

Exploring regulatory mechanisms of CD8⁺ T cell contraction

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A small fraction of the CD8 T cell effector population that responds to an infection progresses through the contraction phase to the memory stage. The factors regulating the extent of contraction are poorly understood. Competition for limited resources has been widely postulated to be the cause of cell death during the contraction phase, but our data show that competition does not affect contraction kinetics. We go on to demonstrate that all effector cells present at the peak of the response have the potential to become bona fide memory cells, thus excluding selection on the basis of functionality. We propose that the fate of a CD8 effector cell is predetermined before the onset of contraction and discuss possible mechanisms of regulation.

infection | *Listeria monocytogenes* | memory | effector | bim

CD8 T cells leave the thymus 4–5 days after undergoing positive selection (1) and become fully mature CD8 T cells shortly after entering the periphery (2). Once a CD8 T cell is activated by antigen recognition in the periphery, it starts to proliferate and differentiate into an effector cell. The CD8 response typically peaks ≈7 days after an infection and is followed by the contraction phase, when 90–95% of the effector cells die in the ensuing days and weeks and the remaining 5–10% become long-lived memory cells (3–6). What cues do CD8 T cells receive that trigger the onset of contraction and regulate the extent of contraction?

The inflammatory environment during the priming phase has been identified as a key factor that influences contraction. A limited amount of inflammation leads to reduced effector proliferation and lack of a contraction phase (7). Mice that lack IL-12 generate a weaker primary response than their WT littermates, but show an increase in CD8 memory cell formation (8). In a setting of inflammation induced by infection, the duration of the initial T cell receptor (TCR) priming stimulus influences the size of the effector population (the longer lasting the signal the bigger the effector pool), but it does not appear to affect the onset or extent of contraction (9). Similarly, the kinetics of pathogen clearance seem to have limited to no impact on the onset or extent of contraction (10, 11). The extent of contraction, i.e., the ratio of the number of cells at the peak of the response to the number at the memory phase, is highly reproducible and comparable within different model systems, yet the mechanisms that control the extent of contraction so tightly are based on speculation.

There are two popular hypotheses that try to explain this phenomenon. The first states that cytokine deprivation caused by having a large population of effector cells competing for limited resources is responsible for cell death during the contraction phase (12–16). Indeed, injecting IL-2 (17) or IL-15 (15) during the contraction phase enhances CD8 T cell survival and limits the extent of contraction. The second hypothesis is based on the observation that the IL7R α^{hi} KLRG-1 $^{\text{lo}}$ (memory precursor cell) effector cell subset present at the peak of expansion preferentially survives contraction, whereas the IL7R α^{lo} KLRG-1 $^{\text{hi}}$ (terminally differentiated effector cell) subset is more prone to cell death (18). It has been proposed that terminally differentiated effector cells cannot make functional memory cells (“decreasing potential model”) (5) and are subsequently weeded out during the contraction phase. The first hypothesis postulates that the extent of contraction is actively regulated during the contraction phase,

whereas the second hypothesis predicts that the fate of the cell is determined before the contraction phase. The two hypotheses are not mutually exclusive as the subset of terminally differentiated cells might be destined to die and cells of the memory precursor subset might compete for limited resources.

Several studies examined the mechanisms causing cell death during contraction using either lymphocytic choriomeningitis virus (LCMV) or herpes simplex virus (HSV) as a model system. These data suggest that cell death during contraction is caspase independent (19) and that it cannot be rescued by overexpression of the prosurvival molecules bcl-2 or bcl-xl in CD8 T cells (20). However, CD8 effector cells that lack the proapoptotic molecule bim (21) are almost completely spared during contraction (22). We established an adoptive cotransfer system exploiting this unique feature of bim^{-/-} CD8 T cells to test the two hypotheses of contraction. Previous studies that examined the parameters that influence the extent of contraction were limited to altering variables like antigen presentation, inflammation, CD8 recruitment, etc., before the contraction phase (7, 10). This makes it impossible to determine what aspects of contraction are controlled in the expansion phase and what aspects are regulated during the contraction phase. Using the bim^{-/-} and WT cell cotransfer system, we overcome this limitation by keeping every parameter constant during the priming phase, leaving the extent of competition between effector cells as the only variable in the contraction phase. This allowed us to directly test the extent to which CD8 T cell contraction is regulated before or after the onset of contraction. Furthermore, we used the system to address whether there is a population of functionally unfit, terminally differentiated effector cells that is eliminated during contraction.

