

Video Article

# Phenotypic and Functional Analysis of Activated Regulatory T Cells Isolated from Chronic Lymphocytic Choriomeningitis Virus-infected Mice

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## Abstract

Regulatory T (T<sub>reg</sub>) cells, which express Foxp3 as a transcription factor, are subsets of CD4<sup>+</sup> T cells. T<sub>reg</sub> cells play crucial roles in immune tolerance and homeostasis maintenance by regulating the immune response. The primary role of T<sub>reg</sub> cells is to suppress the proliferation of effector T (T<sub>eff</sub>) cells and the production of cytokines such as IFN- $\gamma$ , TNF- $\alpha$ , and IL-2. It has been demonstrated that T<sub>reg</sub> cells' ability to inhibit the function of T<sub>eff</sub> cells is enhanced during persistent pathogen infection and cancer development. To clarify the function of T<sub>reg</sub> cells under resting or inflamed conditions, a variety of *in vitro* suppression assays using mouse or human T<sub>reg</sub> cells have been devised. The main aim of this study is to develop a method to compare the differences in phenotype and suppressive function between resting and activated T<sub>reg</sub> cells. To isolate activated T<sub>reg</sub> cells, mice were infected with lymphocytic choriomeningitis virus (LCMV) clone 13 (CL13), a chronic strain of LCMV. T<sub>reg</sub> cells isolated from the spleen of LCMV CL13-infected mice exhibited both the activated phenotype and enhanced suppressive activity compared with resting T<sub>reg</sub> cells isolated from naïve mice. Here, we describe the basic protocol for *ex vivo* phenotype analysis to distinguish activated T<sub>reg</sub> cells from resting T<sub>reg</sub> cells. Furthermore, we describe a protocol for the measurement of the suppressive activity of fully activated T<sub>reg</sub> cells.

## Video Link

The video component of this article can be found at <https://www.jove.com/video/54138/>

## Introduction

Regulatory T (T<sub>reg</sub>) cells express forkhead box P3 (Foxp3) as a transcription factor for their development and function<sup>1</sup>. Additionally, T<sub>reg</sub> cells express various other molecules such as CD25<sup>2</sup>, lymphocyte-activation gene 3 (LAG-3)<sup>3</sup>, glucocorticoid-induced tumor necrosis factor receptor<sup>4</sup>, and cytotoxic T-lymphocyte-associated protein 4 (CTLA-4)<sup>5</sup> on their surface or intracellular region. During chronic infection with various kinds of pathogens such as viruses<sup>6,7</sup>, bacteria<sup>8,9</sup>, and parasites<sup>10-12</sup>, or in the course of cancer development<sup>13,14</sup>, T<sub>reg</sub> cells become differentiated into activated cells, displaying enhanced suppressive function targeting effector CD4<sup>+</sup> and CD8<sup>+</sup> T cells. A number of papers have suggested that expanded and activated T<sub>reg</sub> cells contribute to the impaired CD8<sup>+</sup> T cell response during friend retrovirus (FV) infection<sup>15-17</sup>. FV-induced T<sub>reg</sub> cells inhibit IFN- $\gamma$  or granzyme B expression and cytotoxic reactivity of CD8<sup>+</sup> T cells<sup>15-17</sup>. Moreover, in a herpes simplex virus infection model, it was reported that depletion of CD4<sup>+</sup>CD25<sup>+</sup> T<sub>reg</sub> cells resulted in expansion of virus-specific CD8<sup>+</sup> T cells and severe tissue damage by infiltration of immunopathogenic CD4<sup>+</sup> T cells<sup>18-20</sup>.

Mice infected chronically with the clone 13 strain of lymphocytic choriomeningitis virus (LCMV CL13)<sup>21-24</sup> have been widely used to characterize the phenotype and function of effector T cells (T<sub>eff</sub>) and T<sub>reg</sub> cells during chronic virus infection. During persistent LCMV infection, virus-specific T<sub>eff</sub> cells progressively lose their effector function and become exhausted T (T<sub>exh</sub>) cells. On the other hand, T<sub>reg</sub> cells reinforce their ability to suppress virus-specific T cell response<sup>25</sup>. The decrease in the functioning capacity of the T<sub>eff</sub> cells can be explained by several factors such as upregulation of inhibitory receptors on T<sub>eff</sub> cells, altered function of antigen-presenting cells, production of immunoregulatory cytokines, and increased frequency or enhanced function of T<sub>reg</sub> cells<sup>26</sup>. Among the factors involved in T cell suppression, programmed cell death protein-1 (PD-1)-expressing T<sub>exh</sub> cells and T<sub>reg</sub> cells have been widely considered as the hallmarks of antigen persistence and suppressive environment. Recently, it was reported that blockade of the PD-1 pathway and ablation of T<sub>reg</sub> cells lead to enhanced T cell function and decreased viral load during LCMV chronic infection<sup>27</sup>. Furthermore, T<sub>reg</sub> cells are activated during chronic infection of mice with LCMV<sup>23,25</sup> and their suppressive function is strengthened<sup>25</sup>. PD-1 is highly expressed on T<sub>reg</sub> cells as well as T<sub>exh</sub> cells, and the level of PD-1 expressed by T<sub>reg</sub> cells correlates with the strength of their suppressive function to inhibit T cell proliferation<sup>25</sup>.

