

T Cell Receptor (TCR) Antagonism without a Negative Signal: Evidence from T Cell Hybridomas Expressing Two Independent TCRs

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Summary

Antagonist peptides inhibit T cell responses by an unknown mechanism. By coexpressing two independent T cell receptors (TCRs) on a single T cell hybridoma, we addressed the question of whether antagonist ligands induce a dominant-negative signal that inhibits the function of a second, independent TCR. The two receptors, V α 2V β 5 and V α 2V β 10, restricted by H-2K^b and specific for the octameric peptides SIINFEKL and SSIEFARL, respectively, were coexpressed on the same cell. Agonist stimulation demonstrated that the two receptors behaved independently with regard to antigen-induced TCR downregulation and intracellular biochemical signaling. The exposure of one TCR (V α 2V β 5) to antagonist peptides could not inhibit a second independent TCR (V α 2V β 10) from responding to its antigen. Thus, our data clearly demonstrate that these antagonist ligands do not generate a dominant-negative signal which affects the responsiveness of the entire cell. In addition, a kinetic analysis showed that even 12 h after engagement with their cognate antigen and 10 h after reaching a steady-state of TCR internalization, T cells were fully inhibited by the addition of antagonist peptides. The window of susceptibility to antagonist ligands correlated exactly with the time required for the responding T cells to commit to interleukin 2 production. The data support a model where antagonist ligands can competitively inhibit antigenic peptides from productively engaging the TCR. This competitive inhibition is effective during the entire commitment period, where sustained TCR engagement is essential for full T cell activation.

Key words: T cell receptor antagonist • T cell activation • T cell commitment

T cell activation depends on the specific recognition of peptides presented in the binding groove of MHC molecules (1, 2). The binding of the peptide/MHC ligand to the TCR leads to the tyrosine phosphorylation of TCR-associated proteins, recruitment of kinases and adapters, and activation of intracellular signaling pathways (3–6). Interestingly, T cells can differentially respond to subtle changes in either the MHC or peptide ligand. Substitutions in TCR contact residues can give rise to antagonist ligands that do not elicit any measurable T cell effector functions, but are able to diminish or even abrogate the response to the nominal antigen when both the agonist and antagonist are simultaneously displayed on APCs (7–11).

Various models have been proposed to explain the mechanistic basis for the differential response to antagonist ligands by the same TCR and its translation into altered signaling and T cell activation (12–16). First, a quantitative model has been put forward which proposes that TCR antagonism is due to the fact that antagonist ligands interact

nonproductively with the TCR, and competitively inhibit antigenic ligands from productively engaging the TCR. This competition might disturb the formation of necessary signaling oligomers (7, 12, 17–19). This model is supported by studies showing that several antagonist ligands fail to induce TCR-dependent signaling events such as the generation of sustained Ca²⁺ flux and turnover of the inositol phosphates (7, 9, 17, 20–22) as well as exhibit lower affinities and faster off-rate kinetics for the TCR than antigenic ligands (23–25).

In contrast, various qualitative models have been proposed where antagonist ligands induce a TCR-mediated differential (8, 9, 16, 21, 26–28) or even negative signal (13, 14, 29, 30). One model attributes the induction of a negative signal to an inappropriate conformational change induced by antagonist ligands (14, 31). In contrast, another model proposes that quantitative differences in TCR/ligand binding translate into a negative intracellular signal (29). Studies of positive selection of immature thymocytes

by antagonist ligands in fetal thymic organ culture have shown that these ligands are in fact capable of delivering a signal through the TCR (32–34). Moreover, several groups have demonstrated that altered peptide ligands, with partial agonist or with even pure antagonist properties, were able to initiate some but not all of the early biochemical events associated with TCR signaling (26–28). Although these experiments provided evidence for the induction of differential signaling events induced by partial agonist and antagonist ligands, they were not able to distinguish whether the observed inhibitory effect was due to an induced negative signal or to incomplete activation based on the faster TCR dissociation from the antagonist/MHC complex (24, 25).

In the present study, we have addressed the mechanism of antagonism induced by antagonist ligands using T cell hybridomas coexpressing two independent TCRs. If antagonist ligands induced signals with negative regulatory characteristics, then stimulation through one receptor should be inhibitable by antagonizing the other. Our data clearly demonstrate that the antagonist ligands we examined do not generate such an intracellular dominant-negative signal which inhibits the stimulation of a second independent TCR. The two TCRs, expressed on the same cell, behaved independently with regard to antigen-induced TCR internalization as well as intracellular biochemical signaling. Interestingly, the T cell hybridoma could be inhibited by antagonist peptides far beyond the time point of maximal antigen-induced TCR internalization. The fact that the window of sensitivity to antagonist peptides correlated exactly with the time required for the cells to commit to activation supports a competitive model to explain TCR antagonism.

Materials and Methods

DNA Constructs. The OVA-TCR-1 (OT-I) is comprised of rearranged TCR α (V α 2-J α 26) and TCR β (V β 5-D β 2-J β 2.6) chains and is derived from the K^b-restricted, OVA₂₅₇₋₂₆₄-specific CTL clone, 149.42 (35). The cDNA encoding the TCR α chain was cloned into the G418 resistant retroviral vector, LXSN (36–38). Similarly, the OT-I TCR β chain was cloned into the puromycin resistant retroviral vector, LXSP (39). The TCR β chain (V β 10-D β 2-J β 2.5), when paired with the OT-I V α 2 chain, confers specificity for the Herpes simplex viral peptide, SSIEFARL (40). The V β 10 cDNA was cloned into LXSP as well. The cDNAs encoding CD8 α and CD8 β were kindly provided by P. Cosson (University of Geneva, Geneva, Switzerland). The CD8 α cDNA was cloned into the retroviral vector LXSHD, which expresses a histidinol resistance gene, while the CD8 β cDNA was similarly cloned into the retroviral vector LXSH, containing a hygromycin resistance gene. LXSHD and LXSH were provided by A. Kazlauskas (Harvard Medical School, Boston, MA).

Cells. The TCR- α^{-}/β^{-} T hybridoma, 58, has been described previously (41). The 58CD8 α/β cells were generated by sequentially infecting the original cell line 58 with supernatants containing viruses encoding CD8 α and CD8 β . Subsequently, the TCR α and β chains were transfected by retroviral infection of 58CD8 α/β to obtain cell lines expressing a single TCR (V α 2V β 5 or V α 2V β 10). Similarly, retroviral infection and cell sorting was used to generate a cell line coexpressing two TCRs

(V α 2V β 5 and V α 2V β 10). The P1.32K^b cell line, directed from the cell line P815 by transfection of a DNA construct encoding the MHC class I molecule K^b (42), was used for peptide presentation. All cells were grown in IMDM supplemented with 1.5% heat inactivated FCS, 2 mM l-glutamine, 100 U/ml penicillin, 100 μ g/ml streptomycin, and 50 μ M β -mercaptoethanol. The indicator cell line HT-2 (43) was grown in IMDM containing 5% FCS with the addition of 250 U/ml recombinant IL-2. The ecotropic packaging cell line Bosc23 was purchased from American Type Culture Collection and grown in IMDM containing 10% FCS.

