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## Characterization of CTL Clones Specific for Single Antigen, H60 Minor Histocompatibility Antigen

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Background: Disparities of Minor Hantigens can induce graft rejection after MHC-matched transplantation. H60 has been characterized as a dominant antigen expressed on hematopoietic cells and considered to be an ideal model antigen for study on graft-versus-leukemia effect. Methods: Splenocytes from C57BL/6 mice immunized with H60 congenic splenocytes were used for establishment of H60-specific CTL clones. Then the clones were characterized for proliferation capacity and cytotoxicity after stimulation with H60. Clone #14, #15, and #23 were tested for the TCR binding avidity to H60-peptide/H-2K<sup>b</sup> and analyzed for TCR sequences. Results: H60-specific CTL clones showed different levels of proliferation capacity and cytotoxic activity to H60-stimulation. Clones #14, #15, and #23 showed high proliferation activity, high cytotoxicity, and low activities on both aspects, respectively, and have TCRs with different binding avidities to H60-peptide/H-2K<sup>b</sup> with  $t_{1/2}$  values of 4.87, 6.92, and 13.03 minutes, respectively. The TCR usages were  $V \alpha 12D-3-01+J \alpha 11-01$  and  $V \beta 12-1-01+D \beta 1-01$ +J2-7-01 for clone #14,  $V \alpha 13D-1-02+J \alpha 34-02$  and  $V\beta 13-1-02+D\beta 2-01+J\beta 2-7-01$  for clone #15, and  $V \alpha 16D + J \alpha 45-01$  and  $V \beta 12-1-01 + D \beta 1-01 + J \beta 2-5-01$ for clone #23. Conclusion: The results will be useful for modeling GVL and generation TCR transgenic mouse.

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#### INTRODUCTION

Allograft rejection is the major barrier for the success of solid tissue and stem cell transplantation. Major risk factor for the

rejection is mismatch at MHC locus between donor and recipient. Allograft rejection, however, still occurs even after MHC-matched transplantation for incompatibility of minor histocompatibility (H) antigens (1). Minor H antigens are peptide fragments derived from polymorphic proteins in association with MHC class I and II, able to induce CD4 and CD8 T cell responses (2). Considering the disparities at multiple minor H antigens upon the transplantation between MHC-matched but background-mismatched individuals, antigenic complexity could be enormous but, in reality, immune response focuses on a few dominant H antigens, exhibiting the phenomenon of immunodominance (3,4). In B6 anti-BALB.B system, a multiple minor H antigen-mismatched case, immune responses against a few dominant minor H antigens, H60, H28, H4, and H7, account for more than 85% of total CD8 T cells response and, among those dominant minor H antigens, H60 is dominant over all other minor H antigens, with high frequency of reactive CD8 T cells detected in blood during the B6 anti-BALB response (4).

Not only for the dominancy of specific CD8 T cell response, also for its hematopoietic cell-restricted expression, H60 has been paid attention to and considered as an ideal model antigen for study on graft-versus-leukemia (GVL) effect (5). T cells originated from hematopoietic stem cell donor recognize and react against allo-minor H antigens expressed in leukemic tumor cells residing in the recipient of the stem cells, exerting GVL effect, in addition to their capability to induce graft-versus-host disease (GVHD) through the allo-reaction against minor H antigens expressed on tissues (6-9).

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Keywords: CTL clone, Proliferation, Cytotoxicity, Avidity, TCR usage

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In this respect, CD8 T cells reactive to dominant and hematopoietic cell-restricted H60 would be exploited for tumor therapy and, for this, H60-specific CD8 T cell clones would be useful tool for modeling GVL effect in mouse.

The H60-specific response is initiated by the recognition of 8-a.a. peptide (LTFNYRNL) in the context of H-2K<sup>b</sup> by TCR expressed on CD8 T cells (5). H60 peptide/MHC complexes induce specific CD8 T cell response in CD4 help dependent manner. The help is mediated via CD40-CD40L interaction and is required not only for the induction of primary response but also for the memory cell expansion (10). The dominance of H60-specific response has been ascribed to high precursor frequency of the reactive CD8 T cells and the frequency was estimated to be one out of 10,000 ~ 20,000 cells (11). Spectratyping and sequencing of the CD8 T cells participating in H60-specific responses revealed the involvement of T cells with wide variety of TCRs (12). Based upon these results, we decided to establish CD8 T cell clones reactive to H60 and identify the TCR sequences of the clones, which would be basis for modeling of GVL and generating TCR transgenic mouse, a useful tool for detailed analysis on the mechanism of the GVL effects. In order to do this, we established and characterized H60-specific CTL clones. And the TCR usages of those were identified.

