T Cell-Intrinsic and -Extrinsic Contributions of the IFNAR/STAT1-Axis to Thymocyte Survival

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Abstract

STAT1 is an essential part of interferon signaling, and STAT1-deficiency results in heightened susceptibility to infections or autoimmunity in both mice and humans. Here we report that mice lacking the IFNα/β-receptor (IFNAR1) or STAT1 display impaired deletion of autoreactive CD4+CD8−-T-cells. Strikingly, co-existence of WT T cells restored thymic elimination of self-reactive STAT1-deficient CD4+CD8−-T cells. Analysis of STAT1-deficient thymocytes further revealed reduced Bim expression, which was restored in the presence of WT T cells. These results indicate that type I interferons and STAT1 play an important role in the survival of MHC class I-restricted T cells in a T cell intrinsic and non-cell intrinsic manner that involves regulation of Bim expression through feedback provided by mature STAT1-competent T cells.

Citation: Moro H, Otero DC, Tanabe Y, David M (2011) T Cell-Intrinsic and -Extrinsic Contributions of the IFNAR/STAT1-Axis to Thymocyte Survival. PLoS ONE 6(9): e24972. doi:10.1371/journal.pone.0024972

Editor: Jose Alberola-Illa, Oklahoma Medical Research Foundation, United States of America
Received June 17, 2011; Accepted August 19, 2011; Published September 20, 2011

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Funding: This work was supported by grants from the National Institute of Allergy and Infectious Diseases (AI71233) and The National Multiple Sclerosis Society (RG3948) to MD. The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript.

Competing Interests: The authors have declared that no competing interests exist.
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Introduction

Selection of T lymphocytes expressing antigen receptors that are major histocompatibility complex (MHC)-restricted but tolerant to self-antigens is a fundamental requirement for infection immunity. The relatively small T cell subset of CD4+CD8−-double negative thymocytes gives rise to the CD4+CD8−-double positive thymocytes which account for the majority of T cells in the thymus. Subsequently, CD4+CD8−-thymocytes mature into CD4+ or CD8− single positive T cells via a dual selection process. Positive selection occurs if the TCR of immature thymocytes recognizes the MHC on thymic epithelial cells with sufficient affinity to elicit the transduction of survival and differentiation signals. The result is MHC restriction and eventually the development of single positive mature T cells. However, if the TCRs of positively selected T cells subsequently engage in high affinity interactions with the MHC/peptide complex on stromal cells such as dendritic cells and macrophages, these T cells are deleted from the pool via programmed cell death. Thereby, the process of negative selection eliminates self-reactive thymocytes and facilitates tolerance to self-antigens. Collectively, the outcomes of these two selection processes are required to generate a T cell repertoire that is both self-MHC restricted and self-tolerant, a process that appears to be determined by the avidity between the MHC/peptide complex and the TCR [1,2,3,4].

The signal transducer and activator of transcription 1 (STAT1) is a well-characterized component of the type I (IFNα/β) or type II (IFNγ) interferon-induced signaling pathways [5,6,7]. The physiological importance of STAT1 in-vivo has been made possible through the generation of STAT1-deficient mice by two independent groups [8,9]. As anticipated, STAT1-deficient mice failed to respond to either type I or type II interferons, however, comparison of the individual T cell subsets between wild-type (WT) and STAT1−/− 129Sv/Ev mice provided no evidence for significant differences between these two strains [8,9,10].

Even though all IFNs utilize STAT1 as a signaling mediator, type I and II IFNs exert opposing effects on the progression of the demyelinating, T cell-mediated autoimmune disease multiple sclerosis (MS) [11]. These juxta-posed effects of type I and II IFNs in MS raise the question as to what role STAT1 plays in this pathological process. To address this issue, we had previously employed mice carrying a transgenic T cell receptor specific for myelin basic protein (TCRMBP). Upon T cell activation, TCRMBP-transgenic animals develop experimental autoimmune encephalomyelitis (EAE) which serves as a murine model for MS [12,13]. Analysis of TCRMBP−/− STAT1-deficient mice revealed a dramatically increased susceptibility to EAE development, accounted for at least in part by a defect in the development and functionality of CD4+CD25− regulatory T cells [10]. However, the severity and frequency of spontaneous EAE development in the absence of STAT1 lead us to hypothesize that STAT1 might also contribute to the events that govern the elimination of autoreactive T cells via negative selection.

