Suppression of autoimmune disease after vaccination with autoreactive T cells that express Qa-1 peptide complexes

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The ability of autoreactive T cells to provoke autoimmune disease is well documented. The finding that immunization with attenuated autoreactive T cells (T cell vaccination, or TCV) can induce T cell–dependent inhibition of autoimmune responses has opened the possibility that regulatory T cells may be harnessed to inhibit autoimmune disease. Progress in the clinical application of TCV, however, has been slow, in part because the underlying mechanism has remained clouded in uncertainty. We have investigated the molecular basis of TCV-induced disease resistance in two murine models of autoimmunity: herpes simplex virus-1 (KOS strain)–induced herpes stromal keratitis and murine autoimmune diabetes in non-obese diabetic (NOD) mice. We find that the therapeutic effects of TCV depend on activation of suppressive CD8 cells that specifically recognize Qa-1–bound peptides expressed by autoreactive CD4 cells. We clarify the molecular interaction between Qa-1 and self peptides that generates biologically active ligands capable of both inducing suppressive CD8 cells and targeting them to autoreactive CD4 cells. These studies suggest that vaccination with peptide-pulsed cells bearing the human equivalent of murine Qa-1 (HLA-E) may represent a convenient and effective clinical approach to cellular therapy of autoimmune disease.

Introduction
Autoreactive T cells are part of the normal T cell repertoire (1, 2) and can provoke autoimmune disease after abnormal environmental stimuli, as documented by adoptive transfer of autoreactive T cells in murine models of diseases (3, 4). The cellular mechanisms that may inhibit this process are not well understood. The suggestion that regulatory T cells might be harnessed to inhibit disease came from findings that immunization with attenuated autoreactive T cells conferred resistance to subsequent disease induction (3, 5). This procedure, termed T cell vaccination (TCV) (6), has been examined in several animal models of tissue-specific autoimmune diseases (6, 7) and in small studies of MS patients (8, 9). Progress in the clinical application of TCV, however, has been slowed, in part, because the underlying mechanism of TCV is not well understood.

Recent studies have suggested that the protective effect of TCV may reflect induction of regulatory CD8 cells (10–16). For example, depletion of CD8 cells prior to TCV abrogates its protective effect (12) and T regulatory activity following immunization with lymphocytes is mediated by CD8 cells (17). The mechanism(s) that lead to induction of suppressive CD8 cells remains unclear, however. Although autoreactive T cell receptors (TCRs) have been implicated in the genesis of immunoregulatory cells, the molecular interactions that underlie presentation of autoreactive TCR to the immune system in a way that induces CD8-dependent suppression have not been clarified.

Methods
Mice. Six- to eight-week-old female C.AL-20 (BALB/c-Igh), BALB/c, BALB/c–IFN-γ−/−, BALB/c–IL-10−/−, BALB/c–RAG-2−/−, C.B17–SCID, NOD, 129Sv, and 129SvIFN-γR−/− mice were purchased from The Jackson Laboratory (Bar Harbor, Maine, USA). All mice were housed in microisolator cages in the Dana-Farber Animal Facility and, when appropriate, in biosafety level 2. In all attempts at vaccinating using isolated TCR peptides have not met with uniform success, and the effects of blocking Ab’s against classical MHC class-Ia molecules expressed by autoreactive CD4 cells used for TCV have been inconclusive. Lysis of autologous CD4 cells by CD8 cells from vaccinated MS patients has been reported to be prevented by class Ia Ab (8, 13), while other studies indicate that inhibitory interactions between CD8 cells and Vβ8+ murine CD4 clones is blocked by Ab to the class Ib molecule Qa-1 (17). Although the in vitro responses of CD8 cells is restricted by Qa-1 expressed on target cells (10, 17), the contribution of Qa-1 peptide complexes to the development of inhibitory activity after TCV is not clear.

We have analyzed the effects of TCV in two murine models of autoimmune disease: herpes stromal keratitis (HSK), which is a clinical consequence of corneal infection by herpes simplex virus 1 (HSV-1) and represents a leading cause of human blindness (18), and non-obese diabetic (NOD) mice, a murine model of type 1 diabetes. We show that the therapeutic effects of TCV depend on induction of suppressive CD8 cells that specifically recognize Qa-1 peptide complexes expressed by autoreactive CD4 cells in two murine models of autoimmune disease. These studies provide new insight into the molecular basis of this inhibitory T cell–T cell interaction that is mediated by TCV and suggest new approaches to this cellular therapy.
the experiments each group represents at least eight mice, and each experiment was performed at least twice.

