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Original Article

Rapid G0/1 transition and cell cycle progression in CD8⁺ T cells compared to CD4⁺ T cells following in vitro stimulation

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Running title:

CD4⁺ versus CD8⁺ T cell proliferation

Subject Section : Immunology

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List of Abbreviations:

CFSE, carboxyfluorescein diacetate succinimidyl ester; CTL, cytotoxic T-lymphocyte; FBS, fetal bovine serum; IL-2, Interleukin-2; LN, lymph node; MHC, major histocompatibility complex; mAb, monoclonal antibody; PI, propidium iodide; PKC, protein kinase C; PMA, phorbol 12-myristate 13-acetate; SPL, spleen; TCR, T cell receptor; 7-AAD, 7-amino-actinomycin D

ABSTRACT

T cell population consists of two major subsets, CD4⁺ T cells and CD8⁺ T cells, which can be distinguished by the expression of CD4 or CD8 molecules, respectively. Although they play quite different roles in an immune system, many of their basic cellular processes such as proliferation following stimulation are presumably common. In this study, we have carefully analyzed time course of G0/1 transition as well as cell cycle progression in the two subsets of quiescent T cell population following *in vitro* growth stimulation. We found that CD8⁺ T cells promote G0/1 transition more rapidly and drive their cell cycle progression faster compared to CD4⁺ T cells. In addition, expression of CD25 and effects of its blockade revealed that IL-2 is implicated in the rapid progression, but not the earlier G0/1 transition, of CD8⁺ T cells.

Keywords:

Cell cycle, G0/1 transition, IL-2, T cell proliferation

INTRODUCTION

T cells express a T cell receptor (TCR) by which they recognize an antigen. During development, each T cell progenitor rearranges the genome and expresses the TCR with distinct specificity. Although this diversity of T cells allows them to deal with a wide range of antigens, only a tiny population of T cells can respond to a certain antigen (1). To overcome this insufficiency, they must proliferate to form a bunch of effector T cells. T cells are divided into two major subsets, which are distinguished by the expression of either CD4 or CD8 molecules. Following activation, CD4⁺ T cells function in producing cytokines, while CD8⁺ T cells show cytotoxic effects, indicating their different roles in an immune system. CD4⁺ and CD8⁺ T cells develop from common progenitor cells and recognize the antigen by the TCR in the similar context of major histocompatibility complex (MHC). Accordingly, it is assumed that both CD4⁺ and CD8⁺ T cells employ common mechanisms to regulate most parts of cellular processes including proliferation. T cell proliferation has been traditionally examined by an incorporation of ³H-TdR, which is a precursor of DNA and its usage reflects the DNA synthesis. This technique does not provide much information about the cell cycle progression. In addition, it is not applicable if CD4⁺ and CD8⁺ T cells are in the same culture and required to be analyzed separately. Using carboxyfluorescein diacetate succinimidyl ester (CFSE)-labeled T

cells, Foulds et al. showed that CD8⁺ T cells proliferate more extensively than CD4⁺ T cells in both *in vitro* and *in vivo* (2). Thus, the ability to proliferate seems to be different between CD4⁺ and CD8⁺ T cells, but their cell cycle progressions remain to be carefully compared.

A certain TCR binds to the antigen MHC complex with intrinsic binding affinity and induces various responses in the activated T cells via distinct intra-cellular signaling both qualitatively and quantitatively (3). On the other hand, intra-cellular signaling cascades following the TCR engagement has been relatively clear (4) and the most part of these pathways can be artificially activated by chemical stimulants. One of such stimulants is a combination of phorbol 12-myristate 13-acetate (PMA) and ionomycin (5), stimulating protein kinase C (PKC) and Ca²⁺ influx, respectively. Thus, PMA plus ionomycin activates both CD4⁺ and CD8⁺ T cells strongly in an antigen-independent manner. Furthermore, for the optimal proliferation, both CD4⁺ and CD8⁺ T cells require additional signals derived from costimulatory molecules (6). CD28 is expressed on both T cell subsets and is known to function as a major costimulator (7). A monoclonal antibody (mAb) against CD28 is available and it can provide the T cells sufficient costimulatory signal even under the limited condition of costimulatory ligands.

In this study, we have tried to dissect the proliferation of CD4⁺ and CD8⁺ T cells into

the entry and the subsequent progression of the cell cycle and compare their ability following the same strong stimulation by PMA plus ionomycin and anti-CD28 mAb. We found that CD8⁺ T cells initiate G0/1 transition earlier and drive cell cycle progression faster than CD4⁺ T cells. This rapid proliferation of CD8⁺ T cells was partly due to IL-2, since the blockade of IL-2 receptor signaling reduced the proliferation of CD8⁺ T cells, but not CD4⁺ T cells, without affecting the G0/1 transition. One of most important role of CD8⁺ T cells is to deal with a serious threat of virulent viruses. CD8⁺ T cells might be required to expand quickly compared to CD4⁺ T cells to provide a rapid containment of viruses vigorously expanding following infection.