Transfection and Infection. Bosc23 packaging cell line was transfected as described previously (44). The supernatant containing retroviral particles was used to infect the 58 T cell hybridoma. Briefly, 5×10^5 58 cells were resuspended in 100 μ l IMDM, and 1 ml retroviral supernatant containing 40 μ g DEAE-dextran was added. After 6–12 h, 5 ml of fresh IMDM and the appropriate selective drug was added (1 mg/ml G418 [GIBCO BRL], 3 μ g/ml puromycin [Sigma Chemie], 2 mM histidinol [Sigma Chemie], or 0.5 mg/ml hygromycin [Calbiochem]). The surviving cells were analyzed after 5–7 d and sorted for surface expression by FACS[®]. In all experiments at least two independently generated T cell hybridoma lines were compared, and similar results were obtained. Transfected cells were always maintained in medium containing the selective drugs.

Peptides. The peptides were synthesized at the Basel Institute for Immunology using FastMoc[™] chemistry on 430A peptide synthesizer (Applied Biosystems). The amino acid sequences were the following: SIINFEKL (V α 2V β 5-specific antigen), EIINFEKL (V α 2V β 5-specific antagonist E1), SIINFEPL (V α 2V β 5-specific antagonist P7), SIKFEKL (the control peptide K4), and SSIEFARL (V α 2V β 10-specific antigen).

Antibodies. The anti-V β 5 mAb, MR9-4 (44), anti-CD3 ϵ mAb, 2C11 (45), and anti- ζ mAb, H146-968 (46), were purified from culture supernatants using protein G (Pharmacia). The anti-V α 2.1-specific mAb, B20.1 (47), the anti-V β 10 mAb, B21.5 (48), and the anti-K^b mAb, AF6-88.5 (49), were purchased from Pharmingen. The anti-phosphotyrosine mAb, 4G10, was purchased from Upstate Biotechnology. To detect bound anti- ζ antibodies in Western blots, we used goat anti-rabbit antibodies labeled with horseradish peroxidase (HRPO)¹ from Southern Biotechnology Associates. The blocking anti-K^b mAb, provided by J. Bluestone, was purified from culture supernatants using protein A (50).

Quantitation of TCR Surface Expression. To calculate the relative amount of the two TCRs (V α 2V β 5 and V α 2V β 10) coexpressed on the same hybridoma cell, the expression of each TCR β chain (measured by staining with the appropriate anti-V β mAb) was normalized to the total amount of TCR expressed on the surface (measured by staining with an anti-V α 2 mAb). The ratio of V β 5/V α 2 or V β 10/V α 2 staining on cells expressing a single TCR was taken as 100%. On the surface of the hybridomas expressing two TCRs, V α 2V β 5 and V α 2V β 10 heterodimers accounted for 60 and 40% of the surface TCRs, respectively.

Stimulation Assays. 90 μ l containing 5×10^4 P1.32K^b cells was plated in flat-bottomed 96-well plates and incubated with 10 μ l peptide for 4 h at 37°C. 8×10^4 T hybridoma cells in 100 μ l medium were subsequently added. After a further 25 h of incubation at 37°C, the supernatant was harvested and assayed for IL-2.

Antagonism Assays. P1.32K^b cells were first loaded for 4 h at 37°C with the indicated amount of agonist peptide and unbound

¹Abbreviation used in this paper: HRPO, horseradish peroxidase.

peptide was removed by washing. Peptide loaded cells ($5 \times 10^4/90 \mu\text{l}$) were plated in flat-bottom 96-well plates. $10 \mu\text{l}$ of antagonist peptides, or $10 \mu\text{l}$ of control peptide or medium alone was added and the cultures were incubated at 37°C for 1 h. 8×10^6 T hybridoma cells in $100 \mu\text{l}$ were then added and incubated for 25–27 h at 37°C . The supernatant was harvested and analyzed for the presence of IL-2.

IL-2 Assay. IL-2 content was determined by incubating 2×10^3 HT-2 cells per well in round-bottom 96-well plates with serial dilutions of culture supernatant for 24 h. Alamar blue substrate (Alamar Biosciences) was then added and IL-2 titer was determined by comparison to a standard curve generated using recombinant murine IL-2 (PharMingen) using SOFTmaxPro version 1.1 software.

FACS[®] Analysis of TCR Downregulation. The stimulation of the T cell hybridomas was carried out in parallel and under the same conditions as the antagonism assays mentioned above. To ensure conjugate formation, cells were centrifuged briefly. Similar results were obtained with round- or flat-bottom 96-well plates. After 3 h of stimulation at 37°C , cells were washed once in PBS containing 1% FCS and 0.05% azide, and stained with saturating amounts of PE-conjugated anti-V α 2.1 mAb (B20.1) and, in the case of the double TCR expressing cells, with biotinylated anti-V β 5 (MR9-4) or anti-V β 10 (B21.5) mAbs. For the anti-V β 10 mAb, streptavidin-APC (PharMingen) was used as a second step labeling reagent. The P1.32K^b cells were excluded by staining with anti-K^b FITC labeled mAb (AF6-88.6). Dead cells were excluded by staining with $0.5 \mu\text{g/ml}$ propidium iodide (Molecular Probes). Results were analyzed using CellQuest version 3.1 software.

Tyrosine Phosphorylation Assays. For antigen presentation, the cell line P1.32K^b was loaded either with medium alone or with antigenic peptides, $10 \mu\text{M}$ SSIEFARL or $1 \mu\text{M}$ SIINFEKL, in a 24-well plate overnight (7.5×10^5 cells/well in 1 ml 1.5% FCS IMDM). 10^7 responding T cell hybridomas were then added to each well and incubated at 37°C for 30 min. Cells were harvested and cell pellets were lysed for 20 min in 1 ml of 1% Triton X-100, 10 mM Tris-NCl, pH 7.4, and 150 mM NaCl. To prevent degradation, the lysis buffer was supplemented with 10 mM NaF, 1 mM PMSF, $10 \mu\text{g/ml}$ aprotinin, $10 \mu\text{g/ml}$ leupeptin, 1 mM orthovanadate, and $100 \mu\text{M}$ pervanadate (made from a fresh stock solution of 10 nM orthovanadate and 15 mM H_2O_2). Cell lysates were immunoprecipitated with anti-V β 5, anti-V β 10 mAbs, or anti-CD3 ϵ , respectively, and resolved by SDS-PAGE as described previously (51). Proteins were transferred onto nitrocellulose and the membrane was blocked with 1% BSA/TBS 0.1% NP-40 (BSA stock solution; Pierce) for 20 min at room temperature. Blots were probed with $1 \mu\text{g/ml}$ of the biotinylated antiphosphotyrosine mAb, 4G10 (Upstate Biotechnology) for 1 h at room temperature in 1% BSA/TBS 0.1% NP-40. Bound 4G10 antibodies were visualized with streptavidin-HRPO (1:50,000; Southern Biotechnology Associates) and an enhanced chemiluminescence system (Pierce).