Here, we describe a method to compare the characteristics of activated T<sub>reg</sub> cells isolated from mice infected with LCMV CL13 and resting T<sub>reg</sub> cells isolated from naïve mice. Furthermore, we explain a series of processes to separate activated T<sub>reg</sub> cells and examine their *ex vivo* phenotype, as well as measure their suppressive activity *in vitro*.

## Protocol

In this study, mice were maintained in a specific pathogen-free facility of the Yonsei Laboratory Animal Research Center of Yonsei University. All animal experiments were conducted in accordance with the Korean Food and Drug Administration guidelines using protocols approved by the International Animal Care and Use Committee of the Yonsei Laboratory Animal Research Center at Yonsei University.

### 1. Preparation of Solutions

1. Prepare 2% RPMI media by diluting fetal bovine serum (FBS) to 2% and penicillin-streptomycin to 1% in RPMI
2. Prepare complete RPMI media. To RPMI media add 10% of FBS, 1% of penicillin-streptomycin, 1% of L-glutamine, and 50  $\mu$ M 2-mercaptoethanol.
3. Prepare fluorescence activated cell sorting (FACS) buffer. To do so, supplement phosphate-buffered saline (PBS) with 2% of FBS.
4. Prepare T cell isolation buffer. To do so, supplement PBS with 2% of FBS and 2 mM ethylenediaminetetraacetic acid.

### 2. Isolation of Splenic Lymphocytes

1. Remove the spleens from naïve or LCMV CL13-infected mice as described previously<sup>19</sup> and place them in 60 mm x 15 mm Petri dishes containing 10 ml of 2% RPMI media.  
NOTE: In this study experiments, 5 to 6 week-old C57B1L/6J female mice received  $2 \times 10^6$  plaque-forming units (p.f.u.) of LCMV CL13 through intravenous injection via the tail vein. Sacrifice the mice at day 16 post-infection (16 d p.i.)<sup>25</sup>. Analyze ge-matched naive mice were on the same day.
2. Place a cell strainer into a 50 ml tube and rinse it with 2 ml of 2% RPMI media. Place the spleen on the 70  $\mu$ m cell strainer and grind using the plunger of a syringe. Rinse the cell strainer with 2 ml of 2% RPMI media and remove it from the 50 ml tube.
3. Fill up the tube with 2% RPMI media and centrifuge at 300 x g for 5 min at 4 °C. Discard the supernatant and resuspend the cell pellet in 1 ml of ACK lysing buffer. Incubate the samples at room temperature (RT) for 5 min.  
NOTE: The volume of ACK lysing buffer indicated above is for one spleen. Scale up the volume accordingly for pooled spleen samples.
4. Fill up the tube with 2% RPMI media and centrifuge at 300 x g for 10 min at 4 °C. Discard the supernatant and resuspend the cells at a density of  $1 \times 10^7$  cells/ml in complete RPMI media.  
NOTE: For live cell counting, stain cells with trypan blue and count only live cells that are not stained using hemocytometer.

### 3. Phenotyping of Splenic Conventional T ( $T_{conv}$ ) Cells and $T_{reg}$ Cells

NOTE: Before  $T_{reg}$  cell isolation, examine the phenotype of splenic lymphocytes isolated from naïve or infected mice by staining the cells with various antibodies and analyzing them by flow cytometry.