MATERIALS AND METHODS

T Cell Culture and CFSE analysis

Lymph nodes (LNs: cervical, axillary, brachial, inguinal, mesenteric, periaortic and pancreatic) were isolated from ddY mice aged 8-10 weeks. The mice had been euthanized according to the guidelines of the Animal Care and Use Committee in Kanazawa University. The lymph nodes were crushed to make a single cell suspension (8). LN cells were labeled for 15 min at 37°C with 2 μ M CFSE (Molecular Probes) to track mitotic divisions of T cells following stimulation. After 3 washes with PBS containing 5% heat-inactivated fetal bovine serum (FBS), CFSE-labeled LN cells were inoculated at 1.5×10^6 cells/ml along with the same density of non-labeled splenocytes prepared from the same mice. The cells were co-cultured in RPMI1640 containing 10% FBS supplemented with gentamycin (10 μ g/ml) and stimulated with PMA (2 ng/ml, Sigma-Aldrich) plus ionomycin (20 ng/ml, Sigma-Aldrich). In some cases, T cells were stimulated by anti-TCR β mAb (H57-597, 1 μ g/ml, BioLegend) instead. To provide a costimulatory signal, anti-CD28 mAb from hybridoma PV1 (9) was added to the culture at a concentration of 2 μ g/ml. The cells were harvested from the culture at various times after stimulation and first treated with anti-Fc γ R II/III mAb from hybridoma 2.4G2 (10) to reduce non-specific antibody binding. The cells were then stained with biotin-labeled

anti-CD4 mAb (GK1.5, BioLegend) or anti-CD8 mAb (53-6.7, BioLegend) followed by either PE-labeled (eBioscience) or APC-labeled (SouthernBiotech) streptavidin to distinguish two major T cell populations. The cells were finally treated with 7-amino-actinomycin D (7-AAD, BD Biosciences) or propidium iodide (PI, Sigma) to exclude dead cells from the analysis. The CFSE profile of each population was obtained via flow cytometry using FACSverse (Becton Dickinson) and the mean division number was determined by calculating percent contribution of the initial cohort in each division peak. Initiation of the second and the third cell cycle was estimated by the appearance of the cell division peak that was recognized by the CFSE dilution.

Estimation of the G0/1 transition of quiescent T cells using Pyronin Y staining

In order to determine the transition from G0 to G1, RNA transcription was detected using Pyronin Y (Sigma) staining. The cells were fixed with 4% paraformaldehyde for 15 min and then the cell membrane was permeabilized with 0.02% Saponin (ICN Biomedicals) for 20 min. After that, the cells were treated with 100 µg/ml Pyronin Y for 10 min before FACS analysis.

Expression of IL-2 receptor and blockade of IL-2 signaling

Expression of high affinity IL-2 receptor was determined using an antibody against CD25. CD25 is a part of the IL-2 receptor complex and the one that is responsible to increase binding affinity for IL-2 (11). The cells harvested were stained with anti-mouse CD25 mAb (PC61, IMGENEX) followed by Alexa647-labeled anti-Rat IgG(H+L) Antibody (Invitrogen). CD4 or CD8 was also stained as described above and CD25 expression by those T cells was separately analyzed via FACS. This anti-mouse CD25 mAb has been shown to block binding of IL-2 to the receptor (12), thereby inhibits IL-2 signaling. For this purpose, 10 µg/ml of anti-mouse CD25 mAb was added to the culture at the time of inoculation.

Statistics

Statistical tests were performed using Student's independent *t* test. The results were considered to be significant at $p < 0.05$.

RESULTS

The proliferation of CD8⁺ T cells is greater than that of CD4⁺ T cells

CFSE is a dye that is widely used to examine cell division. Since it is decreased by half in every cell division, it allows us to evaluate the proliferation by calculating the mean division number. Using this strategy, we examined the proliferation at 48 hr after stimulation *in vitro* to see whether any difference between CD4⁺ and CD8⁺ T cells exists (Fig.1). Both T cells were placed in the same culture and stimulated simultaneously, and yet CFSE profiles clearly demonstrated that more cell divisions occurred in CD8⁺ T cells than in CD4⁺ T cells (Fig.1a). The mean division number of CD8⁺ T cells calculated using the CFSE profile was indeed 2-fold higher than that of CD4⁺ T cells (Fig.1b). This greater proliferation of CD8⁺ T cells was also observed when they were stimulated by anti-CD3 mAb (2) or anti-TCR β mAb (Fig.1c), indicating that the stimulation with PMA plus ionomycin correctly reproduces T cell proliferation. Because the T cell population remaining in an undivided state affects the mean division number, we determined the sequential change of the undivided population and the mean division number as well (Fig.2). At 24 hr after stimulation, there was no cell division. Since then, the mean division number gradually increased in both CD4⁺ and CD8⁺ T cells and the latter kept higher than the former up to 60 hr after