Therefore, to explore a possible role for STAT1 in thymic selection, we employed mice carrying a transgenic TCR which recognizes the male-specific HY antigen (designated TCRHY) in the context of H-2Db, a widely used model system to evaluate thymic selection events [14,15,16]. In male animals that harbor the HY antigen, most thymocytes are recognized as self-reactive and deleted from the T cell pool, resulting in the absence of CD4+CD8−-double positive cells and a severe reduction in thymic cellularity [14,16].

The findings presented in this study demonstrate that STAT1, as well as type I interferon, is required for the removal of
autoreactive T cells in a non-cell intrinsic manner, as TCRHY-STAT1–/–, TCRHYIFNAR–/– and TCRHYIFNGR–/– animals revealed striking differences in T cell subsets compared to WT littermates. In the model system we employed, the function of STAT1 in T cell apoptosis correlates directly with Bim expression in CD4+CD8+ double-positive cells. As such, our findings support the notion that T cell - T cell interactions, either through direct cell-to-cell contact or via soluble mediators, are essential for adequate T cell development.

Results

Impaired elimination of autoreactive T cells in the absence of IFNAR and STAT1

In order to investigate a possible function for STAT1 in thymic selection events, STAT1-deficient mice were crossed with TCRHY transgenic animals yielding TCRHYSTAT1–/– animals. Thymocytes were isolated from TCRHY and TCRHYSTAT1–/– transgenic male animals and analyzed by flow cytometry. As previously reported [14,16], male TCRHY mice almost completely lack CD4+CD8+ double positive thymocytes in their thymi due to efficient deletion of these autoreactive T cells (Fig. 1) when compared to non-transgenic WT animals or female TCRHY mice (not shown). Strikingly, the absence of STAT1 resulted in a drastic impairment of this elimination process such that a substantial number of CD4+CD8+ double positive and CD8+ single positive thymocytes were detectable in male TCRHYSTAT1–/– mice (Fig. 1a, 2nd panel). In contrast, deletion of CD4+CD8+ double positive cells in male TCRHYSTAT1+/+ mice (Fig. 1a, 1st panel) or TCRHYSTAT1–/– (not shown) littermates occurred with similar efficiency as observed in TCRHY mice.

As STAT1 functions downstream of the IFNα/β receptor as well as the IFNγ receptor (IFNGR), we decided to explore which if any of these cytokines were accountable for the impaired elimination of the autoreactive T cells. Analysis of the T cell subsets from TCRHYIFNAR–/– and TCRHYIFNGR–/– animals revealed that only TCRHYIFNAR–/– but not TCRHYIFNGR–/– mice display a similar defect in the deletion of autoreactive T cells as TCRHYSTAT1–/– mice (Fig. 1a, 3rd and 4th panel). Thus, the altered distribution of T cell subsets as a consequence of impaired deletion appeared identical in TCRHYSTAT1–/– and TCRHYIFNAR–/– mice, whereas thymic populations in TCRHYIFNGR–/– mice resembled those in TCRHY animals (averages of several mice are enumerated in Fig. 1b). In addition, total numbers of CD8+ cells were increased in the thymi of TCRHYSTAT1–/– and TCRHYIFNAR–/– mice (Fig. 1c, and data not shown), resulting in the concomitant increased presence of TCRHY+ positive cells in the spleen, lymph node, and blood (not shown). To exclude the possibility that the observed differences were due to variations in the expression of the antigen receptor, we analyzed the levels of TCRHY+ cells in the different strains and found no significant difference between TCRHY and TCRHYSTAT1–/– mice (Fig. S1), or between the interferon receptor-deficient animals (not shown). Thus, the presence of TCRHY+CD8+ T cells in the thymi of male mice indicates that STAT1–/– thymocytes are failing to be eliminated.

Impaired deletion of TCRHYSTAT1–/– thymocytes correlates with the presence of STAT1 in hematopoietic cells

Lee et al. previously reported a cell-intrinsic role for STAT1 in the constitutive expression of MHC class I on lymphocytes [17]. Furthermore, an interferon-independent requirement for STAT1

![Figure 1. STAT1 and IFNAR are required for elimination of self-reactive TCRHY thymocytes.](https://doi.org/10.1371/journal.pone.0024972.g001)
in MHC class I expression was found specifically in T cells, such that thymocytes and peripheral T cells from STAT1-deficient mice displayed a more pronounced reduction of MHC class I than those derived from IFNAR/IFNGR double-deficient mice [17]. These findings raise the possibility that potentially reduced expression levels of MHC class I on thymic stromal cells is responsible for the attenuated deletion of autoreactive T cells in the absence of STAT1.