## MATERIALS AND METHODS

### Mice and immunization

C57BL/6 (B6) mouse was purchased from The Jackson laboratory (Bar Harbor, ME, USA). H60 congenic mouse (B6,C-H60/DCR) was a kind gift from Dr. Roopenian (The Jackson Laboratory). The mice were bred and kept under specific pathogen free (SPF) condition at the center for animal resource development of Seoul National University College of Medicine. To induce H60-specific CD8 T cell response in the B6 mice, splenocytes from male H60 congenic mice were injected into female B6 mice via intraperitoneal route  $(2\times10^7 \text{ cells/mouse})$ .

### Antibodies and flow cytometry

Peripheral blood lymphocytes and splenocytes from immunized mice, MLC cells, CTL lines, or CTL clones were incubated at 4°C for 30 min in staining buffer (1×PBS containing 0.1% bovine serum albumin and 0.1% sodium azide) containing PE-labeled H60-tetramer and FITC-labeled CD11a (M17/4; eBioscience, San Diego, CA) and APC-labeled CD8

(53-6.7; eBioscience) antibodies. The stained cells were analyzed using a FACSCalibur equipped with CellQuest software (BD Pharmingen, San Diego, CA).

#### Generation of CTL clones

H60-specific CTL lines were established through mixed lymphocyte culture (MLC) of splenocytes obtained from the immunized B6 mice with those from H60 congenic mouse after irradiation (2,000 cGy) and regular re-stimulation with irradiated H60 congenic splenocytes afterward in DMEM media supplemented with 5% FBS and IL-2 (50 U/ml). Limiting dilution was performed with the H60-specific CTL lines at the 5<sup>th</sup> passage to obtain the H60-specific CTL clones. Then the established clones were maintained through regular re-stimulation with irradiated H60 congenic cells.

#### Proliferation assay of H60-specific CTL clones

CTL clones  $(5\times10^4$  cells) were cultured with irradiated H60 congenic splencoytes  $(5\times10^5$  cells) in the presence of 5 U/ml and 20 U/ml of IL-2. After 48 hr, the co-culture was added with  $^3$ H and further incubated for 18 hr. Then, the cells were harvested and the incorporation of  $^3$ H was measured.

## Cell-mediated lysis assay

The standard  $^{51}$ Cr release assay was performed as described previously (4). In brief,  $^{51}$ Cr-labeled T2-K<sup>b</sup> cells were incubated with  $1\,\mu\mathrm{M}$  concentration of synthetic peptides (LTFNYRNL for H60 and RGYVYQGL for VSV) for 30 min at  $37^{\circ}$ C for peptide loading onto the target cells and then coincubated at  $37^{\circ}$ C with CTL clones (effectors) at E: T cell ratios of 2:1 and 10:1. Lysis of target cells was measured as specific lysis based on the levels of  $^{51}$ Cr released by H60-loaded target cells relative to those by VSV-loaded target cells,

## Inverse RT-PCR and identification of TCR usage of H60-specific clones

Total RNA was isolated from H60-speicific CTL clones and 2  $\mu$ g were used for 1<sup>st</sup> strand synthesis of cDNAs corresponding to TCR alpha chain using a primer 5'-CTGGTTGCTCCAGGCAATGGCCCCATT-3' (TCR  $\alpha$  R), and beta chains using a primer 5'-AGGATTGTGCCAGAAGGTAGCAGAGAC-3' (TCR  $\beta$  R). Second strand synthesis was done with *E. coli* DNA polymerase (Koschem, Daejon, Korea) and *E. Coli* RNAse H (Koschem). For circularization, the ds cDNAs were incubated with T4 DNA ligase at 16°C for overnight. The ligation mixtures were used for PCR amplification using the C  $\alpha$ -specific pri-

mers, TCRAinvF (5'-TCTGGAACGTTCATCACTGACAAA ACT-3') and TCRAinvR (5'-GGTGCT GTCCTGAGACCGAGGATC-3'), and C $\beta$ -specific primers TCRBinvF (5'-TACAAGGAGAGCAA-TTATAGCTACTGC-3') and TCRBinvR (5'-GCTTTTGATGGCT-CAAACAAGGAGACA3'). 35 cycles at an annealing temperature of 60°C were performed. Products of the inverse PCR were then cloned to pGEM-T vector and subsequently subjected to sequence analysis. The TCR usages were identified according to the IMGT database.