stimulation (Fig.2a). In association with this increase, the undivided population was gradually decreased, but slightly larger population was always left behind in the CD4⁺ T cells (Fig.2b). However, most of them were out of this state by 60 hr after stimulation, thus the undivided population was not the critical cause of the difference between CD4⁺ and CD8⁺ T cells. In fact, the mean division number determined in the proliferating cell population that excluded non-divided cells in CD8⁺ T cells showed a greater elevation rate than that in CD4⁺ T cells (Fig.2c). These results suggest that CD8⁺ T cells have the ability to drive the cell cycle faster than CD4⁺ T cells and this property is likely involved in their greater proliferation.

We have studied the G0/1 transition and the cell cycle progression of naive CD4⁺ T cells following stimulation with PMA plus ionomycin and anti-CD28 mAb (13). Using the same approach, we further examined the difference between CD4⁺ and CD8⁺ T cell proliferation. Firstly, we evaluated the entry into G1 from G0 phase by RNA synthesis (Fig.3a and 3b). The cells synthesizing RNA were detected as Pyronin Y positive (Fig.3a) and the minimum time that the first population in CD4⁺ or CD8⁺ T cells entered into G1 phase was determined by the increase of this population (Fig.3b). CD8⁺ T cells initiated RNA synthesis around 9 hr after stimulation, which is approximately one hour earlier than that of CD4⁺ T cells. Thus, the earlier G0/1 transition is also a part of the

mechanism that causes better proliferation of CD8⁺ T cells. Secondly, the onset of the second or the third cell cycle of CD8⁺ T cells was estimated by the appearance of CFSE peak that had half or a quarter of the initial fluorescence intensity (Fig.3c and 3d). From this analysis, we determined the minimum time that CD8⁺ T cells need to complete the first and second cell cycles and they were approximately 2 and 3 hr shorter than those of CD4⁺ T cells, respectively. These results suggest that CD8⁺ T cells are capable of driving each cell cycle in one hour less than the CD4⁺ T cells (13). This finding is consistent with the result of the proliferation rate determined using the mean division number described above (Fig.2c).

IL-2 participates in the greater cell cycle rate, but not the earlier G0/1 transition, in CD8⁺ T cells

Next, we investigated the role of IL-2 on the greater proliferation of CD8⁺ T cells. We first compared expression of CD25, which is a part of a high-affinity IL-2 receptor complex, between CD4⁺ and CD8⁺ T cells at 48 hr after stimulation (Fig.4a). Both CD4⁺ and CD8⁺ T cells expressed CD25 and their levels were not clearly different. From the role of IL-2 on T cell proliferation (14-17), it should work early after stimulation. Therefore, we examined the expression of CD25 until 48 hr (Fig.4b). The

CD25 expression reached its maximum around 24 hr after stimulation and was significantly higher on CD8⁺ T cells than on CD4⁺ T cells at this time point. The higher expression of CD25 on CD8⁺ T cells was also observed at 12 hr after stimulation, but beyond 24 hr, there was no significant difference between CD4⁺ and CD8⁺ T cells. Since CD8⁺ T cells showed higher expressions of CD25 at earlier time points, we further investigated whether CD8⁺ T cells begin to express CD25 earlier than CD4⁺ T cells (Fig.4c). Both CD4⁺ and CD8⁺ T cells began to express CD25 as early as 2 hr after stimulation and there was no significant difference between their onsets, while the expression level on CD8⁺ T cells became clearly higher around 10 hr after stimulation (Fig.4d). Given the fact that T cells enter into the cell cycle around 10 hr after stimulation (Fig.3b), these results suggest that IL-2 seems to participate in the cell cycle progression rather than the earlier G0/1 transition in CD8⁺ T cells. In order to challenge this hypothesis, we took advantage of anti-CD25 mAb (PC61), which is capable to block binding of IL-2 to the receptor (12). It was added to the culture and the effect on the G0/1 transition of CD4⁺ or CD8⁺ T cells was examined (Fig.5). The time of the G0/1 transition was not changed in both T cells, which is consistent with the idea that IL-2 is not responsible for the earlier entry of CD8⁺ T cells into G1 phase. In contrast, 10 µg/ml of anti-CD25 mAb reduced proliferation of CD8⁺ T cells, but not CD4⁺ T

cells, significantly at 48 hr (Fig.6). These results suggest that the cell cycle progression of CD8⁺ T cells is at least partly dependent on IL-2.

