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## STAT5 IS CRITICAL TO MAINTAIN EFFECTOR CD8+ T CELL RESPONSES

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### Abstract

During an immune response, most effector T cells die, while some are maintained and become memory T cells. Factors controlling the survival of effector CD4+ and CD8+ T cells remain unclear. Here, we assessed the role of IL-7, IL-15, and their common signal transducer, STAT5, in maintaining effector CD4+ and CD8+ T cell responses. Following viral infection, IL-15 was required to maintain a subpopulation of effector CD8+ T cells expressing high levels of killer cell lectin-like receptor subfamily G, member 1 (KLRG1) and lower levels of CD127, while IL-7 and IL-15 acted together to maintain KLRG1<sup>lo</sup>CD127<sup>hi</sup> CD8+ effector T cells. In contrast, effector CD4+ T cell numbers were not affected by the individual or combined loss of IL-15 and IL-7. Both IL-7 and IL-15 drove phosphorylation of STAT5 (pSTAT5) within effector CD4+ and CD8+ T cells. When STAT5 was deleted during the course of infection, both KLRG1<sup>hi</sup>CD127<sup>lo</sup> and KLRG1<sup>lo</sup>CD127<sup>hi</sup> CD8+ T cells were lost, while effector CD4+ T cell populations were maintained. Further, STAT5 was required to maintain expression of Bcl-2 in effector CD8+, but not CD4+, T cells. Finally, IL-7 and IL-15 required STAT5 to induce Bcl-2 expression and to maintain effector CD8+ T cells. Together, these data demonstrate that IL-7 and IL-15 signaling converge on STAT5 to maintain effector CD8+ T cell responses.

### Introduction

Maintaining homeostasis of the T cell compartment is critical for normal immune system function. T cell homeostasis is altered during acute viral infection, when antigen-specific CD4+ and CD8+ T cells undergo massive expansion. Shortly thereafter, regulated induction of apoptosis, requiring the pro-apoptotic molecule, Bim, reduces the expanded T cell population and restores homeostasis (1,2). Mechanism(s) that control which effector cells die and which survive and become memory cells remains unclear.

Recent work has shown that heterogeneity within the effector CD8+ T cell pool may allow for identification of some effector T cells with “memory precursor” properties. For example, after acute viral infection, effector CD8+ T cells having increased expression of killer cell lectin-like receptor subfamily G, member 1 (KLRG1) and decreased expression of CD127 (IL-7R $\alpha$ ) had a decreased propensity to generate memory T cells after adoptive transfer (3).

Conversely, KLRG1<sup>lo</sup>CD127<sup>hi</sup> effector CD8<sup>+</sup> T cells had more potential for memory T cell generation when transferred into recipient animals (3). The functional significance of these markers remains unclear as enforced expression of CD127 did not prevent contraction of CD4<sup>+</sup> or CD8<sup>+</sup> T cell responses (4,5). Further, neutralization of IL-7 does not exacerbate contraction of effector CD4<sup>+</sup> T cell responses (6).

In contrast to IL-7, recent studies have shown that IL-15 is critical for effector CD8<sup>+</sup> T cell survival (7,8). Indeed, OT-1 TCR transgenic T cells survived poorly in IL-15-deficient recipient mice after infection with recombinant vesicular stomatitis virus expressing OVA (7). While almost all effector T cells were lost in IL-15-deficient recipients, the loss was greater in the KLRG1<sup>hi</sup> subset (7). Although the authors also provided data suggesting that IL-7 contributed to survival of KLRG1<sup>lo</sup> effector CD8<sup>+</sup> T cells, whether or not these cells were CD127<sup>hi</sup> was unclear due to the use of an anti-IL-7R $\alpha$  neutralizing antibody, which prevented detection of cell surface CD127 (7). In addition, neither anti-IL-7R $\alpha$  antibody, nor transfer of OT-I cells into IL-7-deficient mice instead of normal mice decreased OT-I cell numbers (7). Thus, it remains to be definitively established whether IL-7 plays a necessary, or even a redundant role with IL-15 in survival of subsets of effector CD8<sup>+</sup> T cells. Further, the existence of similar sub-populations of effector CD4<sup>+</sup> T cells, and their potential dependence on these same cytokines, is unknown.

Both IL-7 and IL-15 can activate similar signaling pathways within T cells, including PI-3K/AKT and STAT5. In T cells, both the PI-3K/AKT and Jak3/STAT5 pathways have been shown to be critical for T cell homeostasis (9,10). Further, both pathways have been implicated, but not directly examined, in both the proliferative and cell survival effects of IL-7 and IL-15 *in vivo* (11,12). However, both TCR and cytokine stimulation activate PI-3K/AKT signaling (13,14), which complicates interpretations of the specific role of the PI-3K/AKT pathway on particular stages of T cell homeostasis.

STAT5 exists as two isoforms, a and b, which are encoded by separate but linked genes (10). In T cells, their functions are largely redundant as deletion of either STAT5 gene did not drastically alter T cell homeostasis (15,16). The underwhelming phenotype of the original STAT5a/b-double-deficient mice relative to either *Jak3*<sup>-/-</sup>, *IL-7*<sup>-/-</sup>, and *IL-7R*<sup>-/-</sup> mice suggested that STAT5 signaling was not essential for naïve T cell homeostasis (17). However, recent data suggest that the original STAT5a/b<sup>-/-</sup> mice maintained expression of an N-terminal deleted, but partially functional STAT5 protein (10). The generation of new conditional STAT5a/b-deficient mice has revealed a profound effect of the loss of STAT5a/b on thymocyte development, similar to *Jak3*<sup>-/-</sup>, *IL-7*<sup>-/-</sup>, and *IL-7R*<sup>-/-</sup> mice (10). In addition, cell lineage-specific ablation of STAT5a/b by CD4Cre resulted in the dramatic loss of peripheral naïve CD8<sup>+</sup> and CD4<sup>+</sup> T cells (11). However, the role of STAT5 in effector T cell survival remains unclear. Here, we tested the role of IL-7, IL-15, and STAT5 in the maintenance of effector CD4<sup>+</sup> and CD8<sup>+</sup> T cells during viral infection.

## Materials and Methods

### Mice and viral infection

C57BL/6 were purchased from Jackson Labs or Taconic Farms. IL-15-deficient mice on a C57BL/6 background were purchased from Taconic Farms. STAT5a/b<sup>fl/fl</sup> mice were a generous gift from Dr. Lothar Hennighausen (National Institutes of Health, Washington, DC) and were crossed to C57BL/6 mice and then crossed to B6.Cg-Tg(Mx1Cre)1Cgn/J transgenic mice. All mice were used between 3–8 months of age. Mice were infected intraperitoneally (i.p.) with 2×10<sup>5</sup> pfu of the Armstrong strain of lymphocytic choriomeningitis virus (LCMV). LCMV was grown in BHK-21 cells and viral titers from spleen and liver homogenates were determined by plaque assay on BHK-21 monolayers as

described (18). Animals were housed under specific pathogen free conditions in the Division of Veterinary Services and experimental procedures were reviewed and approved by the institutional animal care and use committee (IACUC) at the Cincinnati Children's Hospital Research Foundation.