Results

Characterization of Bim^{-/-} CD8 T Cells. Bim^{-/-} mice display severe signs of autoimmunity and >50% are terminally ill at 1 year of age (23). B cell and T cell numbers are increased in knockout animals (23) and it is clear that the peripheral T cell phenotype is partially the result of impaired thymic negative selection (24). To avoid complications of using potentially autoreactive polyclonal bim^{-/-} CD8 T cells, we bred bim^{-/-} mice to OT-I TCR transgenic mice. Previous experiments in the literature have been done in a bim^{-/-} environment and it was not clear if the lack of CD8 T cell contraction was solely a CD8 T cell intrinsic characteristic (22, 25). To address this, we transferred 10⁴ naïve bim^{-/-} OT-I or WT OT-I T cells into congenic B6 hosts and infected the mice 1 day later with a priming dose of *L. monocytogenes* secreting ovalbumin (LM-OVA). WT and bim^{-/-} OT-I T cells expanded to the same extent by day 7 following infection (Fig. 1A Top). On day 14, the WT population had undergone contraction as shown by the 10-fold

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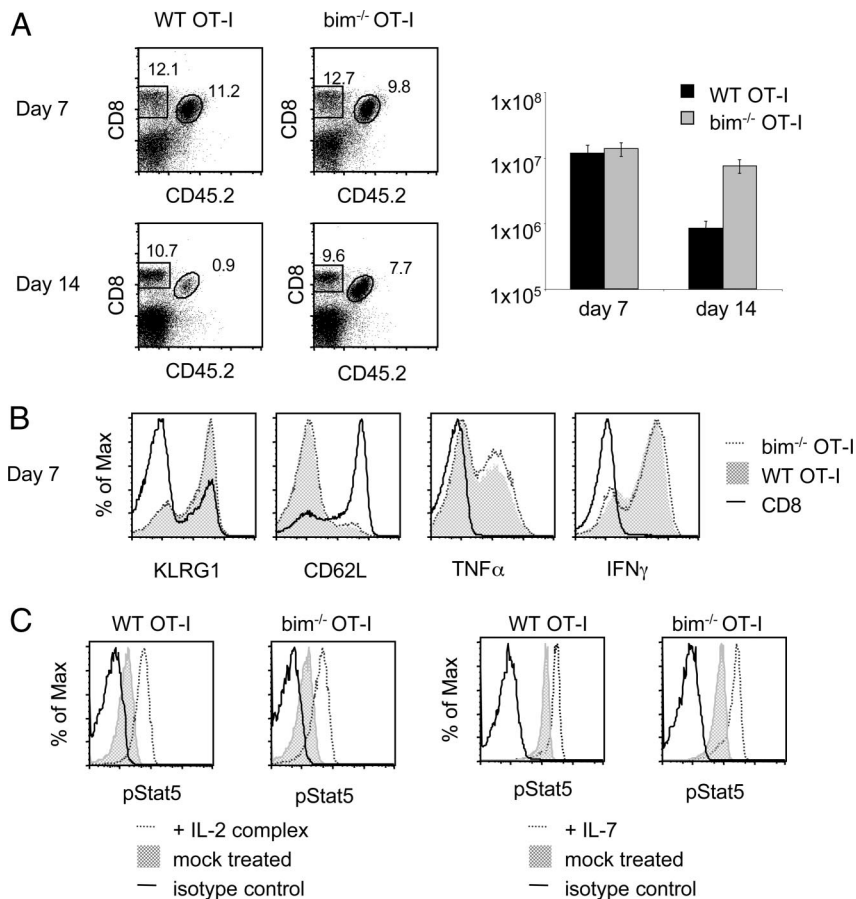