DISCUSSION

In this study, we have characterized the cell cycle progression to determine the cellular basis in the greater proliferation of CD8⁺ T cells compared to CD4⁺ T cells. We have found that CD8⁺ T cells not only drive the cell cycle progression faster, but also enter into the cell cycle state earlier than CD4⁺ T cells. Foulds et al. suggested that the intrinsic mechanism is involved in the difference between CD4⁺ and CD8⁺ T cell proliferation (2). In this regard, the G0/1 transition was not affected by the IL-2 blockade, while it reduced subsequent proliferation in CD8⁺ T cells. Moreover, these findings also suggest that the T cell proliferation is regulated at multiple levels.

There are memory T cells as well as naive T cells in the LN cells. In this study, we determined the minimum times that T cells initiate the G0/1 transition and drive the first and second cell cycle. Veiga-Fernandes et al. reported that the memory T cells exhibit greater proliferation than the naive T cells and CDK6 is involved in this difference (18). The CDK6 is a regulator of G1 phase progression and already preactivated in the resting memory T cells. Thus, we might compare the responses of memory T cells rather than naive T cells. In addition, CDK6 could be the key molecule involved in the difference between CD4⁺ and CD8⁺ T cells. Moreover, in the combined culture of the LN and SPL cells, regulatory cells, which regulate T cell proliferation, are also present. Such

regulatory cells are not just antigen presenting cells, but also Treg cells. Therefore, the difference in the proliferation between CD4⁺ and CD8⁺ T cells reported here is under the influence of these regulatory cells.

In the presence of the antibody against the IL-2 receptor, which can block its binding to the IL-2, the proliferation of CD8⁺ T cells was partially inhibited. In contrast, the same antibody did not affect the proliferation of CD4⁺ T cells in our experiment. Lowenthal et al. have shown that 20 nM of this antibody almost completely inhibits IL-2 binding, therefore 10 µg/ml (~67 nM) of the antibody used in this study is thought to be sufficient to eliminate the effects of IL-2 (12). Although it has been shown that IL-2 promotes the proliferation of CD4⁺ T cells (14, 17) as well as CD8⁺ T cells (14-16), it might not be very important for the CD4⁺ T cell proliferation (19, 20). Thus, CD8⁺ T cells are strictly dependent on the IL-2, which is presumably produced by the CD4⁺ T cells activated in close proximity.

Following activation, CD8⁺ T cells differentiate into effector cells, namely cytotoxic T-lymphocytes (CTLs), which destroy cells infected with viruses. It has been known that IL-2 has multiple effects on CD8⁺ T cells during anti-viral immunity. It not only enhances CD8⁺ T cell proliferation (14-16), but also promotes differentiation toward the CTL (15, 21-23). In order to minimize the collateral damage in our own cells, rapid

expansion of the CD8⁺ T cell clone is extremely important. Thus, CD8⁺ T cells enter into the cell cycle earlier and drive the subsequent progression faster than CD4⁺ T cells in the IL-2 independent and dependent mechanisms, respectively.

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DISCLOSURE

The authors have no conflicts of interest to disclose.

REFERENCES

1. Jenkins M.K., Moon J.J. (2012) The role of naive T cell precursor frequency and recruitment in dictating immune response magnitude. *J Immunol* **188**: 4135–40.
2. Foulds K.E., Zenewicz L.A., Shedlock D.J., Jiang J., Troy A.E., Shen H. (2002) Cutting edge: CD4 and CD8 T cells are intrinsically different in their proliferative responses. *J Immunol* **168**: 1528–32.
3. Gottschalk R.A., Hathorn M.M., Beuneu H., Corse E., Dustin M.L., Altan-Bonnet G., Allison J.P. (2012) Distinct influences of peptide-MHC quality and quantity on in vivo T-cell responses. *Proc Natl Acad Sci USA* **109**: 881–6.
4. Gorentla B.K., Zhong X.-P. (2012) T cell Receptor Signal Transduction in T lymphocytes. *J Clin Cell Immunol* **2012(S12)**: 005
DOI:10.4172/2155-9899.S12-005.
5. Cantrell D. (1996) T cell antigen receptor signal transduction pathways. *Annu Rev Immunol* **14**: 259–74.
6. Van Gool S.W., Vandenberghe P., de Boer M., Ceuppens J.L. (1996) CD80, CD86 and CD40 provide accessory signals in a multiple-step T-cell activation model. *Immunol Rev* **153**: 47–83.