Anti- ζ Blot. After phosphotyrosine detection, HRPO was inactivated by incubating the nitrocellulose membranes with 3% $\text{H}_2\text{O}_2/\text{H}_2\text{O}$ for 15 min at room temperature. Subsequently non-specific binding was blocked with 5% milk/PBS 0.1% NP-40 (nonfat dry milk; BioRad) for 20 min at room temperature. Blots were probed with 100 ng/ml biotinylated anti- ζ mAb H146-968 for 1 h at room temperature. Bound anti- ζ antibodies were visualized with goat anti-hamster HRPO (1:50,000; Southern Biotechnology Associates) and an enhanced chemiluminescence system (Pierce).

Results

Antagonism in V α 2V β 5 (OT-I) TCR Expressing T Cell Hybridomas. To investigate the mechanism of TCR antagonism, we took advantage of the T cell hybridoma line, 58CD α/β TCR- α^-/β^- , which expresses the coreceptor CD8 but no endogenous TCR (41; see Materials and Methods). Using retroviral vectors, we transduced the cDNAs encoding the V α 2V β 5 TCR, which is known as OT-I. This TCR specifically recognizes the ovalbumin peptide, SIINFEKL, bound to the MHC class I molecule K^b (9). Two antagonist peptides, E1 and P7, differing by only one residue from the antigenic peptide, have been described for this TCR (32). Optimal activation of the V α 2V β 5 expressing hybridomas, measured by the secretion of IL-2 or IL-3 and TCR internalization, required high expression of both CD8 α and CD8 β , (52; and data not shown).

The agonist peptide was prepulsed onto the APC to ensure antigen binding as previously described for conventional antagonism assays (9). Antagonism was observed with similar characteristics described for T cell clones (Fig. 1). The V α 2V β 5 expressing cells responded to the antigenic peptide SIINFEKL by secreting IL-2 and IL-3 in an antigen dose-dependent fashion and the presence of the antagonist peptides, E1 and P7, inhibited IL-2 and IL-3 production at suboptimal SIINFEKL concentrations (Fig. 1 A [IL-3 data not shown]). This inhibition of the response was specific for the antagonist peptides since a control MHC K^b-binding peptide, K4, neither inhibited nor potentiated

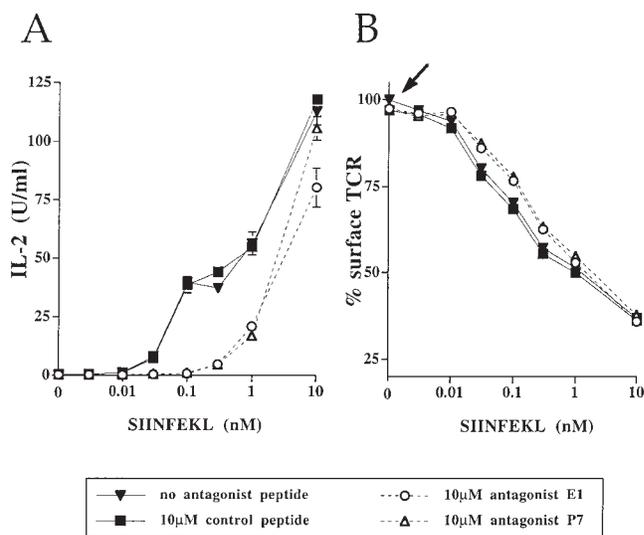


Figure 1. TCR antagonism in 58CD8 α/β T cell hybridomas expressing the V α 2V β 5 (OT-I) TCR. T cell hybridomas were stimulated with their antigen, SIINFEKL, and the effect of the antagonist (E1, P7) and control (K4) peptides on IL-2 response (A) and TCR downregulation (B) was measured. The arrow highlights the data showing that, in the absence of SIINFEKL, neither the antagonists E1 and P7, nor the control peptide K4 induced TCR internalization. The IL-2 response (A) was determined after 25 h of stimulation, while the TCR downregulation (B) was analyzed after 3 h. Data are expressed as mean of triplicates in A or of duplicates in B. This experiment is representative of eight separate experiments.

the IL-2 response elicited by the peptide SIINFEKL alone (Fig. 1 A).

Functionally triggered TCRs are internalized shortly after stimulation. Thus, downregulation of TCRs after exposure to the antigenic peptide, SIINFEKL, was used as a measure for productive TCR engagement (53, 54). As shown in Fig. 1 B, up to 75% of V α 2V β 5 TCRs were downregulated in a dose-dependent fashion after 3 h of antigen exposure. Neither antagonist peptides alone nor the control peptide K4 affected TCR expression (Fig. 1 B, see arrow). However, both antagonist peptides, E1 and P7, modestly inhibited (by 10%) TCR downregulation induced by the antigen, SIINFEKL. Although the reversal of TCR internalization correlated with the inhibition of IL-2 secretion (Fig. 1, A and B), the effect on IL-2 secretion was more pronounced. Surprisingly, the number of internalized TCRs did not necessarily correlate with the amount of secreted IL-2. This was particularly evident when the response to 1 nM SIINFEKL was antagonized by 10 μ M E1 or P7 (compare A and B in Fig. 1 and in Fig. 5). Under these conditions, TCR downregulation was not inhibited, although a significant decrease in IL-2 secretion was observed. These results suggest that in this system there is no strict correlation between the inhibition of antigen-induced TCR internalization and the effect on the IL-2 response.

Expression of Two TCRs Which Share a Common TCR α Chain on the Same Cell. To test the hypothesis that antagonist ligands induce intracellular, dominant-negative signals, we generated a T cell hybridoma line expressing two TCRs, which could be independently activated. This experimental design was used to determine whether positive signals derived from an antigen-stimulated TCR could be blocked by putative inhibitory signals generated by a second, antagonized TCR (Fig. 2). One receptor is the V α 2V β 5 TCR, which is specific for the antigenic peptide, SIINFEKL, and which can be antagonized by the altered peptides, E1 and P7 (32; Fig. 1). The V α 2 chain is also present in the second receptor but is paired instead with a V β 10 chain. This second TCR is specific for a Herpes simplex viral peptide, SSIEFARL, also presented by K b (40).