Fig. 1. Comparison of adoptively transferred WT and *bim*^{-/-} OT-I T cells after immunization. (A) 10⁴ *bim*^{-/-} or WT OT-I T cells were adoptively transferred into CD45.1 congenic B6 hosts and primed with LM-OVA 1 day later. Splensens were analyzed 7 days and 14 days after priming to determine extent of contraction. The number of splenic OT-I T cells on these days is shown in the histogram. (B) Phenotype (KLRG-1, CD62L) and function (TNF α and IFN γ) of host CD8 (black line), *bim*^{-/-} OT-I T cells (dotted line) and WT OT-I (transparent gray) effector cells were determined 7 days after infection with LM-OVA. (C) *Bim*^{-/-} and WT OT-I T cell responsiveness to cytokine stimulation were tested *in vivo* by IL-2 complex treatment of *bim*^{-/-} OT-I or WT OT-I mice (Left) and *in vitro* by determining IL-7 responsiveness (Right) of splenocytes containing memory *bim*^{-/-} OT-I T cells and WT OT-I T cells.

decrease in numbers compared to the day 7 time point. However, *bim*^{-/-} OT-I T cell numbers hardly changed between day 7 and day 14 (Fig. 1A). We concluded that the lack of contraction is indeed entirely CD8 T cell intrinsic and not caused by autoreactive TCRs or extrinsic stimuli from the *bim*^{-/-} environment.

We also compared the phenotype and function of WT and *bim*^{-/-} OT-I T cells on day 7 to ensure that *bim*^{-/-} OT-I T cells are equivalent to WT cells until the onset of the contraction phase. The expression pattern of KLRG-1, CD62L (Fig. 1B) and other phenotypic markers such as IL-7R α and CD27 (data not shown) is identical between *bim*^{-/-} and WT effector cells. Production of IFN γ and TNF α by the two types of effector cells was indistinguishable as well (Fig. 1B), showing that *bim*^{-/-} OT-I T cells are functionally equivalent to WT OT-I T cells. Finally, to ensure that *bim*^{-/-} OT-I T cells are indeed competitors for cytokines we tested their ability to respond to IL-7 signals *in vitro* and IL-2 complex stimulation *in vivo*. We found no difference in the extent of Stat5 phosphorylation between the WT and *bim*^{-/-} group and conclude that *bim*^{-/-} OT-I T cells are viable competitors indistinguishable from their WT counterparts in their ability to bind and respond to cytokines (Fig. 1C).

CD8 T Cells Do Not Contract More in the Presence of Competitors.

After establishing that *bim*^{-/-} OT-I effector cells were valid competitors, we went on to ask if the presence of these noncontracting *bim*^{-/-} OT-I T cells impacts the fate of WT OT-I T cells during contraction. The experimental setup consisted of transferring 10⁴ WT OT-I T cells into one set of B6 hosts, 10⁴ *bim*^{-/-} OT-I T cells into another set of hosts, and a mix of 5 \times 10³ WT OT-I T cells and 5 \times 10³ *bim*^{-/-} OT-I T cells into a third set of hosts. The premise of the setup was that all three groups would be identical until day 7 as all three groups received a total of 10⁴ OVA-specific T cells and the same priming dose of LM-OVA. Thus, T cell

expansion and pathogen clearance should be identical, providing a scenario where the only variable parameter was the number of effector T cells, i.e., the amount of competition for resources, after the peak of the expansion.

We challenged the three different groups (WT only, *bim*^{-/-} only, mix) with LM-OVA and bled the mice 7 days later to confirm successful priming of all animals (Fig. 2A). All three groups had equally sized OT-I populations and *bim*^{-/-} and WT OT-I T cells expanded equivalently in the mix group (Fig. 2A and D). The mice were bled again on day 14 to determine the extent of contraction (Fig. 2B). The OT-I T cells in the *bim*^{-/-}-only group showed little contraction, while the WT-only group had 10-fold fewer OT-I T cells by percentage (Fig. 2B). Mice that received a mixture of WT and *bim*^{-/-} OT-I T cells (mix OT-I group) had significantly fewer OT-I T cells than the *bim*^{-/-}-only group. More than 90% of the OT-I T cells in the mix group were *bim*^{-/-} OT-I T cells (Fig. 2D). The percentage of WT OT-I T cells (of the total OT-I T cell pool) on day 14 ranged from 5% to 15% in different experiments (Figs. 2D, 5, 6, and data not shown). We consistently found that the remaining WT cells of the mix group were \approx 2-fold lower (by percentage) compared to the WT-only group. Because the WT-only group received twice the number of WT cells transferred originally (and had twice the amount on day 7), this suggested that WT cells in the mix group were unaffected by the presence of a large cohort of competitors. To confirm that survival of effector cells was indeed unaffected by a large number of competitors during contraction, we determined absolute cell numbers in the spleen on days 7 and 14 (Fig. 2C). We found that each population expanded equally well and in proportion to its starting population, i.e., there were twice as many OT-I cells present from the *bim*^{-/-}-only and WT-only groups (that received 10⁴ cells each) compared to the *bim*^{-/-} and WT population in the mix group (that received 5 \times 10³