7. McAdam A.J., Schweitzer A.N., Sharpe A.H. (1998) The role of B7 co-stimulation in activation and differentiation of CD4⁺ and CD8⁺ T cells. *Immunol Rev* **165**: 231–47.
8. Ando Y., Yasuoka C., Mishima T., Ikematsu T., Uede T., Matsunaga T., Inobe M. (2014) Concanavalin A-mediated T cell proliferation is regulated by herpes virus entry mediator costimulatory molecule. *In Vitro Cell Dev Biol Anim* **50**: 313–20.
9. Abe R., Vandenberghe P., Craighead N., Smoot D.S., Lee K.P., June C.H. (1995) Distinct signal transduction in mouse CD4⁺ and CD8⁺ splenic T cells after CD28 receptor ligation. *J Immunol* **154**: 985–97.
10. Unkeless J.C. (1979) Characterization of a monoclonal antibody directed against mouse macrophage and lymphocyte Fc receptors. *J Exp Med* **150**: 580–96.
11. Sharon M., Klausner R.D., Cullen B.R., Chizzonite R., Leonard W.J. (1986) Novel interleukin-2 receptor subunit detected by cross-linking under high-affinity conditions. *Science* **234**: 859–63.
12. Lowenthal J.W., Corthésy P., Tougne C., Lees R., MacDonald H.R., Nabholz M. (1985) High and low affinity IL 2 receptors: analysis by IL 2 dissociation rate and reactivity with monoclonal anti-receptor antibody PC61. *J Immunol* **135**: 3988–94.

13. Mishima T., Toda S., Ando Y., Matsunaga T., Inobe M. (2014) Rapid proliferation of activated lymph node CD4(+) T cells is achieved by greatly curtailing the duration of gap phases in cell cycle progression. *Cell Mol Biol Lett* **19**: 638–48.
14. Battaglia A., Buzzonetti A., Baranello C., Fanelli M., Fossati M., Catzola V., Scambia G., Fattorossi A. (2013) Interleukin-21 (IL-21) synergizes with IL-2 to enhance T-cell receptor-induced human T-cell proliferation and counteracts IL-2/transforming growth factor- β -induced regulatory T-cell development. *Immunology* **139**: 109–20.
15. Obar J.J., Molloy M.J., Jellison E.R., Stoklasek T.A., Zhang W., Usherwood E.J., Lefrançois L. (2010) CD4⁺ T cell regulation of CD25 expression controls development of short-lived effector CD8⁺ T cells in primary and secondary responses. *Proc Natl Acad Sci USA* **107**: 193–8.
16. Cousens L.P., Orange J.S., Biron C.A. (1995) Endogenous IL-2 contributes to T cell expansion and IFN- γ production during lymphocytic choriomeningitis virus infection. *J Immunol* **155**: 5690–9.

17. Schorle H., Holtschke T., Hünig T., Schimpl A., Horak I. (1991) Development and function of T cells in mice rendered interleukin-2 deficient by gene targeting. *Nature* **352**: 621–4.
18. Veiga-Fernandes H., Rocha B. (2004) High expression of active CDK6 in the cytoplasm of CD8 memory cells favors rapid division. *Nat Immunol* **5**: 31–7.
19. Jenkins M.K., Khoruts A., Ingulli E., Mueller D.L., McSorley S.J., Reinhardt R.L., Itano A., Pape K.A. (2001) In vivo activation of antigen-specific CD4 T cells. *Annu Rev Immunol* **19**: 23–45.
20. Khoruts A., Mondino A., Pape K.A., Reiner S.L., Jenkins M.K. (1998) A natural immunological adjuvant enhances T cell clonal expansion through a CD28-dependent, interleukin (IL)-2-independent mechanism. *J Exp Med* **187**: 225–36.
21. Pipkin M.E., Sacks J.A., Cruz-Guilloty F., Lichtenheld M.G., Bevan M.J., Rao A. (2010) Interleukin-2 and inflammation induce distinct transcriptional programs that promote the differentiation of effector cytolytic T cells. *Immunity* **32**: 79–90.
22. Bachmann M.F., Wolint P., Walton S., Schwarz K., Oxenius A. (2007) Differential role of IL-2R signaling for CD8+ T cell responses in acute and chronic viral infections. *Eur J Immunol* **37**: 1502–12.

23. Williams M.A., Tyznik A.J., Bevan M.J. (2006) Interleukin-2 signals during priming are required for secondary expansion of CD8⁺ memory T cells. *Nature* **441**: 890–3.