### IL-7 and IL-15 manipulation in vivo

For *in vivo* IL-7 blockade experiments, M25 was grown as ascites, purified by ammonium sulfate precipitation and ion exchange chromatography and injected i.p. at a dose of 3 mg/mouse every other day. Effectiveness of IL-7 blockade was assessed by measuring the numbers of pre-B cells in the bone marrow of each mouse via flow cytometry, using antibodies against IgM, B220, and CD24. For IL-7 and IL-15 delivery experiments, recombinant hIL-7/anti-IL-7 immune complexes were mixed *in vitro* and the equivalent of 2.5 µg of rhIL-7 (Peprotech) was injected i.p. on days 10, 12, 14 after infection. For IL-15 delivery experiments, IL-15/anti-IL-15Rα (R&D Systems, Minneapolis, MN) were mixed *in vitro* and the equivalent of 750 ng IL-15 was injected i.p. on days 10, 12, 14 after infection.

### MHC Tetramer staining and flow cytometry

1–2 million single spleen cell suspensions were stained with different combinations of the following cell surface antibodies: anti-CD4, CD8, CD44, KLRG1, CD127 (from either BD Biosciences or EBioscience) and intracellularly with either anti-Bcl-2 (clone 3F11, produced in house) or anti-STAT5 (Santa Cruz Biotechnologies) as described (19). To assess STAT5 levels, we used a FITC-labeled goat anti-rabbit antibody (Caltag). For detection of p-STAT5 in CD8<sup>+</sup> effector cells, splenocytes were first incubated for 2 hours at 37° C, then stimulated with either mIL-7 or mIL-15 (10 ng/ml) (R&D Systems) for 20 minutes, then stained with D<sup>b</sup>gp33 tetramer and anti-KLRG1 for 30 minutes at 4° C. The cells were then washed, fixed, permeabilized and stained with anti-pSTAT5-APC antibody (BD Biosciences) along with antibody against CD8 and secondary antibody against anti-KLRG1. For detection of p-STAT5 in CD4<sup>+</sup> effector cells, splenocytes were incubated for 2 hours at 37° C, then surface stained with antibodies against CD4, CD44, CD127 for 20 minutes at 37° C, then stimulated with either mIL-7 or mIL-15 (10ng/ml) (R&D Systems) for 20 minutes at 37° C. The cells were then washed, fixed, permeabilized and stained with anti-pSTAT5-APC antibody (BD Biosciences). Data were collected on an LSRII flow cytometer and analyzed using FacsDIVA software. D<sup>b</sup>gp33 tetramers were a generous gift from Dr. Alan Zajac and were coupled to either APC or to PE as previously described (20). I-A<sup>b</sup>gp61 tetramers were produced in house using a baculovirus expression system as described previously (20).

### In vitro cytokine culture and Bcl-2 mRNA analysis

Splenic CD8<sup>+</sup> T cells were purified by negative selection using CD8<sup>+</sup> T cell isolation kits (Miltenyi Biotech) and cultured with IL-7 and IL-15 (R&D Systems) +/- cycloheximide (Sigma) for 3 hours at 37°C. RNA was purified using Triazol and cDNA synthesized and real-time RT-PCR performed on an ABI Prism 7700 Sequence Detection System (Applied Biosystems) with SybrGreen® (Bio-Rad) using the following primer sets L19 forward 5'-CCTGAAGGTCAAAGGGAATGTG-3'; reverse 5'-GCTTTCGTGCTTCCTTGGTCT-3'. Bcl-2 forward 5'-TGGGATGCCTTTGTGGAACAT-3'; reverse 5'-AGAGACAGCCAGGAGAAATCAAAC-3'. L19 cycle counts were used to normalize cDNA levels between samples. The difference in Bcl-2 mRNA levels was calculated as follows: fold change = 1.8<sup>x</sup>, x = the difference in cycle count between cytokine treated and untreated samples after normalization to L19. Samples treated with cytokine + cycloheximide were further normalized to cycloheximide alone treated samples to negate non-specific effects of cycloheximide on Bcl-2 expression.

## IFN- $\alpha/\beta$ Bioassay

Levels of serum type I interferon was assessed by bioassay as described (21). Briefly, dilutions of serum were cultured with an L929 cell line that was stably transfected with an IFN-responsive luciferase construct. After 6 hrs, luciferase activity was measured in a luminometer and actual amounts of type I interferon were calculated based on a standard curve with recombinant IFN- $\alpha/\beta$ .

## Statistical Analyses

Statistical analyses were performed using Student's t-test with Microsoft Excel or with Minitab for Windows Software (Release 14), State College, Pennsylvania.

## Results

### Reciprocal expression of CD122 and CD127 on effector CD8+ and CD4+ T cells

While expression of cytokine receptors by effector CD8+ T cells is dynamic (22,23), few studies have examined cytokine receptor expression on activated, non TCR Tg CD4+ T cells. Here, we assessed the cell surface expression of CD122 (IL-2/15 receptor  $\beta$  chain) and CD127 on effector CD4+ and CD8+ T cells following infection with lymphocytic choriomeningitis virus (LCMV). On day 8 after infection, most LCMV-specific (sp.) CD4+ and CD8+ T cells had decreased expression of CD127, but the reciprocal was true of CD122 (Fig. 1A). Thus, while decreased CD127 expression limited the availability of IL-7 to most effector T cells, these same T cells had substantial ability to compete for IL-15 or IL-2, given their increased expression of CD122.

### IL-15 is critical to maintain Bcl-2 in most effector CD8+ but not CD4+ T cells

As IL-15 is critical for survival of most effector CD8+ T cells (7,8), and effector CD4+ T cells also increased CD122 expression, we next asked whether IL-15 contributed to effector CD4+ T cell survival. First, we assessed the role of IL-15 in maintaining levels of the anti-apoptotic molecule, Bcl-2 within LCMV-sp. CD4+ and CD8+ T cells from wildtype C57BL/6 (BL/6) versus IL-15<sup>-/-</sup> mice. At day 10 after infection, Bcl-2 levels were decreased in both LCMV-sp. CD8+ and CD4+ T cells and the lack of IL-15 led to a significant decrease Bcl-2 levels in LCMV-sp. CD8+, but not CD4+ T cells, (Fig. 1B, C, D). By day 20 after infection, levels of Bcl-2 were increased in effector CD8+ T cells in C57BL/6, but a large population of effector CD8+ T cells failed to increase Bcl-2 levels in IL-15<sup>-/-</sup> mice (Fig. 1B, D) as shown previously (8). In separate experiments, we found that the cells expressing low levels of Bcl-2 in IL-15<sup>-/-</sup> mice also expressed high levels of the killer cell lectin-like receptor subfamily G, member 1 KLRG-1 (data not shown). Interestingly, IL-15 was not required to maintain Bcl-2 expression in effector CD4+ T cells on day 20 after infection (Fig. 1C). Together, these data show that IL-15 is critical for normal Bcl-2 expression in effector CD8+, but not CD4+, T cells.