1. Transfer 50  $\mu$ l of the cells ( $5 \times 10^5$  cells) among total splenic lymphocytes into each well of a new u-bottom 96-well plate and add 150  $\mu$ l of FACS buffer per well. Centrifuge at 300 x g for 2 min at 4 °C.
2. Discard the supernatant. Agitate the cell pellet and add 200  $\mu$ l of FACS buffer per well. Centrifuge at 300 x g for 2 min at 4 °C (2 times).
3. Prepare the antibody cocktail for staining cell surface markers in 50  $\mu$ l of FACS buffer per well with the following antibodies and reagents. Anti-CD4 violet dye, Anti-CD25 green dye, Anti-PD-1 violet dye, Anti-CD8 PerCP-Cy5.5 dye and cell viability detection reagent near-infrared (IR) fluorescent reactive dye.  
NOTE: To further analyze the phenotype of activated  $T_{reg}$  cells, anti-CD103<sup>23,25</sup> can be added for cell surface-marker staining.
4. Resuspend the cell pellet with 50  $\mu$ l of antibody cocktail per well. Incubate for 20 min in the dark at 4 °C. Wash the cells twice with FACS buffer by centrifugation at 300 x g for 2 min at 4 °C.
5. After the final washing step, decant the supernatant and fix the cells for 20 min in the dark at 4 °C with 100  $\mu$ l of fixation buffer prepared according to the manufacturer's protocol.
6. Wash the cells twice with the permeabilization wash buffer (prepared according to manufacturer's protocol) by centrifugation at 300 x g for 2 min at 4 °C.
7. Prepare the antibody solution for intracellular Foxp3 staining, and resuspend the cell pellet with 50  $\mu$ l of Anti-Foxp3 antibody solution.  
NOTE: To further analyze the phenotypes of  $T_{conv}$  and  $T_{reg}$  cells, anti-CTLA can be added at this step.
8. Incubate for 20 min in the dark at 4 °C and repeat Step 3.6. After the final washing step, resuspend the cell pellet in 200  $\mu$ l of FACS buffer and examine the phenotype of the cells by flow cytometer<sup>25</sup>.
9. Gate the live cell population. To analyze the phenotypes of  $T_{conv}$  cells during LCMV CL13 infection, examine the frequency of Foxp3<sup>+</sup>PD-1<sup>+</sup> cells among CD4<sup>+</sup> or CD8<sup>+</sup> T cells.  
NOTE: Based on experimental results, the percentage of Foxp3<sup>+</sup>PD-1<sup>+</sup> is more than 50% in both CD4<sup>+</sup> and CD8<sup>+</sup> T cell populations at 16 d p.i. with LCMV CL13.  
1. To analyze the frequency and phenotypes of  $T_{reg}$  cells, examine the frequencies of Foxp3<sup>+</sup> or Foxp3<sup>+</sup>PD-1<sup>+</sup> cells among CD4<sup>+</sup> T cells.  
NOTE: Based on experimental results, the percentage of Foxp3<sup>+</sup> or Foxp3<sup>+</sup>PD-1<sup>+</sup> cells is more than 20% in CD4<sup>+</sup> T cell population, respectively.

### 4. Isolation of CD4<sup>+</sup>CD25<sup>+</sup> $T_{reg}$ Cells

NOTE: The volumes of all reagents indicated below are for a starting cell number of  $1 \times 10^7$  total splenocytes.

1. Prepare the cells as described in section 2 using naive and LCMV chronically infected mice. Wash the cells by adding 10 ml of T cell isolation buffer. Centrifuge at 300 x g for 10 min at 4 °C. Discard the supernatant completely and resuspend the cell pellet in 40 µl of buffer.
2. For CD4<sup>+</sup> T cell enrichment using the magnetic cell separation system, add 10 µl of biotin-antibody cocktail and mix well. Incubate for 10 min at 4 °C.
3. Add 30 µl of buffer, 20 µl of anti-biotin micro-beads for labeling of non-CD4<sup>+</sup> T cells and 10 µl of CD25-PE antibody for fluorescent labeling of CD25<sup>+</sup> cells. Mix well and incubate for 15 min in the dark.
4. Add 2 ml of buffer and pass the cells through a 40 µm cell strainer into a new 50 ml tube to remove cell debris. Wash the cells by centrifugation at 300 x g for 10 min at 4 °C. Discard the supernatant completely and resuspend the cell pellet in 500 µl buffer at a density of up to 1.25 x 10<sup>6</sup> cells.
5. Apply the cells onto the column and collect the unlabeled cells that pass through the column. Wash the column by adding 2 ml of buffer and centrifuge at 300 x g for 10 min at 4 °C. Discard the supernatant completely and resuspend the isolated CD4<sup>+</sup> T cells in 90 µl of buffer.
6. For magnetic labeling of CD25<sup>+</sup> cells, add 10 µl of anti-PE microbeads, mix well, and incubate for 15 min in the dark at 4 °C.
7. Add 2 ml of buffer and wash the cells by centrifugation at 300 x g for 10 min at 4 °C. Discard the supernatant completely and resuspend the cell pellet in 500 µl of buffer at a density of up to 1 x 10<sup>6</sup> cells.
8. To enrich the CD4<sup>+</sup>CD25<sup>+</sup> T<sub>reg</sub> cells, apply the cells onto the column for positive selection, and wash the column by adding 2 ml of buffer. Repeat the wash three times.
9. When the column is empty after the final washing step, add 1 ml of buffer onto the column, and flush out the magnetically labeled CD4<sup>+</sup>CD25<sup>+</sup> cells using the plunger.
10. Repeat steps 4.8-4.9 to enhance the purity of the isolated CD4<sup>+</sup>CD25<sup>+</sup> T<sub>reg</sub> cells. Wash the cells with FACS buffer by centrifugation at 300 x g for 10 min at 4 °C. Discard the supernatant and resuspend the isolated CD4<sup>+</sup>CD25<sup>+</sup> T<sub>reg</sub> cells at a concentration of 2 x 10<sup>6</sup> cells/ml in complete RPMI media.
11. To check the purity of the isolated-CD4<sup>+</sup>CD25<sup>+</sup> T<sub>reg</sub> cells, use 2 x 10<sup>5</sup> cells from the total isolated-CD4<sup>+</sup>CD25<sup>+</sup> T<sub>reg</sub> cells. Stain the cells with 50 µl of FACS buffer containing anti-CD4 FITC and cell viability detecting reagent near-IR fluorescent reactive dye.  
NOTE: CD25 was already labeled with PE during isolation.
12. Incubate for 20 min in the dark at 4 °C. Wash the cells with FACS buffer by centrifugation at 300 x g for 2 min at 4 °C. After washing, resuspend the cell pellet in 100 µl of FACS buffer and check the purity by flow cytometry.
13. Gate the live-cell population. To analyze the purity of T<sub>reg</sub> cells, confirm the percentage of CD4<sup>+</sup>CD25<sup>+</sup> cells.  
NOTE: Based on experimental results, the percentage of CD4<sup>+</sup>CD25<sup>+</sup> cells among purified cells is over about 80%.