#### Determination of relative avidity

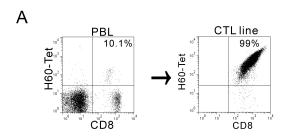
CTL clones were labeled with 50  $\mu$ g/ml of PE-labeled H60-tetramer, incubated at 4°C for 1 hr in FACs buffer and, then, washed extensively. Following this initial tetramer binding, an aliquot of  $10^5$  cells was chased in FACs buffer with incubation at  $37^{\circ}$ C for indicated times. The clones were also stained with FITC-labeled anti-TCR  $\beta$  antibody to monitor TCR levels.

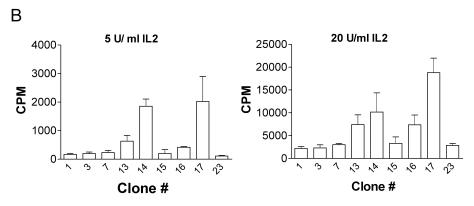
Cells were analyzed by flow cytometry. Mean fluorescence intensity (MFI) values of binding of tetramer and anti-TCR  $\beta$  antibody were plotted. The  $t_{1/2}$  of TCR-peptide/MHC interaction was calculated from specific mean fluorescence intensity using nonlinear regression analysis fitted to classical Michaelis-Menten Kinetics (Prism version 5, GraphPad, San Diego, CA).

#### RESULTS AND DISCUSSION

#### Generation of H60 specific CTL clones

Female B6 mice were immunized with male H60 congenic splenocytes and, on day 14 post-immunization, mixed leukocyte culture (MLC) was set up with irradiated H60 congenic splenocytes. Through five rounds of re-stimulation of the MLC cells with irradiated H60 congenic splenocytes, H60-specific CTL line was established (Fig. 1A). While the frequency of



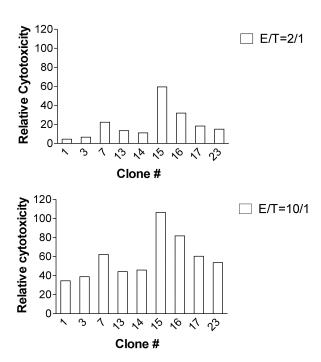


**Figure 1.** Establishment and characterization of H60-specific CTL clones. (A) Female C57Bl/6 (B6) mice were immunized with splenocytes from male H60 congenic mice, and, then, peripheral blood lymphocytes (PBLs) from the immunized mice were examined to see whether H60-specific response was induced by staining with H60 tetramer and subsequent flow cytometry on day 10 post-immunization (left). On day 14, splenocytes from the immunized mice were cultured with irradiated splenocytes from H60 congenic mouse and the mixed lymphocyte culture (MLC) was re-stimulated regularly on weekly basis to generate H60-specific CTL line (right). H60-specific CTL clones were derived from the CTL line on passage 5. (B) Proliferation assay of the H60-specific CTL clones. Established H60-specific CTL clones were examined for their proliferation capacity in reaction to the stimulation with H60 congenic cells in the presence of IL-2 at the concentrations of 5 U/ml (left) and 20 U/ml (right). Howas added to the culture on 48 hr later and, after further induction for 18 hr, cells were harvested and H-incorporation was measured. The assay was performed in triplicates and the data represent three independent experiments.

H60-specific CD8 T cells was in ranges of  $5 \sim 10\%$  of CD8 T cells in blood of the immunized B6 mice at the peak of response, 99% of the established CTL line was positive for H60-tetramer as shown by flow cytometry analysis (Fig. 1A). With this CTL line at the  $5^{th}$  passage, limiting dilution was performed to obtain H60-specific CTL clones. Then, CTL clones, each of which was originated from a single cell via the limiting dilution, were maintained through re-stimulation with irradiated H60 congenic cells on weekly basis and 24 CTL clones reactive to H60 were established.

#### CTL clones have different capacity of proliferation

Among the initial 24 CTL clones obtained, finally, nine CTL clones (#1, 3, 7, 13, 14, 15, 16, 17, and 23) were able to be maintained beyond the passage seven and subjected to clonal characterization. First, each CTL clone was tested for proliferation capacity in response to stimulation with irradiated H60 congenic cells. When <sup>3</sup>H-incorporation by the CTL clones during the stimulation was measured, clone # 14 and



**Figure 2.** Cytotoxicity of H60-specific CTL clones. CTL clones were incubated with <sup>51</sup>Cr-labeled T2-K<sup>0</sup> cells after loading with H60 or VSV peptide at the E:T ratios of 2:1 and 10:1. <sup>51</sup>Cr released into culture supernatant was measured and specific cytotoxicity was calculated in relation to the <sup>51</sup>Cr released in the VSV-peptide-included wells. The assay was performed in triplicates and the mean values were plotted. The data represent three independent experiments.