### IL-7 and IL-15 contribute redundantly to effector CD8+, but not CD4+, T cell survival

After acute LCMV infection, most effector CD8+ T cells were comprised of two major subpopulations (3), a population of KLRG1<sup>hi</sup>CD127<sup>lo</sup> cells and another population that is KLRG1<sup>lo</sup>CD127<sup>hi</sup> (Fig. 2A). Given that KLRG1<sup>lo</sup>CD127<sup>hi</sup> CD8+ T cells expressed both CD127 and CD122, it was logical that IL-7 and IL-15 might be redundant for maintaining this effector subpopulation. To determine the relative redundancy of IL-7 and IL-15 on effector CD4+ and CD8+ T cell survival, we infected groups of either BL/6 or IL-15<sup>-/-</sup> mice and treated them with either isotype control antibody or anti-IL-7 neutralizing antibody between days 10 and 20 after infection.

In BL/6 mice, we found that IL-7 was critical for survival of some KLRG1<sup>hi</sup>CD127<sup>lo</sup> and some KLRG1<sup>lo</sup>CD127<sup>hi</sup> CD8<sup>+</sup> T cells (Fig. 2B). In contrast, IL-15 was critical for survival of most KLRG1<sup>hi</sup>CD127<sup>lo</sup> and some KLRG1<sup>lo</sup>CD127<sup>hi</sup> effector CD8<sup>+</sup> T cells (Fig. 2B). Neutralization of IL-7 in IL-15<sup>-/-</sup> mice further decreased the numbers of both subsets of CD8<sup>+</sup> T cells (Fig. 2B). Further, Bcl-2 levels correlated with the cells loss as the combined effects of IL-7 and IL-15 were required to maintain high levels of Bcl-2 in both subsets of CD8<sup>+</sup> T cells (Fig. S1A,B). At no time was KLRG1 expression observed on LCMV-sp. CD4<sup>+</sup> T cells, although CD127 expression was increased on most of these cells by day 20 after infection (Fig. 2C). Thus, although KLRG1 marked subpopulations of LCMV-sp. effector CD8<sup>+</sup> T cells, the lack of KLRG1 on effector CD4<sup>+</sup> T cells, left CD127 as a potential marker to identify effector CD4<sup>+</sup> subpopulations. Neither the individual or combined loss of IL-7 and IL-15 had a significant effect on the total numbers of CD4<sup>+</sup> gp61-sp. T cells, irrespective of their CD127 expression (Fig. 2C and 2D). To ensure the effectiveness of IL-7 neutralization, we assessed pre-B cells in the bone marrow as described (6). In both BL/6 and IL-15<sup>-/-</sup> mice, administration of M25 caused a ~20-fold loss of IgM<sup>+</sup>B220<sup>int</sup> BM pre-B cells (Fig. S1C). Thus, dynamic regulation of cytokine receptors controls cytokine availability and thereby contributes to effector CD8<sup>+</sup>, but not CD4<sup>+</sup>, T cell survival.

### STAT5 is critical to maintain effector CD8<sup>+</sup> T cells during LCMV infection

STAT5 is known to be a common downstream signaling molecule for both IL-7 and IL-15 (24). To determine if IL-7 or IL-15 could activate STAT5, we cultured effector T cells with the cytokines and assessed STAT5 phosphorylation (pSTAT5) using a phospho-STAT5-specific monoclonal antibody and intracellular flow cytometry. IL-7 drove STAT5 phosphorylation in KLRG1<sup>hi</sup> CD8<sup>+</sup> T cells, and higher pSTAT5 in KLRG1<sup>lo</sup> LCMV-sp. CD8<sup>+</sup> T cells (Fig. 2E), consistent with the increased expression of CD127 on the latter population. Although the pSTAT5 stain worked in conjunction with the D<sup>b</sup>gp33-tetramer stain, it was technically infeasible with the IA<sup>b</sup>gp61-tetramer stain. Instead, we co-stained with CD44, a marker of activation. Recent work has shown that multiple LCMV epitopes are recognized by CD4<sup>+</sup> T cells, and suggest that early after infection, most activated CD44<sup>hi</sup> T cells are LCMV-specific (25). Similar to the results in effector CD8<sup>+</sup> T cells, IL-7 drove modest pSTAT5 in CD127<sup>lo</sup> CD4<sup>+</sup> T cells, but stronger pSTAT5 in CD127<sup>hi</sup> CD4<sup>+</sup> T cells (Fig. 2F). In contrast, IL-15 drove pSTAT5 similarly in all effector CD4<sup>+</sup> and CD8<sup>+</sup> T cells (Fig. 2E and F).

As both IL-7 and IL-15 activated STAT5, we next tested the requirement of STAT5 for survival of effector T cells using conditional STAT5-deficient mice (26). Because cell-lineage specific ablation of STAT5 affects both T cell development and peripheral naive T cell survival (11), we used an inducible method of deletion. As LCMV infection drives very high systemic levels of type I interferon (Fig. S2A), we used a transgenic system in which Cre expression is controlled by the  $\alpha$ IFN-inducible Mx1 promoter, to inducibly delete STAT5 during the course of the response to LCMV. Using a STAT5-specific antibody and intracellular flow cytometry, we found that by day 6 after LCMV infection, most CD4<sup>+</sup> IA<sup>b</sup>gp61-sp. T cells in Mx1Cre-STAT5<sup>fl/fl</sup> mice had significantly decreased expression of STAT5 (Fig. 3A). Similar decreases in STAT5 levels were observed using SDS-PAGE and Western blotting of purified CD4<sup>+</sup> T cell lysates from LCMV-infected Mx1Cre-STAT5<sup>fl/fl</sup> versus STAT5<sup>fl/fl</sup> mice (Fig. 3B). From day 6 to day 8 the frequency of CD4<sup>+</sup> IA<sup>b</sup>gp61-sp T cells that were STAT5<sup>lo</sup> remained constant (Fig. 3C). Further, the total numbers of CD4<sup>+</sup> IA<sup>b</sup>gp61+ T cells were slightly increased in Mx1Cre-STAT5<sup>fl/fl</sup> mice on day 6 after infection, but were unchanged on days 7 and 8 after LCMV infection (Figure 3D). The slight increase in LCMV-sp. CD4<sup>+</sup> T cells in Mx1Cre-STAT5<sup>fl/fl</sup> mice on day 6 was not reproducible (data not shown).