Figure legends

Fig. 1. CD8⁺ T cells proliferate better than CD4⁺ T cells *in vitro*. LN cells were labeled with CFSE and combined with SPL cells. The cells were stimulated with PMA (2 ng/ml) plus ionomycin (20 ng/ml) (a,b) or anti-TCR β mAb (1 μ g/ml) (c) in the presence of anti-CD28 mAb (2 μ g/ml) for 48 hr. (a) The CFSE profile of CD4⁺ or CD8⁺ T cells was separately analyzed via FACS and the representative profile is shown. The number indicated above each peak shows the division number being determined by the intensity of the CFSE fluorescence, which is decreased by half in every cell division. The dotted histogram represents the peak of unstimulated cells. (b) The mean division number was determined as described in the Materials and Methods and the average $\pm$ SEM from six independent experiments is shown. (c) The mean division number was determined as in (b) and the average $\pm$ SEM from three independent experiments is shown. * $p < 0.05$, significantly different.

Fig. 2. CD8⁺ T cells leave undivided state earlier and proliferate faster than CD4⁺ T cells. LN cells were labeled with CFSE and combined with SPL cells. The cells were stimulated with PMA (2 ng/ml) plus ionomycin (20 ng/ml) in the presence of anti-CD28 mAb (2 μ g/ml). At various times after stimulation, the cells were harvested and the

CFSE fluorescence of CD4⁺ (■) or CD8⁺ (○) T cells was separately analyzed via FACS. (a) The mean division number was determined as in Fig.1 and the average $\pm$ SEM from five independent experiments is shown. *p < 0.05, significantly different. (b) The population remaining in the undivided state was determined and the average $\pm$ SEM from five independent experiments except for data at 30 hr, which was performed once, is shown. *p < 0.05, significantly different. (c) The mean division number in the population excluding non-divided cells was determined. The average $\pm$ SEM from five independent experiments is plotted against time and the slope, which was calculated by fitting points to a linear regression curve, is shown.

Fig. 3. CD8⁺ T cells initiate G0/1 transition earlier and drive subsequent cell cycle faster than CD4⁺ T cells. LN cells were labeled with CFSE and combined with SPL cells. The cells were stimulated with PMA (2 ng/ml) plus ionomycin (20 ng/ml) in the presence of anti-CD28 mAb (2 μ g/ml). (a) The cells were stained by Pyronin Y at 6 or 14 hr after stimulation and CD4⁺ and CD8⁺ T cells were analyzed separately. The representative dot plots are shown. The cells in the gate presented in the dot plots are Pyronin Y positive. (b) The Pyronin Y positive population in CD8⁺ T cells (□) was determined at various time points via FACS as in CD4⁺ T cells (■) (13) and the

average $\pm$ SEM from three independent experiments is plotted against time. (c, d) The population in CD8⁺ T cells experienced cell division once (○) or twice (◇) was determined at various time points via FACS as in CD4⁺ T cells (●, ◆) (13) and the average $\pm$ SEM from three independent experiments is plotted against time.

Fig. 4. CD4⁺ and CD8⁺ T cells begin to express CD25 with the same kinetics, but its amount becomes larger on the latter cells since when they enter into the cell cycle. LN cells were labeled with CFSE and combined with SPL cells. The cells were stimulated with PMA (2 ng/ml) plus ionomycin (20 ng/ml) in the presence of anti-CD28 mAb (2 μ g/ml). The cells were harvested at 48 hr (a) or various times after stimulation (b-d) and stained with anti-CD25 mAb. CD25 expression on CD4⁺ or CD8⁺ T cells was separately analyzed via FACS. (a) The representative FACS profile is shown. The dotted histogram represents the staining of unstimulated cells. (b-d) The MFI (b, d) or the positive population (c) of CD25 was determined in CD4⁺ (■) or CD8⁺ (○) T cells and the average $\pm$ SEM from three independent experiments is shown. *p < 0.05, significantly different.

Fig. 5. IL-2 blockade does not change the time of the G0/1 transition in both CD4⁺ and

CD8⁺ T cells. LN cells were labeled with CFSE and combined with SPL cells. The cells were stimulated with PMA (2 ng/ml) plus ionomycin (20 ng/ml) along with anti-CD28 mAb (2 µg/ml) in the presence (○, □) or absence (●, ■) of anti-CD25 mAb (10 µg/ml). The Pyronin Y positive population in CD4⁺ or CD8⁺ T cells was separately determined at various time points via FACS as in Fig.3a and the average ± SEM from three independent experiments is shown.

Fig. 6. IL-2 blockade attenuates the proliferation of CD8⁺ T cells, but not CD4⁺ T cells.

LN cells were labeled with CFSE and combined with SPL cells. The cells were stimulated with PMA (2 ng/ml) plus ionomycin (20 ng/ml) along with anti-CD28 mAb (2 µg/ml) in the presence or absence of anti-CD25 mAb (10 µg/ml) for 48 hr. The mean division number was determined as in Fig.1 and the average ± SEM from four independent experiments is shown. *p < 0.05, significantly different.