Ocular infections and scoring of HSK. Mice were infected with 4 x 10^5 PFU of HSV-1 (KOS strain) in the right eye, and the extent of the disease was scored on different days within 14–15 days after infection as described (19). Disease score is summarized as the HSK index, which equals severity (mean clinical score) multiplied by percentage of incidence multiplied by ten, where severity of clinical stromal keratitis was quantified based on the degree of corneal opacity: 1, less than or equal to 25% of cornea; 2, less than or equal to 50%; 3, less than or equal to 75%; 4, 75–100%. (For all disease severity data points SD was < 0.5.) Incidence was the percentage of mice with a severity score greater than or equal to 1.

CD4+ TCV protocol. CD4+ T cells were purified from the right cervical draining LNs of C.AL-20 mice 10 days after infection of the right eye with 4 x 10^5 PFU HSV-1 (KOS) by magnetic negative selection (Dynabeads; Dynal Inc., Lake Success, New York, USA) using anti-CD8 (53-6.7), anti-B220 (RA3-6B2), anti-Gr-1 (RB6-8C5), and anti–Mac-1 (M1/70) mAb's. CD8 cells (5 x 10^6/mouse) in 200 µl RPMI medium, irradiated at 30 Gy, and resuspended at 1.25 x 10^7 cells/ml in RPMI. Mice were injected in the tail vein with 200 µl of the cell suspension (2.5 x 10^6 cells/mouse), and 3 weeks later were ocularly infected with HSV-1 (KOS). In some experiments cells from the C1-6 CD4+ T cell clone reacting to β2µ peptide and cross-reacting to a γ2ab peptide and cross-reacting to a γ2ab peptide were used as controls. In experiments where sorted CD4+ cells were used, 10^6 purified CD4+ T cells of draining LNs of ocularly HSV-1–infected mice were injected per mouse. For TCV experiments in NOD mice pancreatic LNs from 15-week-old female NOD donors were obtained during the insulitic stage of disease in NOD mice and were treated with conA for 40 hours followed by irradiation (30 Gy) before i.v. injection (2.5 x 10^6 cells/mouse) into 11-week-old female NOD recipients. Vaccinated animals were observed for onset of diabetes; disease progression was measured by serum (or urine, where indicated) glucose levels and mortality.

Cell preparations and purifications were performed in the same way as in TCV for HSK.

In blocking experiments, cells were incubated with a mouse IgG1-Ab that binds to Qa-1β (anti-Qa-1β was kindly provided by M. Soloski, Johns Hopkins University, Baltimore, Maryland, USA), or isotype-matched control Ab (IgGκ; A112-2), or anti–MHC class I Ab (H-2Dβ; 34-2-12) for 1 hour at 4°C after 40 hours of in vitro stimulation with conA (5 µg/ml) and prior to irradiation and injection into mice.

In some experiments, O3 cells were used for TCV after incubation with peptides Vβ8.1 (LLSWVALFL) or L9D (LLSWVALFD) (New England Peptide Inc., Gardner, Massachusetts, USA) as follows: O3 cells were stimulated for 40 hours at 37°C in RPMI/10% FCS with conA (5 µg/ml) before addition of Vβ8.1 peptide or L9D peptide (final concentration 0.5 µg/ml) for 1 hour. After cells were washed three times with RPMI and irradiated (30 Gy), 2 x 10^6 cells (in 200 µl RPMI) were injected per CAL-20 mouse. In one experiment the peptides were dissolved in 25 µl CFA per mouse and subcutaneously injected into the back of anesthetized CAL-20 mice that were 3 weeks later ocularly infected with HSV-1 (KOS) and scored for HSK.

Transfer experiments. For transfer of CD8+ cells, these cells were purified from LNs and spleens 2 weeks after TCV by negative magnetic selection (Dynabeads; Dynal Inc.) with anti-CD4 (GK1.5), anti-B220 (RA3-6B2), anti–Gr-1 (RB6-8C5), and anti–Mac-1 (M1/70) mAb's. CD8 cells (5 x 10^6/mouse) in 200 µl PBS (97% pure CD8 as confirmed by FACS) were injected in the tail vein of recipient animals.