## 5. Isolation of CD8<sup>+</sup> T Cells and Labeling of CD8<sup>+</sup> T Cells

NOTE: The volumes of all reagents indicated below are for a starting cell number of 1 x 10<sup>7</sup> total splenocytes.

1. Prepare cells as described in section 2 using naïve mice. Wash the cells by adding 10 ml of T cell isolation buffer. Centrifuge at 300 x g for 10 min at 4 °C. Discard the supernatant completely, and resuspend the cell pellet in 40 µl of buffer.
2. For CD8<sup>+</sup> T cell isolation using the magnetic cell separation system, add 10 µl of biotin-antibody cocktail for non-CD8<sup>+</sup> T cell labeling and mix well. Incubate for 5 min at 4 °C.
3. Add 30 µl of buffer and 20 µl of anti-biotin microbeads. Mix well and incubate for 10 min at 4 °C.
4. Apply the cells onto the column and collect the unlabeled cells that pass through the column. Wash the column by adding 2 ml of buffer three times.
5. Centrifuge at 300 x g for 10 min at 4 °C. Discard the supernatant completely and resuspend the isolated CD8<sup>+</sup> T cells in 2 ml of buffer.
6. To check the purity of the isolated cells, prepare 50 µl of antibody solution in FACS buffer. Resuspend the 2 x 10<sup>5</sup> cells among isolated CD8<sup>+</sup> T cells in 50 µl of antibody solution.  
NOTE: To prepare antibody solution, add anti-CD8 FITC, and cell viability detection reagent (near-IR fluorescent reactive dye) into FACS buffer.
7. Incubate for 20 min in the dark at 4 °C. Wash the cells with FACS buffer by centrifugation at 300 x g for 2 min at 4 °C. After washing, resuspend the cell pellet in 100 µl of FACS buffer, and check the purity of the cells by flow cytometry.
8. Gate the live-cell population. To check the purity of CD8<sup>+</sup> T cells, confirm the percentage of CD8<sup>+</sup> T cells.  
NOTE: Based on experimental results, the percentage of CD8<sup>+</sup> cells among purified cells is over about 90%.
9. Add PBS to the isolated CD8<sup>+</sup> T cells. Centrifuge at 300 x g for 10 min at 4 °C. Discard the supernatant. Resuspend the isolated CD8<sup>+</sup> T cells at a concentration of 1 x 10<sup>7</sup> cells/ml in PBS.
10. To label the CD8<sup>+</sup> T cells for the *in vitro* suppression assay, dilute cell proliferation tracking violet dye in PBS to obtain a concentration of 5 µM at RT.  
NOTE: The approximate excitation and emission peaks of the cell proliferation tracking violet dye used in the study are 405 and 450 nm, respectively.
11. Mix well equal volumes of cell proliferation tracking violet dye (5 µM) and cell suspension (1 x 10<sup>7</sup> cells/ml of CD8<sup>+</sup> T cells) in a 15 ml tube, and incubate at 20 min at 37 °C. Vortex the tube every 10 min.
12. Fill up the tube with cold complete RPMI media, and leave the tube for 10 min at RT. Centrifuge at 300 x g for 10 min at RT. Discard the supernatant completely, and resuspend the cells at a concentration of 2 x 10<sup>6</sup> cells/ml with pre-warmed complete RPMI media. Incubate the cells for 15 min at RT.

