$ donors restored with CD8⁺ DCs, # 2ip3N plus V γ 1⁺ cells from B6.CD8 $\alpha^{-/-}$ donors restored with CD8⁺ DCs, # 2ip3N plus V γ 1⁺ cells from B6.CD8 $\alpha^{-/-}$ donors restored with CD8⁺ DCs, # 2ip3N plus V γ 1⁺ cells from B6.CD8 $\alpha^{-/-}$ donors restored with CD8⁺ DCs.