In some experiments CD8+ cells from previously vaccinated WT, IL-10−/−, and IFN-γ−/− mice were adoptively transferred into recipient CB.17-SCID mice together with CD8-depleted LN cells from WT BALB/c mice. For the experiments where T cells are transferred into RAG-2−/− recipients 6 days before analysis, 2 x 10^6 cells from donors vaccinated with Vβ6.1 or Vβ8.1 CD4+ cells were admixed with 10^4 conA-activated Vβ6.1 or Vβ8.1 CD4+ cells as indicated.

In vivo depletions. Groups of mice were depleted of CD8 cells prior to TCV and ocular infection by intraperitoneal injections (six times) with monoclonal anti-CD8 Ab (2.43; 25 µg/dose). The depletion (97%) was confirmed by FACS analysis.

**Figure 1**

Qa-1β expression on CD4+ cell vaccine is essential for protection. Effects of monoclonal anti–Qa-1 Ab on TCV in vitro (A) and in vivo (B). (A) C.AL-20 mice were vaccinated with CD4+ cells isolated from HSV-1–infected C.AL-20 mice. A group of mice was vaccinated with the same CD4 cells but incubated with blocking anti–Qa-1 mAb (open squares) or isotype control (filled circles) or were nonvaccinated controls (open circles) as described in Methods. Each group consisted of ten mice, and results are representative of three independent experiments. (B) C.AL-20 mice were intraperitoneally injected with anti–Qa-1 mAb (10 µg/mouse) at the time of TCV (open squares) or isotype control (filled circles) or were nonvaccinated controls (open circles). Two weeks later, vaccinated mice were ocularly infected with HSV-1 (KOS) and scored for HSK. Each group consisted of ten mice, and results are representative of three independent experiments. (C) Qa-1 sorting. C.AL-20 mice were vaccinated with CD4+ cells isolated from HSV-1–infected C.AL-20 mice. These CD4+ cells were sorted according to their level of Qa-1 surface expression (high, filled triangles, versus low, filled squares) or nonvaccinated controls (open circles). Both groups were ocularly infected with HSV-1 (KOS) and scored for HSK as described in Methods. Each group consisted of 10–20 mice. Data are representative of two independent experiments.
Flow cytometry. Flow-cytometric analysis was performed to measure the Qa-1 expression on the surface of the CD4 cells using a Coulter EPICS XL flow cytometer. LN cells or, in sorting experiments, purified CD4 cells were stained with anti-Qa-1b and anti-CD4 phycoerythrin-conjugated (PE–conjugated) Ab (PharMingen, San Diego, California, USA) and with an unconjugated primary Ab to Qa-1 (KOS) followed by an anti-mouse IgG–FITC–conjugated mAb (PharMingen). CD4 cells were measured with a fluorescence-activated cell sorter (FACS) for their Qa-1 expression or sorted according to the level of Qa-1 expression on their surface.

Qa-1 tetramers. Qa-1α and β2-microglobulin were produced as inclusion bodies in Escherichia coli BL21 (DE3) carrying either the pET-23a/2-microglobulin or the β2-microglobulin plasmid. The inclusion bodies were purified and dissolved in urea buffer. The monomeric Qa-1α–2-microglobulin complexes were formed by combining the Qa-1, β2-microglobulin, and the peptide in an arginine-folding buffer. The complexes were then purified on a Superdex-200 gel-filtration column (Pharmacia Biotech Inc., Piscataway, New Jersey, USA) before biotinylation with the BirA enzyme (Avidity, Denver, Colorado, USA). The biotinylated monomeric complexes were purified again with Superdex-200 gel-filtration column. Streptavidin-PE (Molecular Probes Inc., Eugene, Oregon, USA) was added to form the tetrameric reagent in a 4:1 ratio. The tetramer efficiency was measured by HPLC.