Since the two receptors differ only in their TCR β chains, no hybrid receptors of unknown specificities can be formed and the two receptors can easily be followed with V β -specific antibodies. Therefore, all observed effects can be clearly attributed to the respective receptor.

Both Agonists, SIINFEKL and SSIEFARL, and the Antagonists, E1 and P7, Are TCR Specific. First, we determined whether the two receptors, which have a common TCR α chain, are specific for their respective antigenic peptides. T cell hybridomas expressing only one type of TCR, either V α 2V β 5 or V α 2V β 10, were analyzed for their IL-2 response in stimulation assays. As shown in Fig. 3 A, V α 2V β 5 expressing cells responded only to the antigen, SIINFEKL, while V α 2V β 10 expressing cells were stimulated only by the peptide SSIEFARL (Fig. 3 B). Neither of these TCRs was stimulated by the antagonist peptides, E1 and P7, or by the control peptide, K4 (Fig. 3, A and B). Moreover,

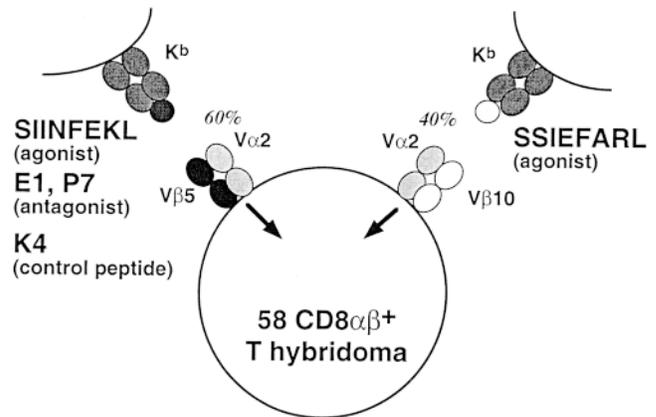


Figure 2. Coexpression of two peptide-specific TCRs on the surface of the same cell. One receptor is the V α 2V β 5 TCR which is specific for the ovalbumin peptide, SIINFEKL, presented by K b . This TCR can be antagonized by the altered peptide ligands, E1 or P7. The K4 peptide represents a control for peptide specificity since it has the similar binding affinity to K b but is not recognized by the V α 2V β 5 TCR. The second receptor is the V α 2V β 10 TCR and is specific for the Herpes simplex viral peptide, SSIEFARL, bound to K b . Both receptors were coexpressed on the T cell hybridoma line 58CD8 α / β .

TCR internalization completely correlated with the IL-2 responsiveness (data not shown).

A reduced IL-2 response to SSIEFARL compared to SIINFEKL (Fig. 3) was observed. However, this difference could not be attributed either to different TCR expression levels or to differences in the ability of these antigens to bind K b . The binding affinities of the peptides to K b , indirectly measured by their abilities to stabilize K b expression on RMA-S cells, were equivalent (data not shown). The lower IL-2 response, induced by the SSIEFARL peptide, was likely due to weak interactions of the V α 2V β 10 TCR with this peptide/MHC complex. Nevertheless, the IL-2 production in response to SSIEFARL demonstrated that the V α 2V β 10 TCR was functional.

To determine if the V α 2V β 5-specific antagonists, E1 and P7, could antagonize the V α 2V β 10 TCR directly, cells expressing only the V α 2V β 10 receptor were stimulated with the cognate antigen, SSIEFARL, in the presence of the V α 2V β 5-specific antagonist peptides, E1 and P7, or the control peptide, K4. None of the peptides inhibited the IL-2 response to SSIEFARL (Fig. 4 A) or the downregulation of the V α 2V β 10 receptor (Fig. 4 B). These results showed that the antagonists of the V α 2V β 5 TCR did not cross-react with the V α 2V β 10 receptor.

Antagonists of SIINFEKL Act Locally and in Cis. Having confirmed the specificity of the two TCRs expressed on separate hybridomas, both receptors were expressed on the same cell. Two-color FACS[®] analysis with anti-V β antibodies was used to calculate that V α 2V β 5 and V α 2V β 10 comprised 60 and 40% of the surface TCRs, respectively (data not shown). Stimulation of the double TCR expressing cells with either the V α 2V β 5- or V α 2V β 10-specific antigen resulted in a low but reproducible IL-2 response (Fig. 5, A and C). Thus, both TCRs were functional when

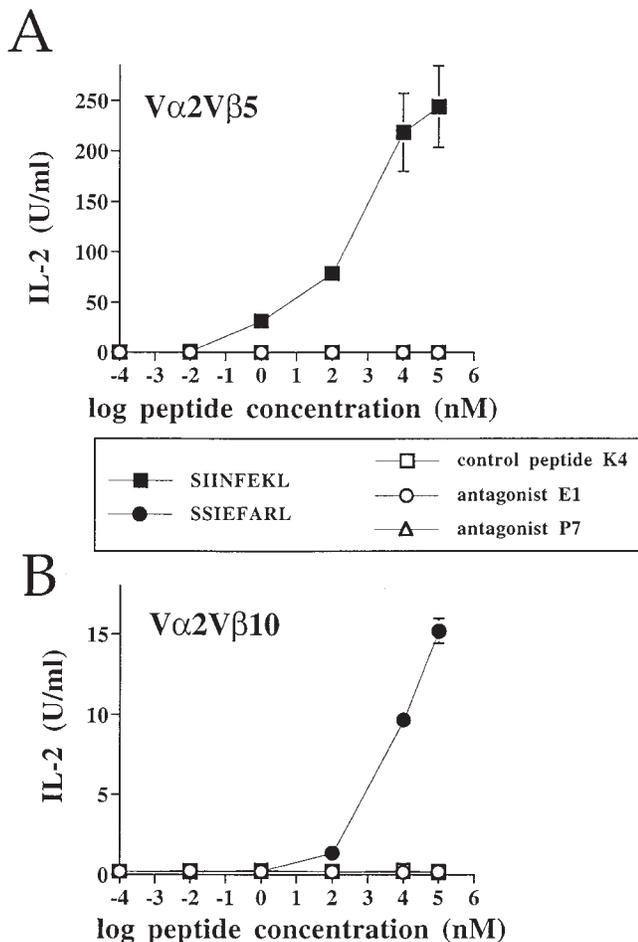


Figure 3. The two TCRs are distinctly specific for their cognate antigenic peptides. The TCRs, V α 2V β 5 (A) and V α 2V β 10 (B), were expressed separately on the cell line 58CD8 α / β and tested for their responsiveness to various peptide analogues in stimulation assays by measuring the IL-2 production. After 25 h of stimulation, supernatants were harvested and analyzed for IL-2. This experiment is representative of two independent experiments.

expressed at the surface of the same cell. An antagonist assay was performed by stimulating the dual expressing hybridoma cells with the V α 2V β 5 antigen SIINFEKL, in the presence of the V α 2V β 5-specific antagonists E1 or P7. Both antagonist peptides, but not the control peptide K4, inhibited IL-2 expression in the dual TCR expressing cells (Fig. 5 A). Additionally, the antigen-induced TCR internalization of V α 2V β 5 receptor was slightly inhibited in the presence of the antagonists E1 and P7 similar to what was observed previously (Fig. 5 B and Fig. 1 B).