#17 proliferated relatively highly (Fig. 2), even when the concentration of supplemented IL-2 was limiting (5 U/ml). The other cells, clone #1, #3, #7, #15, and #23, showed rather minimal, but enough to be maintained for next passage, proliferation by the stimulation. These data demonstrated that even though they were all H60-specific, each clone showed a different level of proliferation capacity in reaction to the H60-stimulation.

## Different capacity of CTL activity of the H60-specific CTL clones

Next, we examined whether cytotoxic activity of the H60-specific CTL clones would be different. In order to do this, the nine CTL clones were subjected to cell mediated lysis (CML) assay by incubating with <sup>51</sup>Cr-labeled and H60-peptide loaded T-2K<sup>b</sup> target cells at two different effector/target ratios of 2:1 and 10:1. In overall, cytotoxicity of the nine CTL clones was higher at the E:T ratio of 10:1 than 2:1 and, among the clones, clone #15 was distinguished for its high CTL activity against H60-loaded T-2K<sup>b</sup> cells. The clone #14, which showed the second highest proliferation activity in reaction to stimulation with H60 congenic cells (Fig. 1B), belonged to the group with low CTL activity. This result combined with that from proliferation assay demonstrated that even though all the CTL clones were specific for H60, they possessed different levels of proliferation capacity and cytotoxicity.

# Different binding affinity of H60-specific CTL clones for peptide/MHC complexes

Then, we wondered whether CTL clones would show different levels of TCR affinity to cognate ligand, H60-peptide/ H-2K<sup>b</sup> complexes. To examine this, the affinity was empirically measured via determining  $t_{1/2}$  of H60-peptide/H-2K<sup>b</sup>-TCR interactions. We selected CTL clone #14, #15, and #23 based upon the clonal characteristics, for high proliferation capacity (#14), high cytotoxicity activity (#15), and low activities in both aspects (#23), respectively. The CTL clones were stained with saturating amount (50  $\mu$  g/ml) of H60-teramer for 1 hr and, after washing, the cells were incubated at 37°C for indicated time period and examined for falling-off the tetramers from their TCR by flow cytometry analysis, measuring the mean fluorescence intensity. To see whether TCR expression itself was influenced, the cells were also stained with anti-TCR  $\beta$ antibody before flow cytometry analysis. When the mean fluorescence intensity (MFI) values of H60-tetramer and anti-TCR  $\beta$ antibody staining were plotted, we found that initial MFI val-

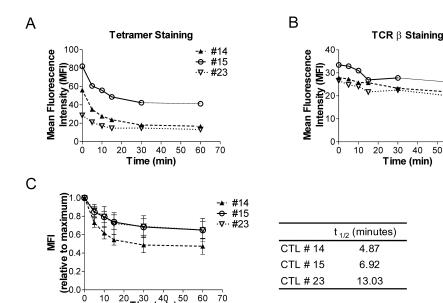


Figure 3. Biochemical features of the interaction of H60-peptide/ H-2K<sup>b</sup>-TCR of H60-specific CTL clones (A) CTL clones, #14, #15, and #23, were reacted with H60tetramer-PE, washed extensively and incubated 37°C for 1 hr. The dissociation of the tetramer from cells during the chase was monitored by flow cytometry. (B) Cells were also stained with anti-TCR  $\beta$  antibody post-chase to detect TCR levels of the cells. The MFI values from the flow cytometry (A and B) were plotted. (C) Relative MFIs were MFI values at indicated time points fractionated by MFI value detected at the beginning of the chase. And the  $t_{1/2}$  was obtained based on this dissociation curve. The data represent two independent experiments.

ues of tetramer staining were different among the three CTL clones (Fig. 3A). The initial MFI value was highest with clone #15 (82) and lowest with clone #23 (30) with #14 in the intermediate (58), even though the TCR level on the cell surfaces were not significantly different in a range of 28~33 MFI values (Fig. 3B). This demonstrated that the three CTL clones, #14, #15, and #23, bound to H60-tetramer with different avidities. When the  $t_{1/2}$  of H60-peptide/H-2K<sup>b</sup>-TCR interactions was calculated (Fig. 3C), the duration of the interaction was shortest in clone #14 (4.87 minutes) and longest in clone #23 (13.93 minutes), demonstrating that even though the TCR of clone #23 has very low initial avidity to the H60-peptide/H-2K<sup>b</sup>, the interaction is stable. Moreover, the results demonstrate that TCR of clone #14, which has high proliferation capacity, has modest initial binding avidity to H60-peptide/H-2K<sup>b</sup> but the duration of the binding is short. In the case of clone #15, the TCR interaction with H60 peptide/H-2K<sup>b</sup> initiated with high avidity and the interaction was rather stable with about 7 minutes of  $t_{1/2}$ . All these results showed that each clone has its own TCR binding avidity to the specific ligand as well as proliferation and cytotoxic capacities.