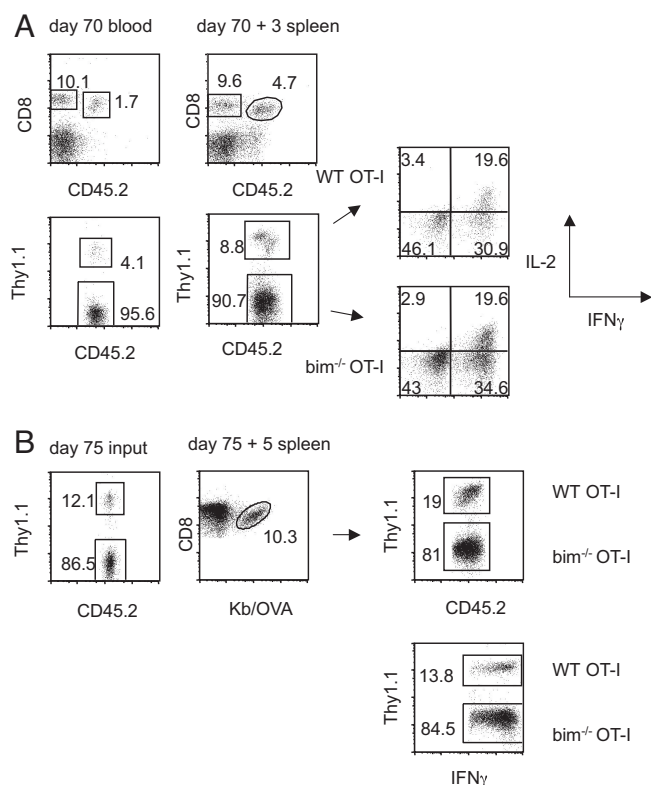


Fig. 6. All *bim*^{-/-} OT-I T cells have the ability to make functional memory cells. Mice received a mix of 5×10^3 WT (Thy1.1⁺) and 5×10^3 *bim*^{-/-} OT-I T cells and were infected with VSV-OVA 1 day later. (A) Mice were bled on day 70 to assess the WT-to-*bim*^{-/-} ratio, infected with LM-OVA, and killed 3 days later. The WT-to-*bim*^{-/-} ratio and ability to make IL-2 and IFN γ were determined. (B) Memory cells (day 75 post-VSV-OVA infection) were sorted and 1×10^4 cells injected (without changing the *bim*-to-WT ratio) into B6 hosts. Mice were infected with LM-OVA and killed 5 days later. OT-I T cells were identified by tetramer staining or as IFN γ ⁺ cells and *bim*^{-/-} and WT cells were distinguished using Thy1.1.

effector cell, and subsequent progression to a memory cell (9). The contraction phase that connects the initial expansion phase with the memory phase has been a black box from which surviving effectors emerge as bona fide long-lived memory cells (3–5). Numerous pieces of evidence suggest that competition for limited resources is a potential cause for the massive CD8 T cell death during the contraction phase. The availability of cytokines regulates T cell homeostasis—an overabundance of cytokines can trigger proliferation and the lack of essential survival cytokines like IL-7 can lead to cell death (30). Indeed, injections of the prosurvival cytokines IL-15 (15) and IL-2 (17) during contraction can rescue effector cells and increase the size of the memory pool. *Bim* and *puma* are members of the proapoptotic BH3-only family that regulates survival vs. cell death in the absence of cytokines (31). As such, cells that lack either *bim* (22, 32) or *puma* (12, 32) progress in greater numbers to the memory stage. Thus it has been widely proposed that cytokine deprivation causes death during contraction (12–16). However, using a cotransfer system of WT and *bim*^{-/-} OT-I T cells, we found that competition at a physiologically relevant level does not regulate contraction of CD8 effector cells. We kept the number of effector CD8 T cells at the peak of the response constant, but increased the number of effector T cells more than fivefold during contraction without affecting number, function, or phenotype of the OT-I WT cells (Figs. 2 and 3). While adding exogenous cytokines at the right time can spare cells from death during contraction and might have therapeutic potential, our