We therefore analyzed the surface expression levels of MHC class I in syngeneic WT, STAT1<sup>−/−</sup>, and IFNAR<sup>−/−</sup> mice. In conformity with Lee et al., we found a dramatic reduction in the surface expression of MHC class I in STAT1<sup>−/−</sup>, but not IFNAR<sup>−/−</sup> thymic T cells compared to WT cells (Fig. S2, left column panels). In striking contrast to the lymphocytes, however, thymic stromal cells including dendritic cells (DCs) and thymic epithelial cells (TECs) derived from either STAT1<sup>−/−</sup> and IFNAR<sup>−/−</sup> mice showed no significant reduction in MHC class I expression levels (Fig. S2, middle and right column panels).

Due to the fact that thymic stromal cells display normal MHC class I levels we considered that STAT1 might be required in hematopoietic cells rather than stromal cells for TCR<sup>HV</sup>-mediated deletion. To test this hypothesis, we lethally irradiated male WT mice prior to transfer of bone marrow (BM) derived from TCR<sup>HV</sup> or TCR<sup>HV</sup>STAT1<sup>−/−</sup> female mice (Fig. 2a, left panels). Conversely, TCR<sup>HV</sup>-derived BM was also transferred into lethally irradiated non-transgenic STAT1<sup>−/−</sup> male animals (Fig. 2a, right panels). In such mice, radiation-resistant TECs were derived from the host animal, whereas hematopoietic cells such as T cells, dendritic cells, or macrophages originate from the transferred BM. After four weeks, flow-cytometric analysis of the thymic T cell populations was performed.

As shown in Fig. 2a, adoptive transfer of TCR<sup>HV</sup> BM into irradiated male WT mice resulted in a similarly efficient deletion of the autoreactive donor T cells as was previously seen in the TCR<sup>HV</sup> donor animals. However, when TCR<sup>HV</sup>STAT1<sup>−/−</sup> BM was used to reconstitute irradiated WT animals, autoreactive T cell deletion was diminished to the same extent as was observed in TCR<sup>HV</sup>STAT1<sup>−/−</sup> mice (Fig. 2a, right panels). Consistent with this observation, adoptive transfer of TCR<sup>HV</sup> BM into irradiated male STAT1<sup>−/−</sup> mice yielded animals with intact T cell elimination (Fig. 2b, left panels), whereas the use of TCR<sup>HV</sup>-STAT1<sup>−/−</sup> mice as BM donors led to severely attenuated elimination of the transferred autoreactive T cells in the host animals (Fig. 2b, right panels).

Collectively, these results clearly demonstrate that elimination of TCR<sup>HV</sup> T cells in male mice is dependent on the STAT1 status of hematopoietic cells, but is independent of the presence of STAT1 in TECs.

To explore whether the observed defects in apoptosis of CD4<sup>+</sup>CD8<sup>+</sup> double positive thymocytes is linked to the presence of the TCR<sup>HV</sup>, we decided to also evaluate the behavior of this T cell subset in response to different stimulations. The CD4<sup>+</sup>CD8<sup>+</sup> double positive population of thymocytes is sensitive to cross-linking of the T cell receptor with anti-CD3 antibody and this approach has also been widely used as an alternative strategy to study molecular mechanisms involved in negative selection in vitro as well as in vivo. When 129Sv/Ev WT and STAT1<sup>−/−</sup> animals were injected with anti-CD3 antibody to induce programmed cell death of thymocytes, STAT1<sup>−/−</sup> cells showed significantly lower sensitivity to the in-vivo anti-CD3 treatment compared to WT thymocytes (Fig. 3a). To determine if there is an intrinsic defect in TCR-induced cell death of STAT1-deficient thymocytes, double-positive cells from non-transgenic 129Sv/Ev and their STAT1<sup>−/−</sup> counterparts were stimulated with plate-bound anti-CD3 antibody and apoptosis was measured after 24 hours. Compared to WT thymocytes, double-positive cells derived from STAT1<sup>−/−</sup>-deficient animals displayed significantly reduced apoptosis following T cell receptor crosslinking (Fig. 3b). STAT1<sup>−/−</sup> T cells are still capable of transducing signals via the TCR as documented by the – albeit somewhat reduced – antigen-induced upregulation of the activation marker CD69 (Fig. 3c). Even though anti-CD3 induced death of double-positive thymocytes in vivo is thought to