## Identification of TCR sequences of the H60-specific CTL clones

Then we thought that such clonal differences in proliferation activity, cytotoxicity, and TCR binding avidity to H60 peptide/H-2K<sup>b</sup> would reflect different TCR usages of the clones. To identify the sequences encoding the TCRs expressed by clone #14, #15, and #23, inverse PCR was performed with the RNAs extracted from each clone and the PCR products were cloned to pGEM-T vector and analyzed for sequence (Fig. 4). The sequence analysis revealed that the three CTL clones had different usages of TCR  $\alpha$  and TCR  $\beta$ . TCR complex of CTL clone #14 consists of V  $\alpha$  12D-3-01 and J  $\alpha$  11-01 for TCR  $\alpha$  chain and V  $\beta$  12-1-01, D  $\beta$  1-01, and J  $\beta$  2-7-01 for TCR  $\beta$  chain. CTL clone #15 consists of V  $\alpha$  13D-1-02 and J  $\alpha$ 34-02, and V  $\beta$  13-1-02, D  $\beta$  2-01, and J  $\beta$  2-7-01 for TCR  $\alpha$ and  $\beta$  chains, respectively. TCR  $\alpha$  and  $\beta$  chains of Clone #23 consists of V  $\alpha$  16D and J  $\alpha$  45-01, and V  $\beta$  12-1-01, D  $\beta$ 1-01, and J  $\beta$  2-5-01, respectively. Based upon the sequence analysis data, we could suggest that the different TCR usages by each clone would have been reflected to differences in the clonal characteristics.

40 50 60

From an antigenic standpoint, H60 has been found to recruit CD8 T cells with diverse TCRs into the specific response, after primary challenge as well as after memory expansion stimulation (12). The results from this study through a series of clonal characterization showed that the CD8 T cells recruited to H60-specific CD8 T cells responses would have each different capacity, in terms of proliferation, cycotoxicity, and TCR avidity to the cognate ligand H60 peptide/H-2K<sup>b</sup>. And the H60-specific CTL clones and TCR information obtained in this study would be useful for understanding the

Α

Clone	Chain	V gene	D gene	J gene
#14	alpha	12D-3-01	-	11-01
	beta	12-1-01	1-01	2-7-01
#15	alpha	13D-1-02	-	34-02
	beta	13-1-02	2-01	2-7-01
#23	alpha	16D	-	45-01
	beta	12-1-01	1-01	2-5-01

В

#14  $\mathsf{TCR}\alpha$  : —-  $\mathsf{TGTACTACTGTGCTCTGAGTGA}$   $\mathsf{GACTCGGGATACAACAAAC}$  ---

 $V\alpha 12D-3-01$   $J\alpha 11-01$ 

#14 TCR\$ : --- CTTCTGTGCCAGCTCTCTC GG CAGGGGC G ATGAACAGTACT ---

Vβ12-1-01 Dβ1-01 Jβ2-7-0<sup>-</sup>

#15 TCR $\alpha$  : \_\_\_\_ CACTTATCTCTGTGCT TTGGCGTT TTCCAATACCAACAA ---

 $V\alpha 13D-1-02$   $J\alpha 34-02$ 

#15 TCR\$ : ---- GCCAGCAGTGA TGGGGGGC ATGAACAGTACTTCGGTC ---

Vβ13-1-02 Dβ2-01 Jβ2-7-01

#23 TCRlpha : --- CTGTGCTATGAGAGAGGG CT ATACAGAAGGTGCAGATAGAC ---

 $V\alpha$ 16D  $J\alpha$ 45-01

#23 TCRB: --- GTACTTCTGTGCCAGC AA GACAGG AGTT AACCAAGACACCCAGT ---

 $V\beta 12-1-01$   $D\beta 1-01$   $J\beta 2-5-01$ 

**Figure 4.** Sequence analysis of H60-specific CTL clones. (A) Annotation of TCR usages of H60-specific CTL clones, #14, #15, and #23, and (B) sequences of the CDR3 regions were shown.

cellular mechanism of graft rejections and GVHD, and also for modeling the GVL effect under MHC-matched transplantation condition.

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#### CONFLICTS OF INTEREST

The authors have no financial conflict of interest.

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