In contrast, the frequency of LCMV-sp. CD8<sup>+</sup> T cells that were STAT5<sup>lo</sup> in Mx1Cre-STAT5<sup>fl/fl</sup> mice was increased on day 6, increased further on day 7, and then decreased by day 8 after infection (Fig. 3E and G). SDS-PAGE and Western blotting again confirmed the loss of STAT5 in purified CD8<sup>+</sup> T cells (Fig. 3F). While the total numbers of LCMV-sp. CD8<sup>+</sup> T cells were not different between Mx1Cre-STAT5<sup>fl/fl</sup> versus STAT5<sup>fl/fl</sup> mice on day 7 after LCMV infection, they were significantly lower in Mx1Cre-STAT5<sup>fl/fl</sup> mice by day 8 after infection. The total number of CD8<sup>+</sup>gp33-sp. T cells on day 8 after infection was also not different between BL/6 and Mx1Cre Tg mice, arguing against a non-specific effect of Cre on the T cell response (data not shown). As previous work showed that STAT5 was critical for perforin expression (27, 28), it was possible that the loss of STAT5 might result in persistent viral infection that could influence CD8<sup>+</sup> T cell responses. However, no significant differences in viral loads were observed from either the livers or spleens of STAT5<sup>fl/fl</sup> nor Mx1Cre-STAT5<sup>fl/fl</sup> mice on days 6 or 7 after infection (Fig. S2B). It was also possible that the diminution of T cell numbers in infected Mx1Cre-STAT5<sup>fl/fl</sup> mice reflected a STAT5 contribution to T cell proliferation. However, using *in vivo* BrdU labeling, the frequency of CD8<sup>+</sup>gp33-sp. T cells that were BrdU<sup>+</sup> was actually increased in Mx1Cre-STAT5<sup>fl/fl</sup> compared to STAT5<sup>fl/fl</sup> mice on day 8 after infection (Fig. 4A). In addition, there was no difference in BrdU uptake in cells that were STAT5<sup>lo</sup> compared to those that were STAT5<sup>hi</sup> at this timepoint (Fig. 4B). Together, these data show that STAT5 is critical to maintain CD8<sup>+</sup>, but not CD4<sup>+</sup>, effector T cells.

### STAT5 is critical to maintain both effector CD8<sup>+</sup> T cell subpopulations

Because IL-15 was critical for KLRG1<sup>hi</sup>CD127<sup>lo</sup> cells, but IL-7 and IL-15 acted in concert to maintain KLRG1<sup>lo</sup>CD127<sup>hi</sup> cells, we next tested the requirement for STAT5 on these two sub-populations. Importantly, deletion of Stat5 was similar in both CD8<sup>+</sup> T cell subpopulations (Fig. S3A). The frequency of KLRG1<sup>hi</sup>CD127<sup>lo</sup> and KLRG1<sup>lo</sup>CD127<sup>hi</sup> CD8<sup>+</sup> T cells was similar in both STAT5<sup>fl/fl</sup> and Mx1Cre-STAT5<sup>fl/fl</sup> mice (Fig. 5A). However, both subsets of effector CD8<sup>+</sup> T cells were significantly reduced in Mx1Cre-STAT5<sup>fl/fl</sup> compared to STAT5<sup>fl/fl</sup> mice (Fig. 5A). As expected, the frequencies of CD127<sup>hi</sup> and total numbers of LCMV-sp. CD4<sup>+</sup> T cells were not different between STAT5<sup>fl/fl</sup> and Mx1Cre-STAT5<sup>fl/fl</sup> mice (Fig. 5B).

### Stat5 is critical for IL-7 and IL-15-driven upregulation of Bcl-2

As IL-15 was critical for Bcl-2 expression (Fig. 1B) and STAT5 was critical for maintenance of effector CD8<sup>+</sup> T cells (Fig. 3G, 3H) we next determined the role of STAT5 in promoting Bcl-2 expression. Even though fewer CD8<sup>+</sup>gp33-sp. cells were STAT5<sup>lo</sup> by day 15 after infection compared to day 8 (31.3%  $\pm$  4.8 versus 46.1%  $\pm$  8.1 respectively, Fig. S3B, C), Bcl-2 levels in LCMV-sp. CD8<sup>+</sup> T cells from Mx1Cre-STAT5<sup>fl/fl</sup> mice were significantly reduced compared to cells from STAT5<sup>fl/fl</sup> mice (Fig. 5C). Further, levels of Bcl-2 were significantly lower in STAT5<sup>lo</sup> (Bcl-2 MFI = 2116  $\pm$  101) compared to STAT5<sup>hi</sup> (Bcl-2 MFI = 3356  $\pm$  158) CD8<sup>+</sup>gp33-sp. T cells from Mx1Cre-STAT5<sup>fl/fl</sup> mice ( $p < 0.0001$ ; Student's t-test). As expected, Bcl-2 levels in CD4<sup>+</sup> gp61-sp. T cells were not affected by STAT5 deletion (Fig. 5C). Moreover, the frequencies of CD4<sup>+</sup> gp61-sp. T cells that were STAT5<sup>lo</sup> were similar between days 8 and 15 (56.5%  $\pm$  6.55 vs. 50.6%  $\pm$  0.98, respectively, Fig. S3B).

To assess the requirement for STAT5 in upregulation of Bcl-2 by IL-7 and IL-15, we cultured spleen cells from d7 LCMV-infected STAT5<sup>fl/fl</sup> or Mx1Cre-STAT5<sup>fl/fl</sup> mice with IL-7 or IL-15 and assessed levels of Bcl-2 within gp33-sp. CD8<sup>+</sup> T cells the next day. Notably, both IL-7 and IL-15 increased expression of Bcl-2 in a dose dependent fashion in T cells from STAT5<sup>fl/fl</sup> mice, while upregulation of Bcl-2 was significantly impaired in T cells from Mx1Cre-STAT5<sup>fl/fl</sup> mice (Fig. 6A and 6B). The slight induction that was observed in

response to IL-7 and IL-15 in Mx1Cre-STAT5<sup>fl/fl</sup> T cells was likely due to incomplete deletion of STAT5 within these cells, as only cells expressing high levels of STAT5 increased expression of Bcl-2 in response to IL-7 or IL-15 (Fig. S4). Combined, these data demonstrate that both IL-7 and IL-15 require STAT5 to induce Bcl-2 within effector CD8+ T cells.