To test the hypothesis that antagonist ligands deliver a negative, trans-acting signal, the double receptor expressing cells were stimulated with the V α 2V β 10-specific antigen, SSIEFARL, in the presence of the V α 2V β 5-specific antagonist peptides E1 or P7. The IL-2 response to SSIEFARL, mediated by the V α 2V β 10 TCR was unaffected in the presence of the V α 2V β 5-specific antagonist peptides E1 or P7 (Fig. 5 C). Furthermore, the V α 2V β 5-specific antago-

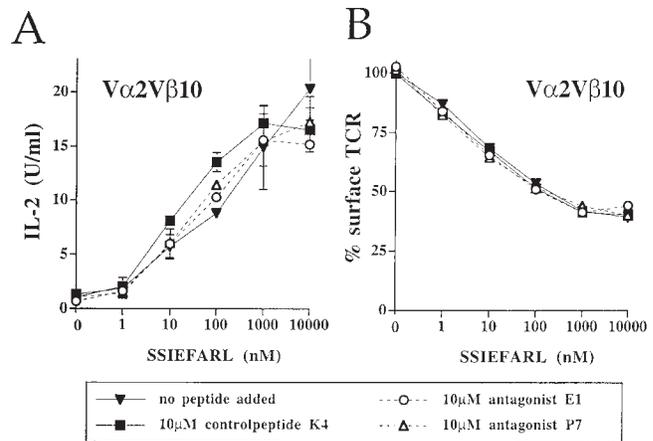


Figure 4. The V α 2V β 5-specific antagonist peptides E1 and P7 do not antagonize the response of the V α 2V β 10 TCR to the Herpes simplex viral peptide, SSIEFARL. V α 2V β 10 58CD8 α / β hybridomas were stimulated with various concentrations of SSIEFARL in the presence of the V α 2V β 5-specific antagonists, E1 and P7, or control peptide, K4. The IL-2 response (A) and antigen-induced TCR internalization (B) were measured in the same experiment and are representative of two independent experiments.

nists did not reverse the downregulation of the V α 2V β 10 TCR (Fig. 5 D).

The results clearly show that antagonizing one TCR did not inhibit the stimulation through a second, independent TCR. Apparently, these antagonist ligands do not induce a dominant-negative signal that acts in trans.

Antigen-engaged TCRs Function Independently in Their Proximal Signaling Pathways. Since the antagonized V α 2V β 5 TCR did not interfere with the response of the stimulated V α 2V β 10 TCR, we investigated the functional independence of the two receptors. To find out if both receptors function independently, both TCRs were monitored for antigen-induced TCR internalization. The stimulation with the V α 2V β 5-specific antigen, SIINFEKL, resulted in downregulation of the V α 2V β 5 TCR but not of the V α 2V β 10 TCR (Fig. 6 A). Similarly, in response to the V α 2V β 10-specific antigen, SSIEFARL, the V α 2V β 10 receptor was downregulated, while no internalization of the V α 2V β 5 receptor was detectable (Fig. 6 B). Similar results were obtained when various cell populations expressing different ratios of the two TCRs were used (not shown), supporting the observations of Valitutti et al. examining T cell clones with disparate expression of the two TCRs (53).

To address whether two different TCRs expressed on the surface of the same cell also behave independently in regard to biochemical signaling, T cell hybridomas expressing the two TCRs were incubated with unpulsed APCs or APCs pulsed with either the V α 2V β 10-specific antigen, SSIEFARL, or the V α 2V β 5-specific antigen, SIINFEKL. From cell lysates, both TCRs were either immunoprecipitated with an anti-CD3 ϵ mAb, or independently retrieved using anti-V β 5- or anti-V β 10-specific mAbs. Immunoprecipitates were resolved by SDS-PAGE and blotted for the presence of phosphorylated tyrosine residues. Only low back-

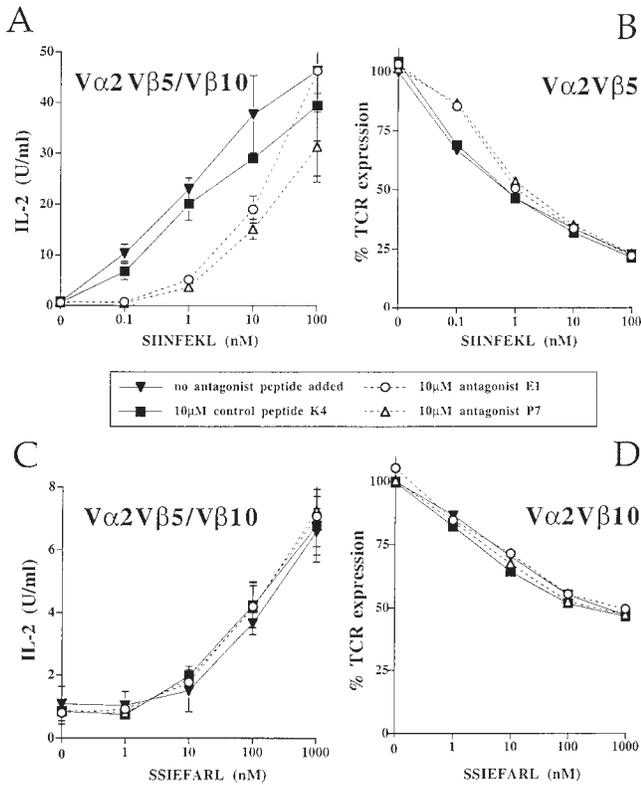


Figure 5. Engagement of one TCR with antagonist peptides does not inhibit the activation through a second independent TCR. Analyzing cells simultaneously expressing both $V\alpha 2V\beta 5$ and $V\alpha 2V\beta 10$ TCRs on the cell surface, the $V\alpha 2V\beta 5$ -specific antagonist peptides, E1 and P7, inhibited the IL-2 production (A) and the TCR downregulation (B) induced by the $V\alpha 2V\beta 5$ -specific antigenic peptide, SIINFEKL. However, the $V\alpha 2V\beta 5$ -specific antagonists, E1 and P7, did not inhibit the IL-2 response (C), nor the TCR downregulation (D) induced by the $V\alpha 2V\beta 10$ -specific antigen, SSIEFARL. The IL-2 response (A and C) and TCR downregulation (B and D) were assayed in the same experiment. The data are representative of six separate experiments.