![Figure 2. STAT1-deficient T cells fail to undergo deletion in the presence of wild type TEC.](image-url)
take place through cytokine secretion by activated CD4+ T cells rather than by direct TCR ligation on double-positive thymocytes, the in-vitro experiments using purified double-positive T cells demonstrate that STAT1-deficient thymocytes are indeed less susceptible to programmed cells death via TCR engagement. This difference was specific to TCR-induced apoptosis as STAT1-deficient thymocytes were still sensitive to dexamethasone induced cell death in vitro and in vivo (Fig. 3b, and data not shown). Collectively, our results obtained from the adoptive transfer studies and anti-CD3 challenges strongly suggest an intrinsic inability of STAT1−/− thymocytes to undergo deletion triggered by TCR-mediated signals.

Deletion of DP thymocytes requires mature T cell feedback

Our findings suggested that the purge of CD4+CD8+ TCRHY T cells in male mice is linked to the STAT1 status of hematopoietic cells, but the question remained whether this requirement for STAT1 was intrinsic to the autoreactive T cells. We reasoned that if STAT1 function during selection events is not T cell intrinsic, impaired deletion of TCRHY,STAT1−/−,CD4+CD8+ thymocytes might be restored by providing STAT1-expressing hematopoietic cells in trans. We therefore performed RAG-KO complementation assays using mixed BMs in which donor-derived thymocytes represent a mixture of cells that arose from the combined transfer of WT and TCRHY,STAT1−/− BMs. Briefly, BM cells derived from the two distinct female donor mice were mixed in a 1:1 ratio, and injected into lethally irradiated male recipients. To avoid possible contamination of developing donor T cells with residual recipient thymocytes that survived the irradiation, we used male RAG−/− mice as recipients. After 4 weeks, the extent of deletion of TCRHY-carrying T cells derived from the transferred BMs was determined by flow cytometry.

As shown in Fig. 4a, when the combined BMs of WT and TCRHY mice were introduced into the irradiated male RAG−/− animals, the elimination of the resulting TCRHY T cells occurred as efficiently as in the original TCRHY mice (left panel). Likewise, transfer of a BM mix consisting of STAT1−/− and TCRHY,STAT1−/− into the irradiated male RAG−/− animals reproduced the defect of the TCRHY,STAT1−/−,CD4+CD8+ thymocytes observed in TCRHY,STAT1−/− mice (Fig. 4a, 2nd panel). However, when the combined BMs of 129WT and TCRHY,STAT1−/− were transferred into the irradiated male RAG−/− recipients, the presence of cells derived from the 129WT BM facilitated the adequate elimination of the autoreactive TCRHY T cells (Fig. 4a, 3rd panel). This result indicates that the presence of STAT1-containing hematopoietic cells is a prerequisite for the deletion of autoreactive TCRHY,STAT1−/− T cells which are intrinsically refractory to elimination.

Impaired T cell apoptosis in STAT1−/− mice correlates with reduced Bim expression