Fig.1

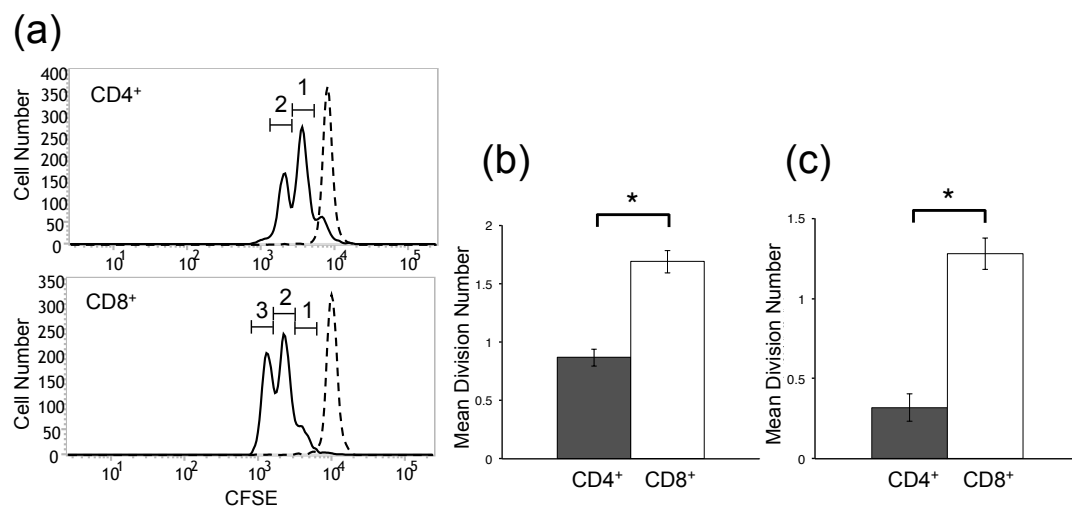


Fig.2

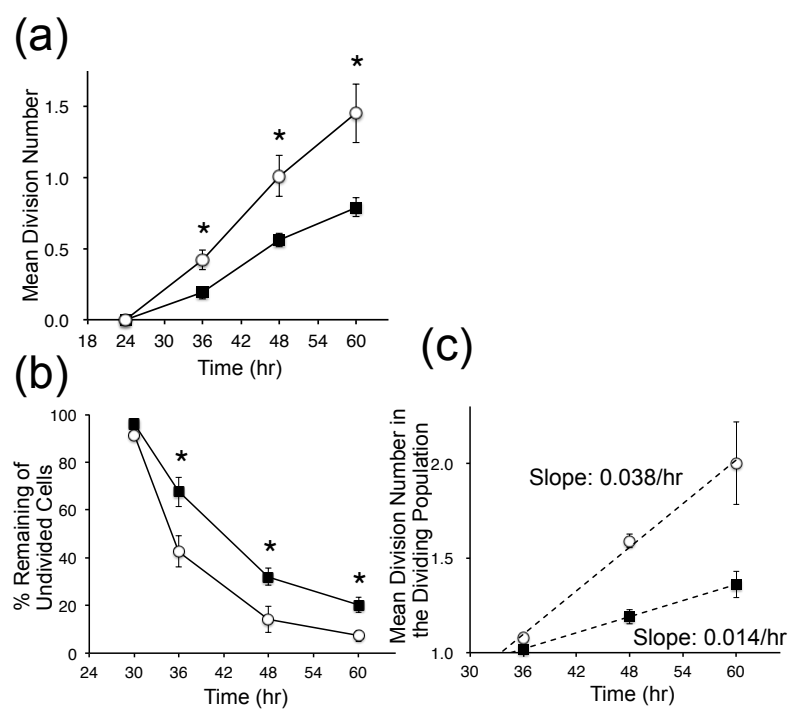


Fig.3

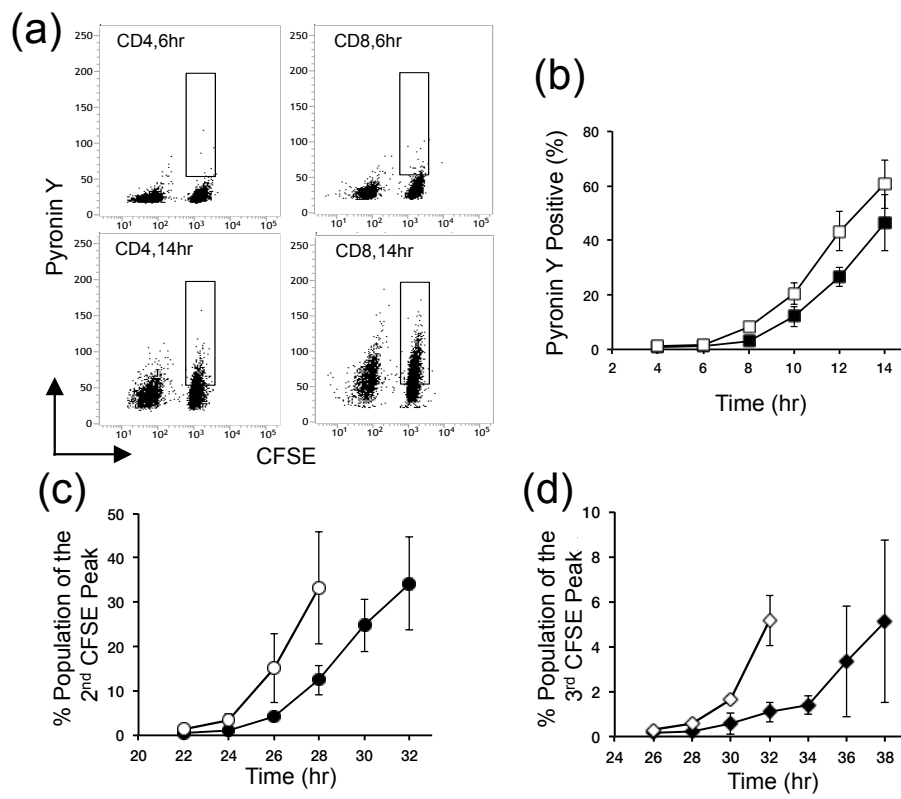