## 6. Setting Up the *In Vitro* Suppression Assay Using CD4<sup>+</sup>CD25<sup>+</sup> T<sub>reg</sub> and CD8<sup>+</sup> T Cells

1. To prepare anti-CD3/CD28-coated beads, transfer the appropriate volume of magnetic beads to a 15 ml of tube (2.5 µl/1 x 10<sup>5</sup> cells). Add an equal volume of PBS and mix. Wash by centrifugation at 300 x g for 2 min at 4 °C and discard the supernatant. Dilute the magnetic beads in complete media (50 µl/well).
2. Aliquot 50 µl of CD4<sup>+</sup>CD25<sup>+</sup> T<sub>reg</sub> cells per well of u-bottom 96-well plate (1 x 10<sup>5</sup> cells/well). Add 50 µl of CD8<sup>+</sup> T cells as responder T (T<sub>resp</sub>) cells per well (1 x 10<sup>5</sup> cells/well). Add 50 µl of diluted anti-CD3/CD28-coated beads into per well.

NOTE: In this step, label and set up control wells as follows: "unstimulated CD8<sup>+</sup> T cell only" with no anti-CD3/CD28-coated beads; "CD8<sup>+</sup> T cell only" with anti-CD3/CD28-coated beads; "CD8<sup>+</sup> T cell only" with anti-CD3/CD28-coated beads; "T<sub>reg</sub> cell only" with anti-CD3/CD28-coated beads. T<sub>reg</sub> cells can be diluted by complete media and co-cultured with T<sub>resp</sub> cells in a different ratio of T<sub>resp</sub> cells:T<sub>reg</sub> cells (1:0.25-1:1).

3. Add 50  $\mu$ l or appropriate volume of media into all wells to total volume of 200  $\mu$ l. Cover the plate with foil and incubate in a CO<sub>2</sub> incubator at 37 °C for 72 hr.

## 7. Analysis of CD8<sup>+</sup> T Cell Proliferation & Cytokine Production from CD8<sup>+</sup> T Cells

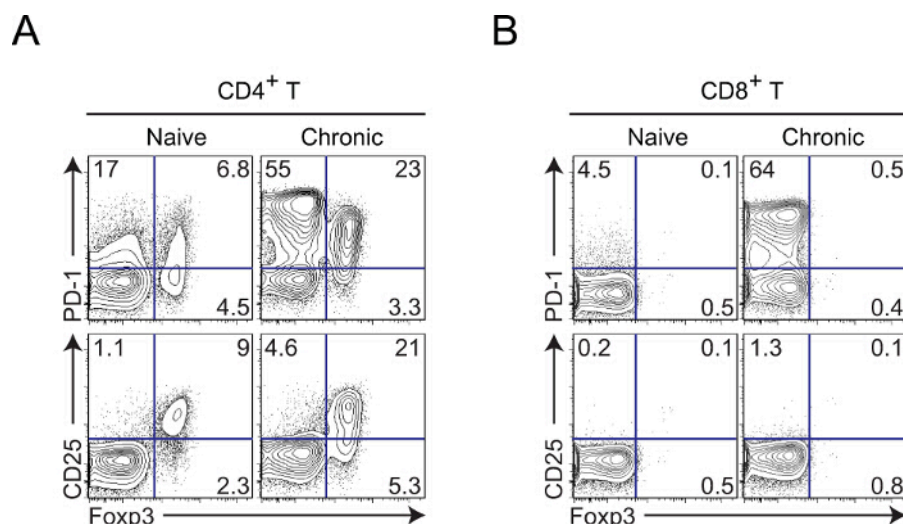
1. For the cytokine production analysis, after 3 days of culture, separate the supernatant of each well into another plate and perform enzyme-linked immunosorbent assay (ELISA).  
NOTE: The supernatant may be aliquoted and stored at -70 °C. In this experiment, anti-mouse IFN- $\gamma$  antibody-coated plate was used to detect IFN- $\gamma$  production according to the manufacturer's protocol. To determine IFN- $\gamma$  production of proliferating CD8<sup>+</sup> T cells on a single cell level, intracellular cytokine staining can be performed.
2. After separating the supernatant from each well, wash the plate containing the cells with FACS buffer and centrifuge at 300 x g for 2 min at 4 °C (3 times).
3. After washing, discard the supernatant. Resuspend the cell pellet with 50  $\mu$ l of antibody cocktail for staining of proliferated CD8<sup>+</sup> T cells. Incubate for 20 min in the dark at 4 °C.  
NOTE: To prepare antibody cocktail, add anti-CD4 FITC, anti-CD8 PerCP-Cy5.5, and cell viability detection reagent (near-IR fluorescent reactive dye) into FACS buffer.  
NOTE: Antibodies against various markers such as CD44 or CD69 can be combined with other antibodies to confirm activation of CD8<sup>+</sup> T cells. Remember that CD8<sup>+</sup> T cells have already been labeled with cell proliferation tracking violet dye at Step 5.10.
4. Wash twice by centrifugation at 300 x g for 2 min at 4 °C. After the final washing step, discard the supernatant, and fix the cells for 20 min in the dark at 4 °C with 100  $\mu$ l of fixation buffer.
5. Wash twice by centrifugation at 300 x g for 2 min at 4 °C. Resuspend the cells with 200  $\mu$ l of FACS buffer and measure the proliferation of cell proliferation tracking violet dye-labeled CD8<sup>+</sup> T cells by flow cytometry.
  1. Gate the CD8<sup>+</sup> T cell population among the live cells. Measure the percentages of divided and undivided cells according to the dilution of the cell proliferation tracking violet dye and CD8<sup>+</sup> T cell dilution according to the following equation. % Inhibition = [(% of proliferated CD8<sup>+</sup> T cells in the absence of T<sub>reg</sub> cells - % of proliferated CD8<sup>+</sup> T cells in the presence of T<sub>reg</sub> cells)/(% of proliferated CD8<sup>+</sup> T cells in the absence of T<sub>reg</sub> cells)] x 100. Further analyze the data by using the flow cytometry software<sup>25</sup>.