The mechanism by which STAT5 controls Bcl-2 expression is controversial. One report suggested that the effects of STAT5 are indirect (i.e. STAT5 inducing expression of another factor that induces Bcl-2 transcription) (29). Other reports have suggested direct effects of STAT5 on Bcl-2 transcription (30,31). To determine which of these mechanism(s) contributes to cytokine-driven Bcl-2 expression in T cells, we cultured purified CD8+ T cells from LCMV-infected BL/6 mice with IL-7 or IL-15 with or without cycloheximide for 3 hours and then measured Bcl-2 mRNA levels by real-time RT-PCR. Interestingly, while both IL-7 and IL-15 both drove significant induction of Bcl-2 within 3 hours, the presence of cycloheximide did not reduce Bcl-2 mRNA levels in response to these cytokines (Fig. 6C and 6D) providing evidence that the effect of STAT5 on Bcl-2 expression is direct (i.e. it did not require new protein synthesis).

### IL-7 and IL-15 depend on Stat5 to maintain effector T cell responses

As both IL-7 and IL-15 required STAT5 to induce Bcl-2 expression *in vitro*, We next determined if IL-7 or IL-15 required for STAT5 to maintain effector T cell responses *in vivo*. To do this, we infected mice with LCMV, and starting on day 10 after infection treated mice with long-acting forms of either IL-7 or IL-15 (6,32) every other day until day 15 after infection. On day 15 we found that, both IL-7 and IL-15 significantly increased numbers of CD8+ gp33-sp. T cells in Stat5<sup>fl/fl</sup> mice (Fig. 6E and 6F). In contrast, IL-7 was unable to significantly increase gp33-sp. CD8+ T cells in Mx1Cre-STAT5<sup>fl/fl</sup> mice (Fig. 6E). While IL-15 did significantly increase gp33-sp. T cells in Mx1Cre-STAT5<sup>fl/fl</sup> mice; the increase was less than in STAT5<sup>fl/fl</sup> mice (Fig. 6F) and the levels of STAT5 were enhanced in Mx1Cre-STAT5<sup>fl/fl</sup> compared to STAT5<sup>fl/fl</sup> controls (data not shown). Thus, IL-15 likely enhanced selection of STAT5<sup>hi</sup> T cells *in vivo*. Combined, these data show that IL-7 and IL-15 largely require STAT5 for maintaining effector CD8+ T cells *in vivo*.

### Discussion

Here we demonstrate that STAT5 is critical to maintain effector CD8+T cells. In this model, Stat5 is deleted in multiple tissues as the Mx1Cre transgene is expressed ubiquitously. It is possible that some of the effects we observe could be due to deletion of Stat5 in non-T cells. For instance, STAT5 signaling in dendritic cells (DCs) may influence survival of effector T cells as it has recently been shown that IL-7 can act on DCs to regulate the size of the CD4 T cell pool under lymphopenic conditions (33). However, whether this axis operates under normal physiologic conditions or during viral infection remains unclear. If there were a non-T cell effect of Stat5 that was dominant, we would have expected to see similar effects on CD4+ and CD8+ T cell responses as expansion of both cells require DCs, and this was clearly not the case. Although we cannot completely rule out non-T cell effects of Stat5, our data clearly show that despite similar deletion of Stat5, CD8+ effector T cells rely on Stat5 considerably more than do effector CD4+ T cells. We were somewhat surprised to find that IL-15 was not required for maintaining effector CD4+ T cell responses, as a recent report showed that IL-15 contributes to survival of memory CD4+ T cells (34). It is possible that IL-15 becomes a more prominent survival factor for memory CD4+ T cells than for effector CD4+ T cells. Indeed, we found that expression of CD122 was quite transient on effector CD4+ T cells (data not shown), and previous work showed that increased expression of CD122 on memory CD4+ T cells (34), may facilitate their dependence on IL-15. We note that our data do not imply that IL-15 is not involved in effector CD4+ T cell survival, just

that it is not strictly required. Thus, mechanisms maintaining effector CD4<sup>+</sup> T cells remain unclear. Previous work has shown that Bcl-3, a NF- $\kappa$ B p50 family member, can promote survival of activated CD4<sup>+</sup> T cells (35). Further, activated CD4<sup>+</sup> T cells have dramatically increased expression of A1 (1,2), and A1 expression can be regulated by NF- $\kappa$ B signaling (36–38). A1 can also antagonize Bim, making A1 a logical candidate for promoting effector CD4<sup>+</sup> T cell survival. Future experiments will evaluate the requirement for A1 in the survival of effector CD4<sup>+</sup> T cells.

Our data in effector CD8<sup>+</sup> T cells are consistent with previous reports showing that in cell lines, IL-7 and IL-15 control expression of Bcl-2 via STAT5 (29,31,39). However, other reports have suggested that STAT5 is not critical for Bcl-2 induction (40,41). In one of these reports, mice were developed in which tyrosine 449 in the IL-7R $\alpha$  gene was mutated to a phenylalanine (41). The Y449F mutation incapacitates both STAT5 and PI-3K activation in response to IL-7 in lymphocytes and thymocytes (39,41). Using T cells from these Y449F mice, the authors showed that although IL-7 was unable to induce detectable pSTAT5, it was able to increase expression of Bcl-2 (41). From these data it was concluded that Bcl-2 upregulation was independent of STAT5 activation. However, at baseline, Bcl-2 levels were significantly lower in peripheral CD4<sup>+</sup> and CD8<sup>+</sup> T cells directly isolated from Y449F mice (41), suggesting that, physiologically, IL-7R $\alpha$ Y449 signaling controls expression of Bcl-2 *in vivo*. Consistent with this, another group showed, in a thymocyte cell line, that Bcl-2 upregulation and STAT5 activation in response to IL-7 signaling required Y449F of IL-7R $\alpha$  (39). Furthermore, mice deficient in JAK3, which is required for IL-7 and IL-15 activation of STAT5, exhibited a profound loss of peripheral CD8<sup>+</sup> T cells but had nearly normal levels of CD4<sup>+</sup> T cells (42–44). Moreover, JAK3-deficiency dramatically impaired Bcl-2 expression in CD8<sup>+</sup>CD4<sup>-</sup>, but not CD4<sup>+</sup>CD8<sup>-</sup> thymocytes (45). Thus, the simplest explanation is that, under physiologic conditions, IL-7 and IL-15 require STAT5 to maintain Bcl-2 expression within CD8<sup>+</sup> T cells.