To determine whether the protective effects of TCV depend on expression of the class Ib molecule Qa-1 by activated CD4 cells we examined the effects of monoclonal anti–Qa-1 Ab on protection. Vaccination using cells that had been incubated with anti–Qa-1 Ab failed to efficiently inhibit disease compared with CD4 cells incubated with an isotype-matched control Ab, which induced strong disease resistance (Figure 1A). Injection of monoclonal anti-Qa-1 blocking Ab in vivo along with TCV also largely abrogated its suppressive effects (Figure 1B). Finally, vaccination with FACS-sorted Qa-1α–CD4 cells (mean fluorescence intensity greater than 100) yielded complete (100%) protection, while vaccination with Qa-1α–CD4 cells (mean fluorescence intensity less than 40) yielded only partial (50%) protection. These results suggest that TCV-induced protection is dependent on Qa-1 expression on CD4 cells.
did not exert detectable disease protection (Figure 1C). Both Qa-1hi and Qa-1lo CD4 subpopulations were equally activated according to expression of activation-associated surface markers CD44 and CD62, as judged by forward light scatter upon FACS analysis (not shown). Possibly, Qa-1hi cells may represent a subpopulation of activated CD4 cells that displays relatively high levels of non-Qdm peptides, including those derived from the TCR. An additional property of these cells that may be relevant to their immunogenicity is the increased levels of surface TCR (but not other surface receptors, such as CD4; data not shown).

**TCV-induced protection depends on TCR Vb peptide–Qa-1 complexes expressed by CD4 cells**

The autoreactive C1-6 TCR (Vβ8.1Vα11) that confers HSK has been defined in polyclonal populations and tested in mice that carry the C1-6 TCR transgene (21). Vaccination using purified Vβ8+ CD4 cells from normal (Balb/c) mice resulted in 100% protection from HSK, while vaccination using Vβ8-depleted CD4 cells or purified Vβ2+ CD4 cells did not protect from HSK (Figure 2A). We compared the protective effects of three different Balb/c Vβ8+ TCRs: Vβ8.1+class II–restricted C1-6 TCR (20, 21); DO11.10, a Vβ8.2+ TCR specific for an Ova-derived peptide/class IIβ (22); and O3, a Vβ6+ TCR specific for a second Ova-derived peptide/class IIβ (23). Vaccination with activated, irradiated Vβ8.2+ DO11.10 cells and Vβ8.1+ C1-6 cells successfully inhibited HSK, while vaccination with Vβ6+ O3 cells did not (Figure 2B).

**TCV-induced protection depends on Vβ peptide binding to Qa-1b**

These results suggested that the TCV ligand comprised a Vβ8-derived peptide bound to Qa-1b on activated CD4 cells. Analysis of Vβ8.1 revealed a single motif (LLSWVALFL) that might allow refolding of purified Qα-1b (but not MHc class I molecules). Qα-1b–folded activity with this peptide was confirmed according to HPLC analysis of Vβ8 peptide: Qa-1 complexes and tetramers (24). This finding allowed a direct test of the role of this Vβ–Qa-1 ligand in TCV: Did loading of Vβ6+ O3 cells with the Qa-1b–binding, Vβ8-derived peptide (which were themselves inactive; Figure 2B) confer protective activity on this clone? We found that vaccination with activated irradiated Vβ6+ CD4 cells incubated with Vβ8.1 peptide resulted in 100% protection (Figure 2C), while vaccination with Vβ6+ CD4 cells incubated with a peptide homologue containing a L→D amino acid exchange (LLSWVALFD) that abolished peptide-induced folding of Qa-1b had no protective effect (Figure 2C). The possibility that the Vβ8.1 peptide itself might confer protection independent of its interaction with Qa-1 was tested by injection of Vβ8.1 peptide admixed with Qa-1+ O3 cells or CFA; these stimuli had no effect on HSK (data not shown).

**TCV-based inhibition of diabetes in NOD mice**

We asked whether vaccination with pancreatic LN T cells obtained during the insulitis stage of disease in NOD mice might preempt subsequent Iddm. This was the case, as measured by disease progression (serum glucose) and mortality (Figure 3). Pathogenic islet-reactive T cells are enriched in draining pancreatic LNs, particularly early in disease in NOD mice (25). NOD mice vaccinated with CD4 cells from pancreatic LNs but not inguinal or mesenteric LNs are protected from disease progression (Table 1).