## Representative Results

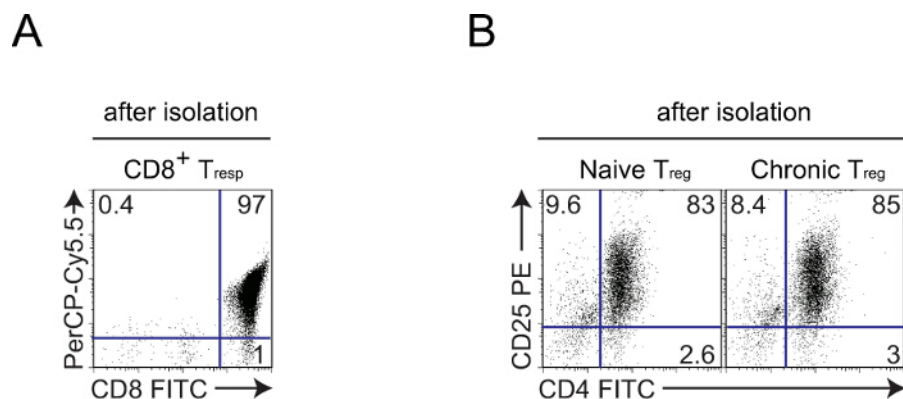
We generated mice with persistent virus infection by injecting them with  $2 \times 10^6$  p.f.u. of LCMV CL13 intravenously. To investigate the phenotypic changes in T<sub>reg</sub> cells and T<sub>conv</sub> cells during chronic virus infection, splenic lymphocytes obtained from naïve and infected mice were stained with various antibodies and analyzed by flow cytometry. At 16 d p.i., PD-1 was upregulated in both Foxp3<sup>+</sup>CD4<sup>+</sup> T<sub>conv</sub> (**Figure 1A**, upper panel) and Foxp3<sup>+</sup>CD8<sup>+</sup> T<sub>conv</sub> (**Figure 1B**, upper panel) cells. The frequency of Foxp3<sup>+</sup>CD4<sup>+</sup> T<sub>reg</sub> cells was two times higher in the LCMV CL13-infected mice than in the naïve mice (**Figure 1A**). In particular, most of the Foxp3<sup>+</sup>CD4<sup>+</sup> T<sub>reg</sub> cells displayed the activated phenotype, expressing high levels of PD-1 at this time point (**Figure 1A**).

For the *in vitro* suppression assay, a considerable number of CD8<sup>+</sup> T<sub>conv</sub> cells and CD4<sup>+</sup>CD25<sup>+</sup> T<sub>reg</sub> cells were required. To obtain  $1 \times 10^7$  cell proliferation tracking violet dye-labeled CD8<sup>+</sup> T cells, splenocytes from two spleens of naïve mice were pooled. To obtain  $2 \times 10^6$  or  $3 \times 10^6$  of CD4<sup>+</sup>CD25<sup>+</sup> T<sub>reg</sub> cells, at least three spleens from naïve and LCMV CL13-infected mice, respectively, were pooled. CD8<sup>+</sup> T cells as T<sub>resp</sub> cells were separated successfully without significant contamination with other immune cells (**Figure 2A**). T<sub>reg</sub> cells could also be isolated, with a dominant population of CD25<sup>+</sup>CD4<sup>+</sup> T cells with more than 80% purity (**Figure 2B**).