Our data also suggest that STAT5 maintains Bcl-2 directly as cycloheximide failed to block Bcl-2 induction by IL-7 and IL-15. A previous paper showed in a pro-B cell line that cycloheximide blocked induction of Bcl-2 by IL-2, leading the authors to conclude that the effects of STAT5 on Bcl-2 induction by IL-2 were largely indirect (29). However, in that study the effect of cycloheximide was only partial and these results were not reported to be significant. Our results in primary T cells suggest that the effects of IL-7 on Bcl-2 induction are direct and do not require new protein synthesis. These data are supported by a recent report showing that, in mast cells, STAT5 can bind to a site in intron 2 of the Bcl-2 gene (30). Further work will be required to determine if cytokines promote STAT5 binding to this site in intron-2 in activated CD8<sup>+</sup> T cells.

These data have implications for the development of T cell memory that emerges from the effector pool. We and others have previously shown that the pro-apoptotic molecule, Bim, is critical for limiting the numbers of effector T cells that can become memory T cells (20,46,47). Here our data suggest that IL-7 and/or IL-15 signaling through Stat5 is likely critical to maintain levels of Bcl-2 sufficient to combat Bim and develop into memory T cells. Thus, cytokines may have use as potential vaccine adjuvants. Optimal vaccines would promote robust CD4<sup>+</sup> and CD8<sup>+</sup> T cell responses that would be long-lived. As we and others have shown, common gamma chain cytokines can significantly enhance CD4 and CD8 T cell responses (6,7,48). Although these cytokines enhanced short-term T cell responses, their effects were transient once the cytokines were withdrawn. This makes sense given that survival of nearly all populations of T cells is achieved, at least in part, via competition for limiting amounts of cytokines. However, we note that, in general these studies were done in viral infections in which the T cell responses are extraordinarily robust. It remains unclear if cytokine adjuvants, given short-term, may enhance long-term memory

responses under conditions in which less robust T cell responses are generated. This approach could be beneficial for enhancing sub-optimal vaccine responses which currently require several rounds of boosting.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

## Acknowledgments

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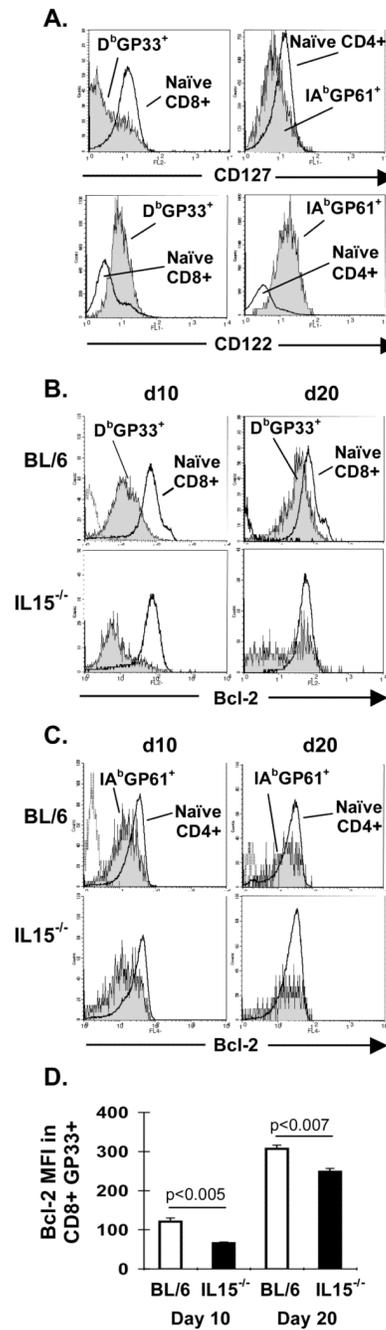
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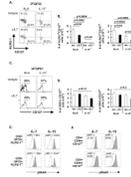
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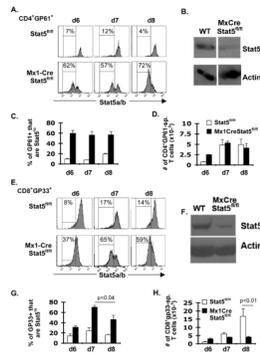


**Figure 1. Dynamic cytokine receptor expression in LCMV-specific CD4<sup>+</sup> and CD8<sup>+</sup> T cells**  
**(A)** Groups of BL/6 mice were uninfected (N = 2) or infected i.p. (N = 5) with LCMV ( $2 \times 10^5$  pfu), sacrificed at day 8 after infection and spleen cells were stained with MHC class I (left panels) and MHC class II tetramers (right panels). Results show the levels of CD127 (top panels) or CD122 (bottom panels) in naïve total CD4<sup>+</sup> or CD8<sup>+</sup> (open histograms) versus D<sup>b</sup>gp33-sp. or IA<sup>b</sup>gp61-sp. (gray histograms). **(B, C)** Groups of BL/6 (N = 5) or IL-15<sup>-/-</sup> (N = 5) mice were infected with LCMV, sacrificed on day 10 or 20 and Bcl-2 levels in T cells were assessed by intracellular flow cytometry. Results show the levels of Bcl-2 within **(B)** D<sup>b</sup>gp33-sp. versus **(C)** IA<sup>b</sup>gp61-sp. (gray histograms) compared to total naïve **(B)** CD8<sup>+</sup> or **(C)** CD4<sup>+</sup> (open histograms). Dashed line histograms in upper panels

represent isotype control staining. (D) Results show the mean fluorescence intensity of the Bcl-2 signal in CD8+ gp33-sp. T cells in either BL/6 or IL-15<sup>-/-</sup> mice on either day 10 or 20 after infection. Data are representative of 3 independent experiments.