**TCV-associated induction of suppressive CD8 cells**

The CD8 cell response is essential for TCV-induced protection against HSK because CD8 depletion abolished the protective effect of TCV on HSK (Figure 4A), and purified CD8 cells from donors vaccinated

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**Table 1**

TCV-induced protection of DM in NOD mice

<table>
<thead>
<tr>
<th>Source of LN T cells</th>
<th>Cervical</th>
<th>Pancreatic T cells</th>
<th>Pancreatic CD4 cells</th>
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<tbody>
<tr>
<td>NOD (female) mice</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Nonvaccinated</td>
<td>17 wk</td>
<td>30 wk</td>
<td>20 wk</td>
</tr>
<tr>
<td>Mesenteric</td>
<td>17 wk</td>
<td>30 wk</td>
<td>20 wk</td>
</tr>
<tr>
<td>Mortality</td>
<td>50%</td>
<td>90%</td>
<td>0%</td>
</tr>
<tr>
<td>Elevated glucose</td>
<td>75%</td>
<td>100%</td>
<td>60%</td>
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NOD female mice (11 weeks old) were vaccinated with T or CD4 cells as described in Methods. Each experimental group consisted of 40–90 mice.

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**Figure 4**

TCV induces CD8+ T suppressor cells: analysis in HSK. (A) CD8 depletion before TCV. Depletion of CD8 T cells abolishes TCV-induced protection. C.AL-20 mice were depleted of CD8 T cells by intraperitoneal injection of depleting anti-CD8 Ab (filled circles) or isotype control (filled squares) prior to TCV and ocular infection with HSV-1. A group of mice was not vaccinated for comparison (open circles). Each group consisted of ten mice, and results are representative of three independent experiments. (B) CD8 cell transfer of disease resistance. Protection can be transferred by CD8 cell transfer from vaccinated mice. C.AL-20 mice received CD8 cells from donors vaccinated with either keratogenic Vβ8+ C1-6 cells (filled circles) or OVA-specific Vβ6+ O3 cells (open circles). The recipients were ocularly infected with HSV-1 (KOS) and scored for HSK. Each group consisted of ten mice, and results are representative of three independent experiments.
We have analyzed TCV in the context of HSK and NOD disease because these models have several advantages over other murine models of autoimmunity. Since HSV-1 (KOS) infection of BALB/c mice is routinely followed by a CD4-dependent autoimmune attack against corneal tissue, this model allowed analysis of TCV-induced protection against the autoimmune portion of an antiviral response after infection. We also analyzed the effects on the spontaneous development of diabetes in NOD mice. Both murine models allow testing of the efficacy of TCV on the development of autoimmunity in settings that more closely mimic the clinical development of autoimmunity in the absence of deliberate immunization with self peptide. In both cases, in vivo depletion of CD8 cells prior to TCV abolished its protective effects, prompting us to further probe the role of CD8 cells in regulating disease development. Transfer of disease protection into naive recipient mice by CD8 cells from donors that had received TCV delineated this subset as a central mediator of the inhibitory effects of TCV. The ability of CD8 cells from mice vaccinated with Vβ6+ or Vβ8+ CD4 cells to inhibit the expansion of CD4 cells in a Vβ-specific fashion indicated that a direct CD8–CD4 inhibitory interaction may be sufficient to mediate the disease-inhibitory effects of TCV. It remains possible that other (auxiliary) regulatory cells generated by TCV may also contribute to protection. Resistance to antigen-induced experimental autoimmune encephalomyelitis (EAE), for example, is also associated with regulatory CD4+ T cells that may recognize peptides from autoreactive TCR (26) and enhance generation of CD8 T cells (27).