Eradication of self-reactive TCRHY,STAT1−/−,CD4+CD8+ thymocytes can be restored by the presence of STAT1-containing mature T cells, albeit the resistance of non-transgenic STAT1−/−
T cells to anti-CD3 induced apoptosis (Fig. 3) indicates that T cell intrinsic factors are involved in impaired apoptosis in the absence of STAT1. To determine which features are responsible for the impaired elimination of auto-reactive STAT1\(^2\)/\(^2\) thymocytes we chose to examine the levels of pro- and anti-apoptotic Bcl-2 family members in STAT1\(^2\)/\(^2\) thymocytes compared to 129WT thymocytes. The pro-apoptotic Bcl-2 family member Bim has been shown to be pivotal to thymocyte negative selection [19] and mice that lack Bim succumb to T cell mediated autoimmune disease [20]. When we analyzed Bim levels of total thymocytes following 0 or 6 hrs in culture with or without anti-CD3 stimulation, we found that Bim levels were strikingly reduced in thymocytes derived from STAT1\(^2\)/\(^2\) mice as compared to their wild-type counterparts (Fig. 5a). Splice variants of Bim (BimS and BimES) were still induced following TCR stimulation in the absence of STAT1 thymocytes, but maximal Bim levels in STAT1\(^2\)/\(^2\) thymocytes still remained well below those observed even in unstimulated WT cells (Fig. 5a, top panel). Bcl-2 levels were slightly lower in STAT1\(^2\)/\(^2\) thymocytes (Fig. 5a, 2nd panel from top), whereas no difference was detectable in the levels of the anti-apoptotic family member Mcl-1 (Fig. 5a, 3rd panel from top). As Mcl-1 is considered the direct antagonist to Bim to promote cell survival [21,22], it seemed reasonable to conclude that the survival of STAT1\(^2\)/\(^2\) CD4\(^+\)CD8\(^+\) thymocytes is promoted by normal Mcl-1 levels in the presence of reduced Bim expression. Consistent with this finding is the observation that STAT1\(^2\)/\(^2\) mice develop lymphoproliferative disease as evidenced by significantly enlarged spleens in aged STAT1\(^2\)/\(^2\) mice compared to two age-matched WT animals (Fig. S3).

Deletion of TCR\(^{HV}\)STAT1\(^2\)/\(^2\) CD4\(^+\)CD8\(^+\) thymocytes can be restored by the presence of STAT1-containing mature T cells, one would therefore anticipate that these WT T cells facilitate an increase in Bim expression in the TCR\(^{HV}\)STAT1\(^2\)/\(^2\) CD4\(^+\)CD8\(^+\) thymocytes. We thus analyzed Bim expression levels by intracellular FACS staining of TCR\(^{HV}\)STAT1\(^2\)/\(^2\) thymocytes obtained from mixed bone marrow chimeras as described in Fig. 4. Corroborating the results acquired from the non-transgenic animals, the expression levels of Bim in TCR\(^{HV}\) thymocytes derived from TCR\(^{HV}\)STAT1\(^2\)/\(^2\)/STAT1\(^2\)/\(^2\) chimeras were significantly lower than in those from TCR\(^{HV}\)/WT chimeric animals (Fig. 5b, top panel). Consistent with the effects on deletion of TCR\(^{HV}\)STAT1\(^2\)/\(^2\) CD4\(^+\)CD8\(^+\) thymocytes, the presence of WT, but not of TCR\(^2\)/\(^2\) BM-derived cells elevated Bim expression in TCR\(^{HV}\)STAT1\(^2\)/\(^2\) thymocytes (Fig. 5b, bottom panel). In summary, our findings demonstrate an intrinsic inability of auto-reactive STAT1\(^2\)/\(^2\) thymocytes to undergo deletion, an event that mandates Bim expression which in turn appears to be controlled by a STAT1-dependent, but cell-extrinsic process.

**Discussion**

Deletion occurs when the TCRs of CD4\(^+\)CD8\(^+\) thymocytes interact with a peptide-MHC complex with high affinity, leading to apoptotic cell death [1,3,4,23]. Our previous work had demonstrated that STAT1-deficiency dramatically increases the incidence of autoimmune disease [10]. In addition to decreased CD4\(^+\)CD25\(^+\) regulatory T cell function, it seemed plausible that the absence of STAT1 also leads to impaired deletion of auto-
reactive T cells in the thymus. Consequently, we decided to investigate whether STAT1 is associated with the central tolerance and to explore this hypothesis, we employed TCRHV transgenic mice defective in STAT1. The TCRHV is a low affinity receptor compared to other transgenic TCR model systems. The early expression of the TCRHV T cell receptor during thymic development (DN to DP stage) and the universal expression of the HY antigen have provoked questions regarding the immunological relevance of this model. Notwithstanding its limitations, the TCRHV is a widely utilized and effective model in the study of thymic events that control T cell selection processes. Our results revealed that STAT1, in addition to the type I IFN receptor, is indeed required for the removal of autoreactive CD8+ cells in the TCRHV model system (Fig. 1a).