ground phosphorylation of TCR-associated proteins was observed in T cell hybridomas exposed to unpulsed APCs (Fig. 6 C, lanes 1–3). In contrast, incubation with APCs pulsed with 10 μ M SSIEFARL induced a strong phosphorylation of various proteins which coimmunoprecipitated with the TCR. In anti-CD3 ϵ and anti-V β 10 immunoprecipitates, tyrosine-phosphorylated isoforms of ζ (p21 and p23), CD3 ϵ , ZAP-70, and a 36-kD phosphoprotein, most probably LAT, were observed (Fig. 6 C, lanes 4 and 6). To test for the phosphorylation of unstimulated bystander TCRs, the $V\alpha 2V\beta 5$ complex was immunoprecipitated from lysates of cells which had been stimulated with the $V\alpha 2V\beta 10$ -specific antigen, SSIEFARL. There was no detectable induction of phosphorylation of any protein species precipitated with the $V\alpha 2V\beta 5$ TCR complex (Fig. 6 C, lane 5). Similar results were obtained with the reverse experiment. In stimulations with 1 μ M of the $V\alpha 2V\beta 5$ -specific antigen, SIINFEKL, only anti-CD3 ϵ and anti-V β 5 immunoprecipitates showed the induction of phospho- ζ (p21 and p23), CD3 ϵ , ZAP-70, and the 36-kD phosphoprotein

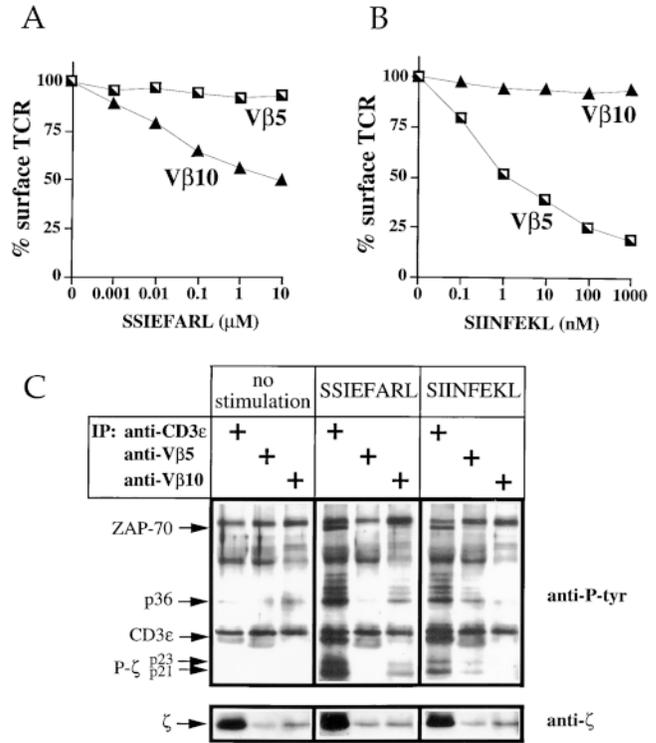


Figure 6. Antigen-engaged TCRs behave independently with regard to antigen-induced TCR internalization and biochemical signaling. The double TCR expressing hybridomas were stimulated with the $V\alpha 2V\beta 5$ -specific antigen, SIINFEKL (A), and the $V\alpha 2V\beta 10$ -specific antigen, SSIEFARL (B). TCR downregulation of both receptors was analyzed. The data are representative of three independent assays. To study the induction of TCR-associated phosphoproteins (C), P1.32K^b APC were pulsed overnight with medium alone (no stimulation) or with the antigenic peptides, 10 μ M SSIEFARL or 1 μ M SIINFEKL. 10^7 double TCR expressing cells, $V\alpha 2V\beta 5/V\beta 10$ 58CD8 α/β , were stimulated for 45 min at 37°C. Proteins in cell lysates were immunoprecipitated with anti-V β 5 mAb (MR9-4), anti-V β 10 mAb (B21.6), or anti-CD3 ϵ mAb (2C11), respectively, separated on a 12% polyacrylamide gel, and transferred to nitrocellulose. The blot was probed with the anti-phosphotyrosine mAb, 4G10, and then with the anti- ζ mAb, H146-968. The arrows on the left identify the major tyrosine-phosphorylated species.

(Fig. 6 C, lanes 7 and 8). Immunoprecipitation of the bystander TCR, $V\alpha 2V\beta 10$, did not coprecipitate any of the same phosphorylated proteins (Fig. 6 C, lane 9). As the subsequent decoration of the same blot with anti- ζ showed, both anti-V β -specific antibodies immunoprecipitated similar amounts of ζ . The anti-CD3 ϵ mAb was sixfold more efficient in coprecipitating ζ than both V β -specific mAb (Fig. 6 C, bottom).

To determine the sensitivity of detection, the eluate of cells stimulated with 10 μ M antigenic peptide was serially diluted (twofold) to determine the detection limits of the phosphoproteins (data not shown). These experiments showed that as little as 5% of the phosphorylated ζ (shown in Fig. 6 C, lane 4) could be detected. Thus, the biochemical activation of bystander TCRs is very limited, if it occurs at all. The combined results demonstrate that the two TCRs expressed on the same cell are biochemically and functionally independent.

The Antagonists E1 and P7 Do Not Alter the Kinetics of TCR Internalization. Since antagonist ligands do not induce a dominant-negative signal, antagonist peptides may outcompete agonist peptides for interaction with the TCR. As shown in Fig. 1 B antagonist ligands only slightly inhibited the antigen-induced TCR downregulation. In fact, the extent of TCR internalization in the presence of antagonist peptides did not reflect the profound inhibition of IL-2 secretion observed (Fig. 1). To assess whether antagonist peptides perturb T cell stimulation by affecting the kinetics of agonist-induced TCR downregulation, we monitored TCR internalization over time (Fig. 7). Antagonist peptides affected the extent but not the kinetics of antigen-induced TCR internalization. The full extent of TCR downregulation was reached after 2–3 h of sustained antigen exposure.

Susceptibility to Antagonism Parallels the Time of Commitment to T Cell Activation. To better understand the events that led to antagonism, the time period when cells are sensitive to antagonist ligands was determined. T cell hybridomas, expressing the single TCR, V α 2V β 5, were stimulated with the antigenic peptide, SIINFEKL, and antagonist peptides were added at various time points thereafter. The addition of antagonist peptides could fully inhibit the IL-2 response up to 12 h after addition of the antigen, SIINFEKL (Fig. 8). This effect was peptide-specific as the control peptide, K4, did not abrogate the IL-2 response.