data clearly show that it is not competition for cytokines or any other limiting resource that determines the extent of contraction of CD8 T cells in a physiological setting. This includes clonotype-specific competition, which has been shown to play a key role in CD4 T cell viability (33). If contraction is not actively regulated by competition, is cell death predetermined for functionally unfit cells that get subsequently weeded out during contraction?

We show that all *bim*^{-/-} effector cells at the peak of the response can give rise to functional and fit memory cells that are maintained equivalently to their WT counterparts over the course of 10 weeks (Fig. 2D). This is in contrast to a previous study (25), probably because of the facts that our *bim*^{-/-} memory cells had a defined TCR specificity and were neither autoreactive themselves nor maintained in an autoreactive environment. In agreement with the aforementioned study, we found that the lack of *bim* results in a higher percentage of IL7R α ^{lo} KLRG-1^{hi} cells at day 14 compared to the WT population (Fig. 4B). This is because IL7R α ^{lo} KLRG-1^{hi} cells contract more than their IL7R α ^{hi} KLRG-1^{lo} counterparts (18), although it is noteworthy that this is a trend rather than a rule as WT IL7R α ^{hi} KLRG-1^{lo} cells contract substantially as well (Fig. 4C). *Bim*^{-/-} OT-I T cells do acquire a memory phenotype following the contraction phase, suggesting that the large cohort of doomed effector cells is capable of further differentiation (Fig. 5).

The WT-to-*bim*^{-/-} ratio is stable over time (Fig. 2D), suggesting that the change in the phenotype of *bim*^{-/-} OT-I cells is not because of a preferential loss of the IL7R α ^{lo} KLRG-1^{hi} *bim*^{-/-} population nor of an increase in the turnover of *bim*^{-/-} IL-7R α ^{hi} KLRG-1^{lo} cells, as this would skew the WT-to-*bim*^{-/-} ratio. However, while the WT-to-*bim*^{-/-} ratio remained constant, the overall percentage of OT-I T cells in the blood decreased even 30 days after infection (data not shown). This late decrease in memory CD8 T cells, which is equal in WT and *bim*^{-/-} cells, may be explained by slow elimination of IL7R α ^{lo} KLRG-1^{hi} cells. Further experiments will be required to obtain a definitive answer.

The equivalent functional ability of the IL7R α ^{lo} KLRG-1^{hi} and IL7R α ^{hi} KLRG-1^{lo} subsets is undisputed, but the proliferative potential upon restimulation is controversial (18, 26, 34–37). Our data agree with the notion that IL7R α ^{lo} KLRG-1^{hi} cells are not impaired upon restimulation (Fig. 6). In fact, our data show that both the IL7R α ^{hi} KLRG-1^{lo} cells that preferentially survive contraction and IL7R α ^{lo} KLRG-1^{hi} cells that tend to be eliminated during contraction are indeed functional (Fig. 6). The presence of a large fraction of doomed, nonfunctional *bim*^{-/-} OT-I T cells would have resulted in a major shift in the ratio of *bim*^{-/-} to WT cells following recall, which was not observed in recall responses (Fig. 6). This suggests that every single effector T cell present at the peak of the response has the intrinsic ability to become a functional memory cell if allowed to survive contraction.

Our initial experiments demonstrated that contraction is predetermined and not influenced by the number of T cells present during contraction, but there is no specific marker that identifies surviving CD8 T cells before the contraction phase and correlates strictly with their fate during contraction. A recent study demonstrated that the KLRG-1 phenotype of effector cells is determined ≈ 2 days before the onset of contraction, but a thorough analysis of effector cells during the expansion phase failed to reveal another differentially regulated marker (34). Similarly, microarray characterization of the two KLRG-1 subsets did not give more clues regarding the regulation of life vs. death during contraction (34).