IFNs and STAT1 are recognized to play an important role in modulating MHC class I expression [17], and the fact that TCRHV-mediated deletion was perturbed in the absence of both STAT1 and IFNAR1 (Fig. 1a) indicated the involvement of type I IFN signals. According to Lee and colleagues, STAT1 is required for the constitutive expression of MHC class I molecules on lymphocytes [17]. Indeed, we observed drastically reduced levels of MHC class I expression on STAT1+/− T cells, however, we did not find a significant decrease in H-2b expression in T cells derived from IFNAR−/− mice (Fig. S2). As deletion was impaired in both TCRHV/STAT1−/− and TCRHV/IFNAR−/− mice, where the former had reduced, but the latter displayed normal expression levels of MHC on their thymocytes (Fig. S2), no good correlation exists between the observed defects in deletion of STAT1−/− and IFNAR−/− autoreactive T cells, and lymphocytic H-2b expression. We were also unable to detect any substantial reduction in MHC class I expression levels on thymic stromal cells which are considered the key APCs mediating negative selection. These results imply that the expression levels of MHC class I on thymocytes of TCRHV/STAT1−/− and TCRHV/IFNAR−/− mice are unlikely to be (solely) responsible for the defects in T cell selection in these animals.

Most importantly, the coexistence of WT thymocytes restored the impaired deletion of TCRHV/STAT1−/− thymocytes in male BM chimeras (Fig. 4). However, BM of TCR−/− mice which lack mature T cells failed to support the elimination of the autoreactive TCRHV/STAT1−/− T cells, strongly suggesting that mature T cells contribute to the efficiency of the selection process. Two groups had independently demonstrated a role for T cell - T cell interactions in the positive selection of CD4+ T cells by generating transgenic mice expressing MHC class II on their thymocytes [24]. More recently, it was found that mature single-positive T cells recirculate to the thymus and contribute to the positive selection of thymocytes [25]. Thus, while it appears that positive selection can be induced by MHC expressing T cells, our finding that cell derived from β2M−/− bone marrow can still support deletion of TCRHV/STAT1−/− thymocytes in mixed BM chimeras suggest that T cell – T cell communication in support of negative selection occurs independent of MHC class II (Fig. 4). It is also important to remember that the donor thymocytes were derived from female mice that do not harbor the HY antigen and can consequently not display the HY peptide antigen in their MHC class II molecules. Therefore, not the MHC class I/antigen complex, but other signals provided by donor-derived T cells via type I IFNs signaling are required for the elimination of TCRHV thymocytes in non-cell intrinsic manner.

Bim, Bax and Bak play important roles in T cell development and homeostasis [19,20,26]. Bim in particular is important for negative selection through the T cell receptor promoting the removal of self-reactive T cells [19]. Considering that STAT1−/− double positive cells also displayed resistance towards AICD elicited by TCR activation via antibody-mediated CD3 crosslinking (Fig. 5), a possibility existed that STAT1 regulates pro-apoptotic factors. Strikingly, we found that expression of the pro-apoptotic factor Bim was significantly reduced at the protein level in thymocytes from STAT1−/− mice compared to WT mice (Fig. 5a). Bim expression in TCRHV-STAT1−/− T cells could be restored to WT levels in WT+TCRHV/STAT1−/− but not in TCRHV-STAT1−/− chimeric mice (Fig. 5b). Although it has recently been suggested that Bim does not play a role in negative selection, Bim was necessary for elimination of double positive cells in the cortex [27], the stage at which TCRHV T cells are deleted [28]. It is also worthwhile noting that the flow cytometric profiles of male TCRHV/STAT1−/− thymic cell populations (Fig. 1) closely resemble those of TCRHV/Bim−/− thymi [19]. In addition,
Materials and Methods

Animals

STAT1−/− [9], IFNAR−/− [30], IFNγR−/− [31], TCR−/− [18] and TCRHY [16] mice have been described previously. RAG1−/− mice and β2M−/− mice were obtained from Jackson Laboratory (Bar Harbor, Maine). STAT1−/−, IFNAR−/− and IFNγR−/− mice were crossed with TCRHY transgenic mice yielding TCRHYSTAT1−/−, TCRHYIFNAR−/− and TCRHYIFNγR−/−, respectively. The expression of TCRHY was verified by FACS analysis. Animals were between 6 and 10 weeks of age at the time of the experiments. All mice used in these experiments were housed in a pathogen-free environment and were bred and cared for in accordance with University of California, San Diego Animal Care Facility regulations. All studies involving animal have been approved by the “The University of California San Diego Institutional Animal Care and Use Committee” (Protocol S02194).