To compare the suppressive function of naïve and chronic T<sub>reg</sub> cells, the isolated T<sub>reg</sub> and T<sub>resp</sub> cells were incubated together under stimulation with anti-CD3/CD28-coated beads at 37 °C for 3 days. % inhibition of CD8<sup>+</sup> T<sub>resp</sub> cell proliferation was increased in a T<sub>reg</sub> cell dose-dependent manner (**Figure 3A**). When CD8<sup>+</sup> T<sub>resp</sub> cells were co-cultured with chronic T<sub>reg</sub> cells in a ratio of 1:1, proliferation of CD8<sup>+</sup> T<sub>resp</sub> cells were significantly inhibited (**Figure 3B**). IFN- $\gamma$  production by CD8<sup>+</sup> T<sub>resp</sub> cells were also significantly inhibited when CD8<sup>+</sup> T<sub>resp</sub> cells were co-cultured with activated T<sub>reg</sub> cells from chronically infected mice rather than with resting T<sub>reg</sub> cells from naïve mice in a T<sub>reg</sub> cell dose-dependent manner (**Figure 3C**).



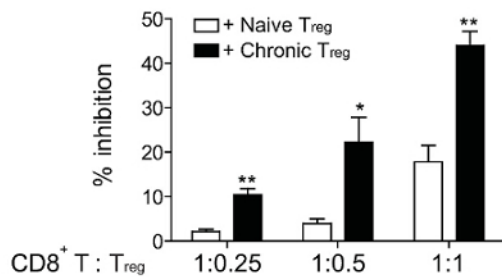
**Figure 1: Phenotypes of CD4<sup>+</sup> and CD8<sup>+</sup> T cells in the spleen of naïve mice or LCMV CL13-infected mice at 16 d p.i.** (A) Expression of PD-1 or CD25 on Fxp3<sup>+</sup>CD4<sup>+</sup> T<sub>conv</sub> and Fxp3<sup>+</sup>CD4<sup>+</sup> T<sub>reg</sub> cells. (B) Expression of PD-1 or CD25 on Fxp3<sup>+</sup>CD8<sup>+</sup> T<sub>conv</sub> cells. Splenic lymphocytes were stained with antibodies against CD4, CD8, CD25, PD-1, and Fxp3. Data are representative of three independent experiments. Panel A has been modified from<sup>25</sup> as a reference. [Please click here to view a larger version of this figure.](#)



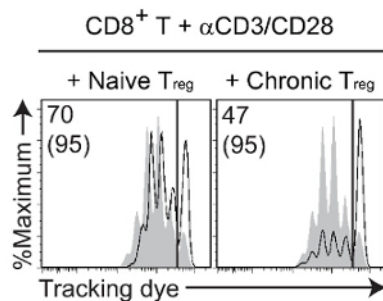
**Figure 2: Purities of T<sub>resp</sub> cells isolated from naïve mice and T<sub>reg</sub> cells isolated from naïve or LCMV CL13-infected mice.** (A) Percentage of CD8<sup>+</sup> T cells after isolation. (B) Percentage of CD25-expressing CD4<sup>+</sup> T<sub>reg</sub> cells after isolation. Each quadrant in (B) was determined by the expression levels of CD4 and CD25. Data are representative of three independent experiments. [Please click here to view a larger version of this figure.](#)



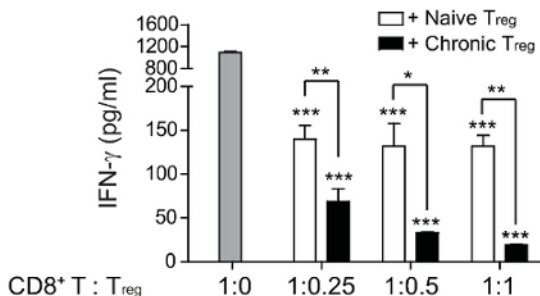
A



B



C



**Figure 3: Effect of activated T<sub>reg</sub> cells on CD8<sup>+</sup> T cell response.** (A) % inhibition of CD8<sup>+</sup> T cells co-cultured with T<sub>reg</sub> cells in a T<sub>reg</sub> cell dose-dependent manner. (B) Proliferation of CD8<sup>+</sup> T cells co-cultured with T<sub>reg</sub> cells in 1:1 ratio. (C) IFN-γ production of CD8<sup>+</sup> T cells co-cultured with T<sub>reg</sub> cells in a T<sub>reg</sub> cell dose-dependent manner. Proliferation of CD8<sup>+</sup> T cells was measured by dilution of cell proliferation tracking violet dye in proliferated CD8<sup>+</sup> T cells and IFN-γ secretion was evaluated by ELISA. Cell proliferation tracking violet dye-labeled CD8<sup>+</sup> T cells were stimulated with anti-CD3/CD28-coated beads for 3 days in the absence or presence of T<sub>reg</sub> cells isolated from either naive or chronically infected mice. In the histogram, the numbers without and with parenthesis indicate the proliferation percentage of CD8<sup>+</sup> T cells co-cultured with and without T<sub>reg</sub> cells, respectively. Filled gray peaks in the histogram indicate proliferation of CD8<sup>+</sup> T cells co-cultured without T<sub>reg</sub> cells. Each group was designed in triplicates. The bars represent mean + SEM. This figure has been modified from<sup>25</sup> as a reference. [Please click here to view a larger version of this figure.](#)