How could contraction be regulated? Epigenetic modifications have been shown to play a key role in CD8 T cell function and differentiation (38). They are key regulators of many biological processes that involve lineage commitment or differentiation. Cell death could be the default pathway during contraction and a combination of epigenetic modifications and transcriptional changes could allow a subset of cells to survive and progress to the memory stage. This could be done by unequal segregation of proteins in the daughter cells after mitosis (39, 40), leading to the generation of

a T cell subset that is poised to survive, which could explain how the fate of a cell during contraction can be predetermined and tightly regulated. The notion of asymmetric cell division has been established in developmental processes and has recently been suggested to play a role in CD8 T cell differentiation as well (41).

Experimental Procedures

Mice. *Bim*^{-/-} mice (stock no. 004525) and C57BL/6.SJL (stock no. 002014) mice were obtained from The Jackson Laboratory (Bar Harbor, ME) and housed in specific pathogen-free conditions in the animal facilities at the University of Washington (Seattle, WA). OT-I TCR transgenic mice congenic for Thy1.1 were bred and maintained in the same facilities. OT-I TCR transgenic mice were bred to *bim*^{-/-} mice to generate *bim*^{-/-} OT-I TCR transgenic mice. C57BL/6.SJL (CD45.1+) host mice were infected at 8–12 weeks of age. Data shown for each experiment are from at least three independent experiments with at least three animals per group. All experiments were done in accordance with the Institutional Animal Care and Use Committee guidelines.

Adoptive Transfer and Cell Sorting. Naïve CD44^{low} OT-I T cells were isolated from lymph nodes using a CD8 isolation kit (Miltenyi) plus anti-CD44^{bio} (IM-7) as previously described (42). A total number of 1×10^4 OT-I T cells (WT only, *bim*^{-/-} only, or a 1:1 mix) per recipient was transferred intravenously (i.v.).

Infections. *L. monocytogenes* that expresses a secreted form of OVA (LM-OVA) (43) was grown as previously described (44). For primary infections, mice were injected i.v. with 2×10^3 cfu LM-OVA 1 day after adoptive transfer of OT-I T cells. For secondary infections, mice received 2×10^5 cfu LM-OVA and were killed 3–5

days later. For priming with VSV-OVA (45) and vaccinia virus (VAC)-OVA (46) mice were injected i.v. with 1×10^6 pfu and 2×10^6 pfu, respectively.

Flow Cytometry. Recipient mice were killed at the time points indicated and single-cell suspensions were prepared from the spleen, and in some experiments from lymph nodes and lungs after perfusion of the animal. Cells were typically stained with anti-CD8, anti-CD62L, anti-CD27, anti-KLRG-1, anti-IL-7R α , anti-Thy1.1, anti-CD45.2 (eBioscience and BD). For intracellular staining, cells were prepared with the Cytofix/Cytoperm kit in the presence of brefeldin A (BD) and stained with anti-IFN γ , IL-2, TNF- α , anti-Thy1.1, anti-CD45.2, and anti-CD8 (eBiosciences and BD). For BrdU incorporation, 1 mg BrdU was injected i.p. 12 and 24 h before harvesting (2 mg total). Cells were stained using anti-BrdU-APC Abs according to manufacturer's protocol (BrdU Flow Kit; BD). Cells were analyzed using a FACSCanto (BD) and analyzed using FLOWJO (TreeStar) software.

Stat5 Phosphorylation. Mice were treated with IL-2 complex for 1 h and the amount of Stat5 phosphorylation was determined as previously described (47). Briefly, splenocytes were fixed in 1.6% formaldehyde/PBS for 30 min at room temperature, followed by permeabilization with ice-cold methanol. The fixed and permeabilized cells were stained in 2% FCS/PBS with anti-phospho-STAT5 (Y694; Phosflow, BD) mAb. Mouse IgG1 mAb (BDZ) was used as the isotype control. For *in vitro* experiments, single-cell suspensions of spleens were prepared and resuspended in prewarmed RP10; IL-7 was added to a final concentration of 1 ng/ml and incubated for 30 min at 37 °C before analysis.

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