**Surface ligands on autoreactive CD4 cells induce regulatory CD8 cells**

EAE-inducing T cell lines specific for different components of myelin basic proteins (MBPs) did not prevent disease induced by different MBPs (28). In contrast, TCV against EAE cross-protected against autoimmune arthritis (29). These and other studies indicated that the protective effects of TCV do not depend on the antigen specificity of CD4 cells used for vaccination. Qa-1 can present a number of foreign (30) and self peptides (31). Although regulatory CD8 cells generated by TCV in MS patients did not recognize MBP-reactive CD4 clones bearing TCR chains that differed from the immunizing MBP, these CD8 clones recognized several distinct MBP-reactive CD4 clones that shared the same (Vβ15) chain (14). Jiang et al. have demonstrated the important role of Vβ-specific anti-MBP clones in the induction of TCV-based protection: vaccination with

**Discussion**

The precise mechanism underlying the effects of TCV has not been elucidated since 1981 when Cohen and colleagues coined the term to describe the administration of activated attenuated autoimmune T cells (6). Although these and other studies have documented the inhibitory effects of TCV on the progression of autoimmune disease in experimental animal models, the underlying mechanism has remained poorly understood. As a result, approaches based on TCV have had varying degrees of success in clinical trials. The studies described here are aimed at a mechanistic understanding of TCV. They suggest that TCR Vβ-derived peptides associated with Qa-1 (the murine homologue of HLA-E) on activated autoreactive CD4 cells can activate CD8-dependent suppression and inhibit autoimmunity.
three different V\(\beta\)8+ MBP-reactive CD4 T cell clones protected against disease induced by another MBP-reactive V\(\beta\)8+ CD4 T cell clone, while V\(\beta\)6+ MBP-reactive CD4 T cell clones did not (12).

Previous studies of HSK induced by HSV-1 (KOS) have established that pathogenic autoreactivity resides in a numerically small subpopulation of CD4 cells that express a V\(\beta\)8.1+ TCR and cross-react with an HSV-1 viral epitope (19, 20). Vaccination with activated (polyclonal) V\(\beta\)8+ CD4+ T cells from nonimmune mice conferred complete protection against HSK as did vaccination with V\(\beta\)8.2+ DO11.10 cells (reactive to an OVA peptide) and V\(\beta\)8.1+ C1-6 cells (see Figure 2B). Protection against HSK was not achieved after vaccination with conA-activated CD4+ T cells (containing the full spectrum of V\(\beta\)\_s) or purified V\(\beta\)2- or V\(\beta\)6+ cells. Both V\(\beta\)8.1 and V\(\beta\)8.2 share an identical Qa-1+–binding motif (LLSWVALFL), which allows refolding of Qa-1–β\(\alpha\)m, but not MHC class I–β\(\alpha\)m molecules in vitro (24). A V\(\beta\)6+ clone of CD4 cells that expresses high levels of Qa-1 after activation (but lacks a Qa-1+–binding motif within the V\(\beta\)6 protein) was programmed to induce HSK protection after pulsing with the LLSWVALFL peptide, but not an L9D mutant peptide (which does not allow refolding of Qa-1–β\(\alpha\)m; see Methods for incubation conditions). Since immunization with the V\(\beta\)8-derived peptide alone (in CFA) was not protective, these data indicate that cellular vaccination provides a peptide–Qa-1 complex that may specifically activate and expand Qa-1–restricted suppressive CD8 cells that can target autoreactive CD4 cells that express Qa-1-containing ligands. Although these results delineate the essential role of a Qa-1–V\(\beta\)8 peptide complex in TCV-induced resistance, they do not define the potential role(s) of other Qa-1+–binding peptides, including naturally occurring Qdm and HSP-60. Although Qdm, HSP-60, and other Qa-1–peptide complexes are not sufficient for TCV-based protection in the absence of the V\(\beta\)8–Qa-1 ligand, they may, for example, regulate peptide exchange. We postulate that a shift in the proportions of the high-affinity Qdm–Qa-1 complex in favor of the lower-affinity HSP-60–Qa-1 complex 48 hours after activation of CD4 cells may facilitate exchange between exogenous peptides and the Qa-1 molecule (32–34).