To determine whether the time period during which T cells are susceptible to antagonist ligands overlapped with the time period required for T cell commitment, the K^b-specific antibody, Y3, was used to specifically block the in-

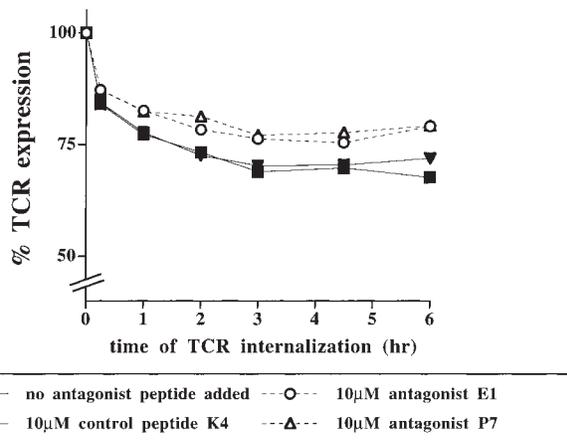


Figure 7. The kinetics of antigen-induced TCR internalization are unaltered in the presence of antagonist ligands. P1.32K^b APCs were pulsed for 4 h at 37°C with 100 pM antigenic peptide SIINFEKL. Unbound peptides were removed and the cells were incubated for 1 h with 10 μM antagonist peptides E1 or P7, or 10 μM control peptide K4, or medium alone (no peptide added). The responders, hybridomas expressing the V α 2V β 5 TCR, were added at various time points. T cell-APC conjugates were formed by centrifugation and all samples were harvested and stained at the same time. TCR surface expression was detected by PE-labeled anti-V α 2 (B20.1) and expressed as percentage of mean channel fluorescence of the unstimulated cells. The data are representative of four experiments.

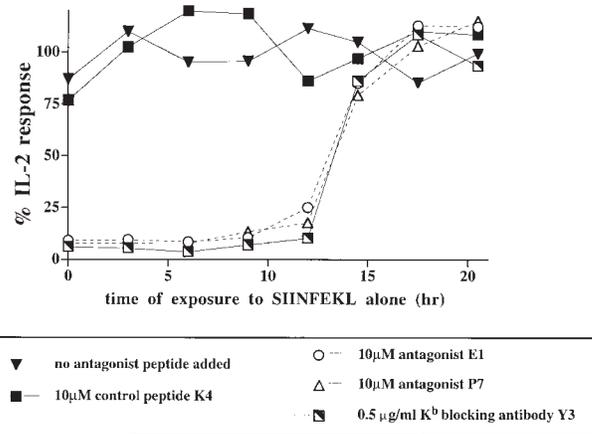


Figure 8. Susceptibility to antagonist ligands corresponds to the time of commitment. V α 2V β 5 58CD8 α / β cells were stimulated with P1.32K^b cells which had been prepulsed with 100 pM SIINFEKL peptide for 4 h at 37°C. At various time points after the initiation of the stimulation, antagonist peptides E1 or P7, control peptide K4, anti-K^b blocking antibody Y3, or medium alone were added, and the cells were resuspended. After 25 h of culture, the supernatant was harvested and assayed for IL-2 production. For more convenience, the kinetic analysis was carried out in two parts separated by 12 h. The mean of secreted IL-2 obtained in the samples where no antagonist peptides were added was set as 100%. The data are representative of four independent experiments.

teraction of the V α 2V β 5 TCR with the SIINFEKL/K^b complex. The time required for sustained TCR engagement, determined using the blocking antibody, Y3, correlated exactly with the period of susceptibility to antagonist peptides (Fig. 8). The cells were irreversibly committed to IL-2 secretion only after 12 h of continuous exposure to the antigen ligand, in the absence of antagonist peptides or blocking mAbs.

Thus, antagonist ligands were able to inhibit cellular responses long after the responding T cell had interacted with its cognate antigen and achieved a reduced steady-state level of TCR expression. The T cell was fully sensitive to the inhibitory effects of antagonist ligands until it did not require further engagement with the antigen/MHC complex and was irreversibly committed to respond.

Discussion

T cell antagonism has been implicated as a potentially important mechanism in T cell activation, thymic development, escape from an antiviral immune response, and autoimmunity. Although the actual mechanism by which antagonists exert their effect is still unresolved, various models seek to explain the paradox of how the recognition of altered peptide/MHC ligands leads to altered T cell responses. Quantitative models postulate that antagonist ligands inhibit T cell responses by competing for TCR engagement with the antigenic peptide. Due to their lower affinity and faster off-rate kinetics of TCR binding (23–25), antagonists are ineffective in initiating downstream signaling pathways but nevertheless reduce the rate of successful TCR engagements by the antigenic peptide (7, 12, 17,

19, 54, 55). In contrast, qualitative models characterize antagonists as ligands capable of delivering differential (8, 9, 16, 21, 26–28) or even negative signals to the T cell which subsequently inhibit T cell activation (13, 14, 31).

By coexpressing two TCRs of independent specificity on the same cell (Fig. 2), we were able to directly address the issue of whether a dominant-negative signal is induced by antagonist ligands. If antagonist peptides generate an intracellular dominant-negative signal, downstream from the site of TCR engagement, then antagonist ligands would be expected to inhibit the ability of a second, independent TCR to induce a cellular response. Our experimental system clearly shows that in cells expressing two TCRs, activation through one TCR is not inhibitable by engaging a second TCR with antagonist ligand (Fig. 5 C). These results are supported by the following observations, that (a) the antigenic peptide for the $V\alpha 2V\beta 10$ TCR, SSIEFARL, is a weak agonist (whose weak stimulatory capacity is definitely not due to weak MHC binding affinity as determined by RMA-S stabilization, data not shown). Therefore, a weak stimulation through $V\alpha 2V\beta 10$ should have been easily inhibitable by a putative negative signal through the $V\alpha 2V\beta 5$ TCR; (b) the ratio of the two TCRs expressed on the same cell favored the signals through the antagonized TCR since $V\alpha 2V\beta 5$, recognizing the antagonist, represented 60% of the TCR on the cell surface; and (c) all peptides were presented on the same APCs (agonist was loaded first to ensure its binding to K^b), therefore even a putative negative signal with only short-range efficacy should have been detected.