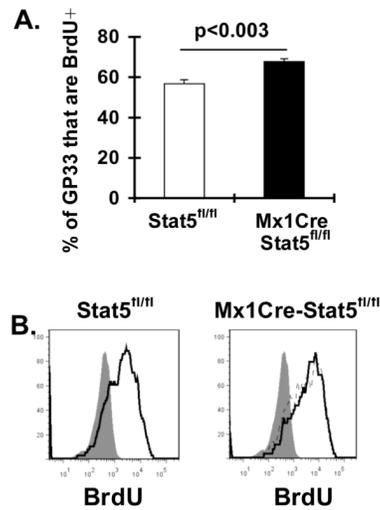


**Figure 2. IL-7 and IL-15 contribute redundantly for effector CD8+, but not CD4+ T cell survival** Groups of BL/6 or IL-15<sup>-/-</sup> mice (N = 5/group) were infected with LCMV and treated i.p. with either isotype control (MPC11) or  $\alpha$ -IL-7 (M25) antibody (3 mg/mouse on days 11, 13, 15, 17, 19) and sacrificed on day 20. Spleen cells were stained with D<sup>b</sup>gp33-tetramers and with antibodies against CD8, KLRG1, and CD127 or with I-A<sup>b</sup>gp61-tetramers and with antibodies against CD4, CD16/32, and CD127. Dot plots show (A) KLRG1 by CD127 staining in D<sup>b</sup>gp33-sp. CD8<sup>+</sup> T cells from either BL/6 (left panels) or IL-15<sup>-/-</sup> mice (right panels) treated with either isotype control (top panels) or  $\alpha$ -IL-7 antibody (bottom panels). (B) Graphs show the total numbers of KLRG1<sup>hi</sup>CD127<sup>lo</sup> versus KLRG1<sup>lo</sup>CD127<sup>hi</sup> D<sup>b</sup>gp33-sp. CD8<sup>+</sup> T cells. p values shown are from a Student's t-test analysis. (C) Histograms show CD127 staining of I-A<sup>b</sup>gp61-sp. CD4<sup>+</sup> T cells from BL/6 or IL-15<sup>-/-</sup> mice treated with either isotype control or M25 antibody. Gray histogram shows staining of I-A<sup>b</sup>gp61-sp. T cells with an isotype control antibody for CD127. (D) Graphs show total numbers of CD127<sup>hi</sup> versus CD127<sup>lo</sup> I-A<sup>b</sup>gp61-sp CD4<sup>+</sup> T cells from BL/6 or IL-15<sup>-/-</sup> mice treated with either MPC11 or M25 antibody mice. Data are representative of 3 independent experiments. (E, F) Groups of BL/6 mice (n=3) were infected with LCMV and sacrificed on 8 days later. pStat5 levels were determined in splenocytes following stimulation with cytokines (10ng/ml for 20 minutes) in either (E) D<sup>b</sup>gp33-specific CD8<sup>+</sup> T cells or (F) CD4<sup>+</sup> CD44<sup>hi</sup> cells. Histograms show pSTAT5 levels in cells treated with either media alone (open histograms) or with IL-7 or IL-15 (gray histograms). Data are representative of 2 independent experiments.



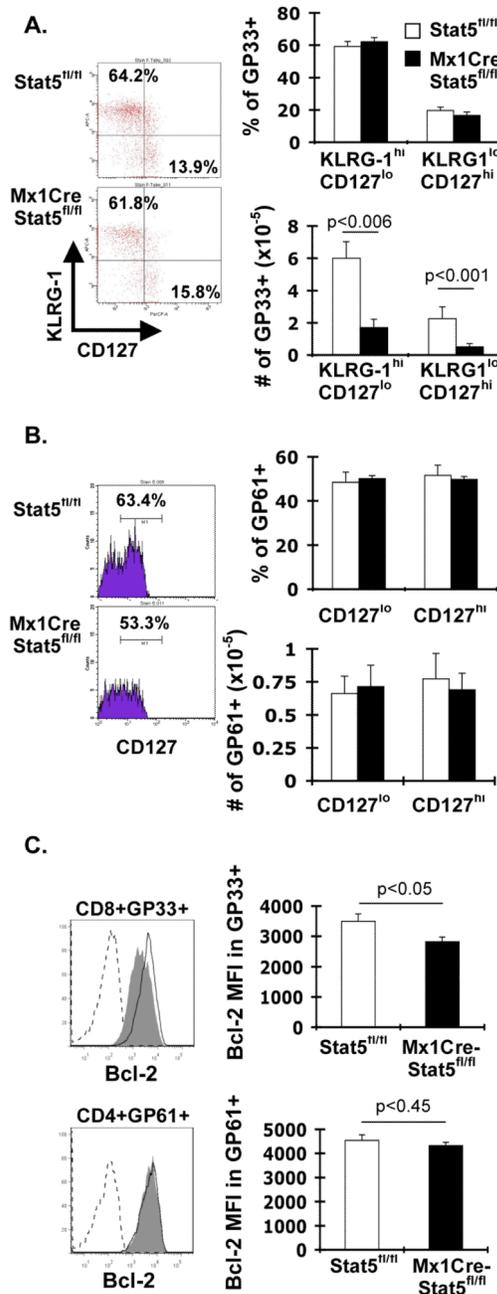
### Figure 3. STAT5 is critical for maintaining effector CD8+, but not CD4+, T cells

Groups of STAT5<sup>fl/fl</sup> or Mx1Cre-STAT5<sup>fl/fl</sup> mice (N = 3–5/group) were infected with LCMV and sacrificed 6–8 days later. Spleen cells were surface stained with either IA<sup>b</sup>gp61 or with D<sup>b</sup>gp33 tetramer along with antibodies against CD8 or CD4 and intracellularly with antibody against total STAT5. (A, C, E, G) Frequencies of MHC tetramer+ cells that were STAT5<sup>lo</sup> (C,G) and total overall numbers of tetramer+ T cells (D, H) are shown. Data from day 8 were pooled between 2 independent experiments. Shown are p values from a Student's t-test analysis. (B, F) Groups of mice of either STAT5<sup>fl/fl</sup> or Mx1Cre-STAT5<sup>fl/fl</sup> mice (N = 3–5/group) were infected with LCMV and sacrificed 8 days later. Total CD4+ (B) or CD8+ (F) were purified using a panT cell isolation kit (Miltenyi Biotech) and 1×10<sup>6</sup> cell equivalents were subjected to SDS-PAGE and Western blotting for either STAT5 or for actin. Data in figures 3B and 3F were done in separate experiments. The bands displayed in figures 3B for actin and STAT5 were from separate lanes run on the same gel. Data are representative of 2 independent experiments.