## Discussion

Although only a small number of T<sub>reg</sub> cells exist in mice and humans, it is important to understand their function as they play a crucial role in regulating the immune response and maintaining immune tolerance. The number and suppressive functions of T<sub>reg</sub> cells increases during a chronic virus infection<sup>15-20</sup> as well as cancer progression<sup>13,14</sup>. This is probably due to continued antigen stimulation. To evaluate the T<sub>reg</sub> cells function under antigen persistence and disease development, their suppressive activity needs to be measured.

Here, we describe a protocol to analyze the *ex vivo* phenotype of T<sub>reg</sub> cells and measure their suppressive activity using an *in vitro* co-culture system of CD8<sup>+</sup> T cells and T<sub>reg</sub> cells. Critical steps in the current protocol are analysis and isolation of T<sub>reg</sub> cells. Live and freshly isolated T<sub>reg</sub> cells show clear suppressive activity *in vitro*. We also examined the phenotypes of T<sub>conv</sub> and T<sub>reg</sub> cells during a chronic virus infection. T<sub>conv</sub> cells, *i.e.*, Foxp3 CD4<sup>+</sup> and CD8<sup>+</sup> T cells showed a decrease in cellular activity after the chronic virus infection (Figure 1). The PD-1 expression was upregulated in both the T<sub>conv</sub> cell populations. Increase in T<sub>reg</sub> cell number and up-regulation of PD-1 expression are the hallmarks of immune

response during a chronic virus infection. Additionally, anti-bodies for other inhibitory receptors such as TIM-3 and CTLA-4 can be used in combination with PD-1 to examine the T cell phenotype by FACS after chronic virus infection.

In order to investigate the suppressive activity,  $T_{resp}$  cells and  $T_{reg}$  cells were co-cultured at a ratio of 1 to 1. Co-culture with  $T_{reg}$  cells from chronically infected mice significantly reduced  $CD8^+$  T cell proliferation and IFN- $\gamma$  secretion when compared to  $T_{reg}$  cells from naïve mice. This implies that the proliferation and cytokine production of  $CD8^+$  T cells co-cultured with  $T_{reg}$  cells are inversely related to the suppressive function of  $T_{reg}$  cells. Although this protocol recommends the use of a magnetic cell separation system for  $T_{reg}$  cell isolation, contamination with non- $T_{reg}$  cells cannot be ruled out. The  $T_{reg}$  cells obtained from Foxp3-GFP reporter mice<sup>28-30</sup> are better candidates to accurately investigate the function of  $T_{reg}$  cells than magnetic cell separation system, as CD25 is often overexpressed on activated  $CD4^+$   $T_{conv}$  as well as  $T_{reg}$  cells.

The suppressive function of  $T_{reg}$  cells has been evaluated by various *in vitro* suppression assays<sup>16,28,31-34</sup>. The advantage of the *in vitro* suppression assay is that it evaluates the direct effect of  $T_{reg}$  cells on the inhibition of the  $CD8^+$  T cell response, because only  $T_{resp}$  cells are co-cultured with  $T_{reg}$  cells. In addition to  $CD8^+$  T cells, the  $CD25^+$   $CD4^+$  T cells can be used instead of  $T_{resp}$  cells<sup>25</sup>. The effects of  $T_{reg}$  or  $T_{conv}$  cells on immune cells such as dendritic cells or natural killer cells can be evaluated by varying the experimental conditions.

Molecules such as the  $CD103^+$ , LAG-3<sup>36</sup>, glycoprotein A repetitions predominant<sup>37</sup>,  $CD43^+$ ,  $CD11a^+$ , or killer cell lectin-like receptor subfamily G member 1<sup>38</sup> have been used as markers of activated  $T_{reg}$  cells in specific diseases. In this protocol, we used PD-1 as an indicator to identify activated  $T_{reg}$  cells during chronic virus infection. This protocol can be used for multifaceted analyses (phenotypic and functional) of  $T_{reg}$  cells under specific conditions.

## Disclosures

S.-J.H. has a patent and receives patent royalties related to the PD-1 pathway. The other authors have no financial conflicts of interest.

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