Flow-cytometric analysis

For immunostaining, single cell suspensions were prepared from thymus, spleen or peripheral lymph nodes. After removal of red blood cells, approximately 10^6 cells were suspended in FACS buffer (PBS, 1% FCS, 0.02% NaN3) and stained for 20 min in the dark on ice. FITC-anti-CD4 (GK1.5), FITC-anti-B220 (RA3-6B2), PE-anti-CD62L (MEL-14), PE-anti-CD69 (H.12F3), PE/Cy7 or PE-anti-CD8 (53.6.7), APC-anti-CD3 (145-2C11) and biotin-anti-TCR (T3.70) were obtained from eBioscience (San Diego, CA). PE-anti-H-2Dk (KH95), biotin-anti-CD11c (HL3), biotin-anti-CD44 (IM7), biotin-anti-Rat IgG2a (RG7/1.30) and purified anti-Ep-CAM (G8.8) were purchased from BD Biosciences (San Jose, CA). APC-streptavidin was used as a secondary antibody. Histograms are derived from CD3 gated dendritic cells or B220 gated thymic epithelial cells, respectively. Dotted lines in the middle and lower panels represent MHC class I expression in WT cells, and the mean fluorescence intensity (MFI) is indicated. Representatives of at least three independent experiments are shown.

In vitro and In vivo cell death

Thymi were isolated from 129WT and STAT1−/− mice and single-cell suspensions prepared. Cells were plated at 1×10^6/ml in 24-well plates coated with 10 μg/ml anti-CD3 (eBiosciences) or 100 nM Dexamethasone (Sigma). Cells were harvested 24 hours following initiation of culture and apoptosis was measured by flow-cytometric analysis of sub-G0/G1 peaks after propidium iodide (PI) staining in 1 mM Tris (pH 8), 0.1% Triton, 0.1% sodium citrate, 0.1 EDTA, and 50 μg/ml PI. For in vivo studies, mice were injected intraperitoneally with either 20 μg anti-CD3 or Dexamethasone at 1 mg/kg and thymocyte populations were analyzed 48 hours later and compared to PBS treated mice. Double positive thymocytes from female TCRHY and TCRHYSTAT1−/− mice were incubated with increasing concentrations of smyc (738-746) peptide, AnaSpec (Fremont, CA), and analyzed 18 hours later for CD69 expression by flow cytometry.

Supporting Information

Figure S1 TCR expression on TCRHY and TCRHYSTAT1−/− thymocytes. Total thymocytes from TCRHY and TCRHYSTAT1−/− male mice were stained with monoclonal antibody T3.70 recognizing the Vβ8 chains of the TCRHY transgenic TCR. Dashed and solid lines represent TCRHY and TCRHYSTAT1−/− thymocytes, respectively. (TIF)

Figure S2 MHC class I expression in WT and STAT1−/− thymi. Flow cytometric analysis of thymocytes derived from WT, STAT1−/− and IFNAR−/− mice stained with MHC class I H-2^b-specific antibodies. Histograms are derived from CD3^+ gated T cells, CD11c^+ gated dendritic cells or B220^+ Ep-CAM^+ gated thymic epithelial cells, respectively. Dotted lines in the middle and lower panels represent MHC class I expression in WT cells, and the mean fluorescence intensity (MFI) is indicated. Representatives of at least three independent experiments are shown. (TIF)

Figure S3 Lymphoproliferative disease in STAT1−/− mice. a) Splenocytes from four ~36 week old STAT1−/− and two age matched WT mice are shown for comparison. b) Flow-cytometric analysis of splenic T cells from WT and STAT1−/− mice: Splenocytes from the indicated mice were gated on CD3^+ cells, and analyzed for CD62L and CD44-expressing subpopulations. (TIF)

Acknowledgments

We would like to thank Drs. A. Goldrath, S. Hedrick and R. Rickert for many helpful discussions and advice.

Author Contributions

Conceived and designed the experiments: HM DO YT MD. Performed the experiments: HM DO YT. Analyzed the data: HM DO YT MD. Wrote the paper: DO MD.
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