Studies attempting to address the role of classical MHC class I molecules in TCV have yielded conflicting results (8, 13, 35). Qa-1 was initially described as a surface molecule expressed on activated (polyclonal) V\(\beta\)8+ T cells from nonimmune mice and was initially described as a surface molecule expressed on activated (polyclonal) V\(\beta\)8+ T cells from nonimmune mice. An additional property of Qa-1 hi cells was the ability to bind class I molecules in TCV have yielded conflicting results (8, 13, 35). Qa-1 reflects the situation in vivo, however. Since optimal protection was achieved after conA-dependent vaccination with the LLSWVALFL peptide, but not an L9D mutant peptide; see Methods for incubation conditions. Since immunization with the V\(\beta\)8-derived peptide alone (in CFA) was not protective, these data indicate that cellular vaccination provides a peptide–Qa-1 complex that may specifically activate and expand Qa-1–restricted suppressive CD8 cells that can target autoreactive CD4 cells that express Qa-1-containing ligands. Although these results delineate the essential role of a Qa-1–V\(\beta\)8 peptide complex in TCV-induced resistance, they do not define the potential role(s) of other Qa-1+–binding peptides, including naturally occurring Qdm and HSP-60. Although Qdm, HSP-60, and other Qa-1–peptide complexes are not sufficient for TCV-based protection in the absence of the V\(\beta\)8–Qa-1 ligand, they may, for example, regulate peptide exchange. We postulate that a shift in the proportions of the high-affinity Qdm–Qa-1 complex in favor of the lower-affinity HSP-60–Qa-1 complex 48 hours after activation of CD4 cells may facilitate exchange between exogenous peptides and the Qa-1 molecule (32–34).

Studies attempting to address the role of classical MHC class I molecules in TCV have yielded conflicting results (8, 13, 35). Qa-1 was initially described as a surface molecule expressed on activated CD4 cells that interacted with suppressive CD8+ cells (36–39), either alone or through induction of regulatory CD4 cells (26). Qa-1+–restricted inhibitory activity of CD8 cells (Figure 6) may likewise reflect in vivo elimination of activated CD4 cells that display Qa-1–peptide complexes. Although the precise suppressor-effector mechanism remains to be determined, these data establish the central role of Qa-1–peptide complexes in the induction of TCV-based protection in two murine disease models and suggest potentially effective approaches to TCV in clinical settings, including the use of “universal” HLA-E+ cell lines pulsed with target peptides as a potentially convenient and effective approach to immunosuppressive cellular therapy.

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Regulatory sublineages of CD4 and CD8 cells
Analysis of regulatory T cells in tissue-specific autoimmune disease has implicated a CD25+ subpopulation of CD4 cells in some models and IL-4–producing NK T cells in others (41). Previous studies of murine EAE, however, showed that TCV might result in induction of CD8+ T cells that lyse or damage autoreactive CD4+ T cells in vitro (5, 17). The tempo of this inhibitory response may be contrasted with naturally occurring CD4CD25+ regulatory T cells that can interrupt expansion of self-reactive T cells. The development of Qa-1+–restricted regulatory CD8 activity more closely resembles the development of CD8 CTL rather than regulatory CD4 cells. Once generated, CD8 CTL lyse virally infected target cells through TCR-dependent recognition of viral peptides–class Ia MHC, while Qa-1+–restricted CD8 cells may eliminate a subpopulation of activated autoreactive CD4 cells through TCR-dependent recognition of self- peptide–Qa-1 complexes (42, 43). In contrast, CD4CD25+ regulatory activity generally acts at the early stages of primary responses (44, 45), either alone or through induction of regulatory CD8 cells (26).

Analysis of T cells expressing a TCR transgene specific for insulin associated with Qa-1 has demonstrated that these cells are selected in the thymus through interactions with Qa-1+–self peptides expressed primarily on hematopoietic cells. Both selection and antigen-recognition depend on expression of the TCR\(\alpha\)/\(\beta\) and do not require NK2 receptors (46). CD8 cells that recognize Qa-1 associated with the insulin peptide efficiently kill Qa-1–expressing lymphoblastoid target cells in the presence of intact soluble insulin, that is following transporter associated with antigen presentation–independent (TAP-independent) processing and presentation of insulin-derived peptide by Qa-1 on activated target cells (47). Possibly, Qa-1+–restricted inhibitory activity of CD8 cells (Figure 6) may likewise reflect in vivo elimination of activated CD4 cells that display Qa-1–peptide complexes. Although the precise suppressor-effector mechanism remains to be determined, these data establish the central role of Qa-1–peptide complexes in the induction of TCV-based protection in two murine disease models and suggest potentially effective approaches to TCV in clinical settings, including the use of “universal” HLA-E+ cell lines pulsed with target peptides as a potentially convenient and effective approach to immunosuppressive cellular therapy.
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