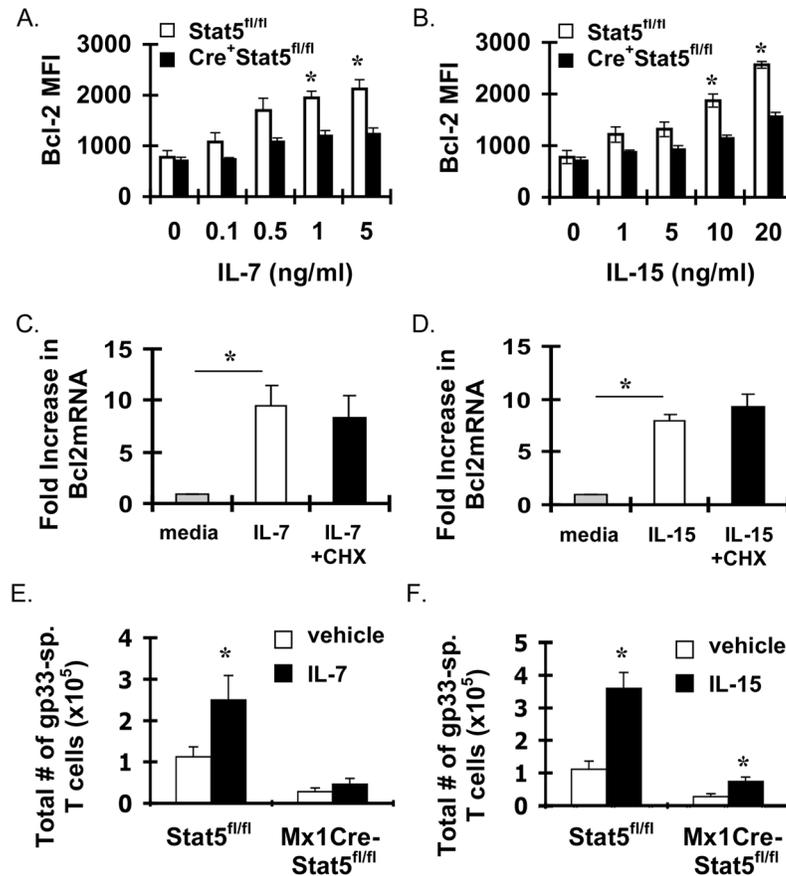
**Figure 4. In vivo proliferation of CD8+gp33-sp. T cells in Mx1Cre-Stat5<sup>fl/fl</sup> mice on day 8 after infection**

Groups of either Mx1CreStat5<sup>fl/fl</sup> versus Stat5<sup>fl/fl</sup> (N=5 mice/group) mice were infected i.p. with LCMV. On day 7 after infection, mice were injected i.p. with 0.8mg of BrdU in the morning and evening and sacrificed on day 8 after infection. One control mouse did not receive BrdU. Spleen cells were stained with antibodies against CD8, KLRG1, CD127 and intracellularly against STAT5 and BrdU. (A) Results show the frequency of CD8+ GP33-sp. T cells were BrdU+ from either Stat5<sup>fl/fl</sup> versus Mx1CreStat5<sup>fl/fl</sup> mice. (B) Representative histograms show BrdU staining intensity in CD8+gp33-sp. T cells from an LCMV-infected STAT5<sup>fl/fl</sup> mouse (left histogram, 60% BrdU+) versus a Mx1CreStat5<sup>fl/fl</sup> mouse after gating on STAT5<sup>hi</sup> (dark line, 67% BrdU+) versus STAT5<sup>lo</sup> (dashed line, 68% BrdU+) cells. The shaded histogram shows BrdU staining intensity in CD8+gp33+ T cells from the control mouse not injected with BrdU.



**Figure 5. STAT5 is critical for maintaining Bcl-2 levels in effector CD8+, but not CD4+, T cells**  
 Groups of either STAT5<sup>fl/fl</sup> or Mx1Cre-STAT5<sup>fl/fl</sup> mice (N = 5/group) were infected with LCMV and sacrificed 15 days later. (A) Representative dot plots show KLRG1 versus CD127 staining in gated CD8+ D<sup>b</sup>gp33-sp. T cells from either STAT5<sup>fl/fl</sup> (top plot) or Mx1Cre-STAT5<sup>fl/fl</sup> (bottom plot). Graphs show the frequency (top) and total numbers (bottom) of effector CD8+ subpopulations. Numbers of KLRG1<sup>hi</sup>CD127<sup>lo</sup> and KLRG1<sup>lo</sup>CD127<sup>hi</sup> cells were significantly decreased in Mx1Cre-STAT5<sup>fl/fl</sup> mice (Students t-test). (B) Representative histograms show CD127 staining in gated CD4+ IA<sup>b</sup>gp61-sp. T cells from either STAT5<sup>fl/fl</sup> (top histogram) or Mx1Cre-STAT5<sup>fl/fl</sup> (bottom histogram). Graphs show the frequency (top) and total numbers (bottom) of effector CD4+ subpopulations. (C) Representative histograms show Bcl-2 staining in gated CD8+ gp33-sp.

T cells (top histogram) or IA<sup>b</sup>gp61-sp. T cells (bottom histogram) from either STAT5<sup>fl/fl</sup> (open histogram) or Mx1Cre-STAT5<sup>fl/fl</sup> (shaded histogram). Isotype control staining is shown by the dashed line. Graphs show the mean fluorescence intensity (MFI) of the Bcl-2 stain in CD8+ D<sup>b</sup>gp33-sp. (top graph) or in CD4+ IA<sup>b</sup>gp61-sp. cells (bottom graph) from either STAT5<sup>fl/fl</sup> or Mx1Cre-STAT5<sup>fl/fl</sup> mice. Bcl-2 levels were significantly decreased in CD8+ D<sup>b</sup>gp33-sp. T cells from Mx1Cre-STAT5<sup>fl/fl</sup> mice (Students t-test). Data are representative of 2 independent experiments.



**Figure 6. IL-7 and IL-15 require Stat5 to drive Bcl-2 expression *in vitro* and to maintain effector T cell responses *in vivo***

Groups of *STAT5<sup>fl/fl</sup>* versus *Mx1Cre-STAT5<sup>fl/fl</sup>* mice (N = 3 mice/group) were infected with LCMV and sacrificed 7 days after infection. Spleen cells were cultured with the indicated concentrations of (A) IL-7 or (B) IL-15 overnight. Cells were then stained with D<sup>b</sup>-gp33 Tetramers, antibodies against CD8, KLRG1, and intracellularly with an  $\alpha$ -Bcl-2 antibody. Results show the MFI of the Bcl-2 stain within CD8+gp33-sp. T cells +/- SEM. (C, D) BL/6 mice were infected with LCMV, sacrificed on day 7 and purified splenic CD8+ T cells were cultured in triplicate with IL-7 or IL-15 with or without 2  $\mu$ M cycloheximide (CHX) for 3 hours. Cells were harvested, RNA was isolated, and cDNA generated and subjected to real-time RT-PCR for Bcl-2. Results show the fold change in Bcl-2 mRNA expression in response to (C) IL-7 or (D) IL-15 +/- S.D. (E, F) Groups of *STAT5<sup>fl/fl</sup>* versus *Mx1Cre-STAT5<sup>fl/fl</sup>* mice (N = 5 mice/group) were infected with LCMV and treated with either PBS, with (E) IL-7/ $\alpha$ -IL-7 immune complexes or with (F) IL-15/IL-15R $\alpha$  complexes on days 10, 12, 14 and sacrificed on day 15 after infection. Spleen cells were stained with D<sup>b</sup>-gp33 tetramers, antibodies against CD8, KLRG1, and IL-7R $\alpha$ . Results show the total number of D<sup>b</sup>-gp33-sp. T cells +/- SEM. \* = statistically significant difference between cytokine treated vs untreated control (Student's t-test). Data are representative of 2 independent experiments.