The fact that we observed no functional interference between different TCRs expressed on the same cell and no bystander TCR downregulation upon antigenic stimulation (Fig. 6, A and B) suggests that these two TCRs function independently, supporting the findings of Valitutti et al. on T cell clones (53). This is further supported by the biochemical data in hybridomas expressing two TCRs. Phosphorylated ζ and recruited phospho-ZAP-70 were only coprecipitated from TCRs engaged with their respective antigen (Fig. 6 C). These experiments argue that the phosphorylation of engaged TCRs does not spread to unengaged TCRs. It remains possible that individual cells expressing two different TCRs are intrinsically able to engage only one type of TCR. However, the simultaneous exposure of the cells to both antigenic peptides, the $V\alpha 2V\beta 5$ -specific SIINFEKL and the $V\alpha 2V\beta 10$ -specific SSIEFARL, presented on the same APC, induced concomitant internalization of both TCRs, indicating that both TCRs are functional on the same cell (data not shown).

Taken together, our data indicate that these antagonist ligands do not act by generating dominant-negative signals which inhibit the response of the entire cell. Therefore, we favor a quantitative model to explain TCR antagonism. Antagonist ligands have faster off-rate kinetics for TCR engagement than agonist ligands (23–25). Thus, antagonist ligands engage the TCR for a longer time than null peptides, but not long enough to induce the full signaling cascade for activation, as first suggested by the kinetic

proofreading model of McKeithan (18). Although this ineffective engagement does not lead to full activation, it does generate differential biochemical signals reflected by the differential phosphorylation patterns of ζ ($p21 > p23$; reference 28). However, as the data from our experimental system have shown, these differential signals did not represent an intracellular, negative signal per se. As a consequence of ineffective TCR engagement, antagonist ligands prevent agonist ligands from serially engaging and triggering enough TCRs to reach the critical threshold for activation (53). This competition for TCR engagement can be observed in the inhibition of TCR downregulation in the presence of antagonist ligands (19; Fig. 1 B and Fig. 5 B). Furthermore, the quantitative mechanism of antagonist action also explains the observation that T cell responses are less efficiently inhibited when antagonists and agonist peptides are presented on different APCs (9; data not shown). Conditions which reduce the direct competition between agonist and antagonist ligands reduce the efficiency of TCR antagonism. Antagonist ligands may also influence the reorganization of TCRs which may be required for full activation (55–57).

Although our data indicate that these antagonist ligands do not induce an intracellular negative signal, downstream from TCR engagement, there are potentially some limitations to the interpretation of our results. It is possible that for some unknown reason, the $V\alpha 2V\beta 10$ receptor could not be inhibited. In this regard, the $V\alpha 2V\beta 10$ TCR induced a strong degree of tyrosine phosphorylation but relatively little IL-2 secretion. This is somewhat surprising with regard to previously published data where a strong correlation between the degree of tyrosine phosphorylation and the extent of the functional responses was observed (26, 27, 58, 59). Whether this indicates that the $V\alpha 2V\beta 10$ receptor is intrinsically incapable of responding to a negative signal is difficult to know. A reciprocal study using antagonists for the $V\alpha 2V\beta 10$ TCR and agonist ligands for the $V\alpha 2V\beta 5$ receptor would have been informative, but antagonists for SSIEFARL (ligand for the $V\alpha 2V\beta 10$ receptor) have not been identified. It is also possible that antagonist peptides deliver negative signals which only inhibit other identical TCRs. If these putative negative signals are cis-limited, then we would not have been able to detect them using a second, independent receptor.

It is also theoretically possible that antagonist ligands generate dysfunctional TCRs, e.g., by inducing a conformation that disables the receptor from further activation. However, preexposure to antagonist peptides for 4 h alone did not render the $V\alpha 2V\beta 5$ expressing hybridomas cells unresponsive to further antigenic stimulation (data not shown), which is in accord with the findings of Preckel et al. for $CD8^+$ T cell clones (19). Moreover, since the signaling capacity of a second independent TCR was unaffected, it is also unlikely that antagonist peptides alter T cell responses by consuming limited amounts of signaling intermediates.

There are two reports of antagonists that can inhibit T cell responses when present at substoichiometric ratios relative to the antigenic peptide (10, 11). These types of antagonist peptides may not function by competition and con-

ceivably could deliver a negative signal to the responding T cell. Thus, there may be different classes of antagonist peptides that act through different mechanisms. One class that was studied here must be present in molar excess to exert an inhibitory effect and uses a competitive mechanism to inhibit T cell responses. Other classes of TCR antagonists, which are effective at substoichiometric ratios, may work by a dominant-negative mechanism.

Our data also show that neither the extent of inhibition (Fig. 1 B) nor the kinetics (Fig. 7) of TCR internalization fully accounted for the inhibition of IL-2 response in our experimental system. In fact, these data demonstrate that antagonist peptides can fully exert their inhibitory effect well after the initial phase of TCR downregulation, at a stage where there is no further net loss of surface TCRs (Fig. 8). Furthermore, the blocking studies with the anti-K^b antibody, Y3, demonstrated that the time window of susceptibility to antagonism correlated exactly with the time of commitment.

Previous studies have already pointed out the importance of a sustained signal for T cell commitment (60–62). Depending upon the amount of available antigen, the type of APC, the number of adhesion and costimulatory molecules present on the APC, and the state of differentiation of the responding cell, the length of time required for T cell commitment varied (63). For the V α 2V β 5 expressing T cell hybridoma, 12 h of continuous engagement with the agonist ligand was required for a commitment to IL-2 secretion. On the other hand, internalization of TCRs reached a steady-state after 2–3 h of antigen exposure (Fig. 7). After this point there may be continued TCR downregulation, but there is no further net loss of TCR from the

cell surface. These data suggest that antagonist peptides competitively inhibit agonist ligands from productively engaging TCR during the entire commitment period. Therefore, it is conceivable that small changes in the rate of TCR engagement, integrated over time, may have a significant impact on the final outcome of T cell activation.

Nevertheless, antagonist as well as partial agonist ligands induce biochemical signals, which are different from the signaling events generated by antigenic ligand used over a broad range of concentrations (28). Altered peptide ligands have been shown to induce altered phosphorylation of the ζ chain without stable phosphorylation of CD3 ϵ and ZAP-70 (26–28). That partial agonists induce differential biochemical signaling events as well as a long lived anergy in CD4⁺ T cell clones (20, 26, 64) has led some investigators to postulate that altered peptide ligands generate qualitatively different signals (26). In principle, the anergic state could result from the consumption of limiting signaling intermediates or from an imbalanced expression of transcription factors (65). Since T cell hybridomas, which are not IL-2 dependent for their proliferation, were used in these studies, we could not directly address this issue. Nevertheless, our experiments suggest that the inhibition of IL-2 production, a feature which characterizes most anergic T cells, is not mediated by a negative signal.

In summary, the data are consistent with the idea that the antagonist ligands studied here do not deliver negative signals, but rather compete with antigenic ligands for TCR engagement. This competition can effectively inhibit T cell responses by reducing the rate of TCR engagement throughout the entire commitment period.

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