Fig.4

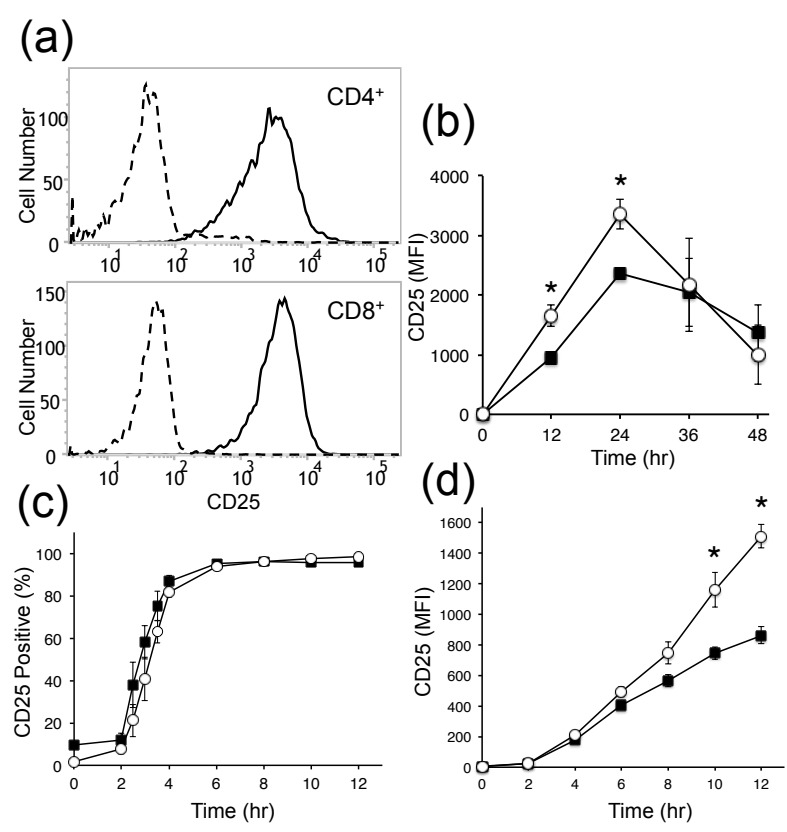


Fig.5

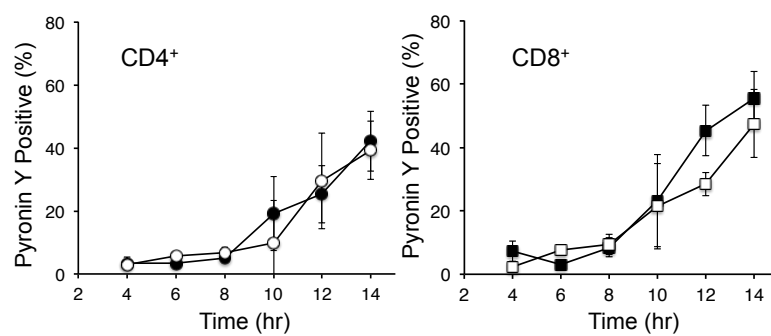


Fig.6

