Induction and Exhaustion of Lymphocytic Choriomeningitis
Virus-specific Cytotoxic T Lymphocytes Visualized Using
Soluble Tetrameric Major Histocompatibility Complex
Class I–Peptide Complexes

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Summary
This study describes the construction of soluble major histocompatibility complexes consisting
of the mouse class I molecule, H-2D[b], chemically biotinylated β2 microglobulin and a peptide
epitope derived from the glycoprotein (GP; amino acids 33–41) of lymphocytic choriomeningitis virus (LCMV). Tetrameric class I complexes, which were produced by mixing the class I complexes with phycoerythrin-labeled neutravidin, permitted direct analysis of virus-specific
cytotoxic T lymphocytes (CTLs) by flow cytometry. This technique was validated by (a)
staining CD8+ cells in the spleens of transgenic mice that express a T cell receptor (TCR) specific
for H-2D[b] in association with peptide GP33–41, and (b) by staining virus-specific CTLs in the
cerebrospinal fluid of C57BL/6 (B6) mice that had been infected intracranially with LCMV-
DOCILE. Staining of spleen cells isolated from B6 mice revealed that up to 40% of CD8+ T
cells were GP33 tetramer+ during the initial phase of LCMV infection. In contrast, GP33 tet-
ramers did not stain CD8+ T cells isolated from the spleens of B6 mice that had been infected
2 mo previously with LCMV above the background levels found in naïve mice. The fate of vi-
rus-specific CTLs was analyzed during the acute phase of infection in mice challenged both in-
tracranially and intravenously with a high or low dose of LCMV-DOCILE. The results of the
study show that the outcome of infection by LCMV is determined by antigen load alone. Fur-
thermore, the data indicate that deletion of virus-specific CTLs in the presence of excessive an-
tigen is preceded by TCR downregulation and is dependent upon perforin.

The ability to clear infections with noncytopathic vi-
ruses is predominantly attributed to CD8+ CTLs.
CTLs recognize infected cells via an interaction between
TCRs and their corresponding ligands, class I MHC mole-
cules (1). MHC class I molecules are expressed on the cell
surface in association with self or pathogen-derived pep-
tides that are generated intracellularly by proteolytic degra-
dation of the parent proteins (2). After recognition of an
infected cell, naïve CTLs become activated, proliferate, and
attain not only the ability to lyse infected cells, but also the
ability to produce IFN-γ (3). Although IFN-γ may have a
direct antiviral effect (4, 5), it has also been shown to im-
prove the efficiency of antigen presentation by class I mole-
cules (6–9) thereby promoting the induction of a CTL re-
sponse and improving the efficiency with which CTL can
recognize their infected targets.

CTLs have been shown to be essential for the recovery
of mice from the acute phase of infection with the noncy-
topathic lymphocytic choriomeningitis virus (LCMV; ref-
ERENCE 10). So far, it has not been possible to follow the ki-
netics of appearance and disappearance of antigen-specific
effectors CTLs during the acute phase of both low- and
high-dose LCMV infections in non-TCR-transgenic mice.
A recent study by Altman et al. (11) described a method,
using tetrameric soluble MHC class I–peptide complexes,
for the identification of antigen-specific CD8+ cells in the
PBMCs of HIV-infected humans. This study describes an adaptation of this method for the identification of antigen-specific CD8+ cells in B6 (H-2b) mice infected with LCMV. In this case, soluble peptide–MHC complexes were generated using the mouse class I heavy chain D, chemically biotinylated human β2 microglobulin (β2M) and the LCMV peptide epitope glycoprotein (GP)33–41 (GP33-KAVYNFATC). Fluorescence-labeled tetrameric complexes were subsequently produced by mixing the biotinylated complexes with phycoerythrin-labeled neauradinin. Peptide GP33–41 (GP33) was used for the purposes of this study since, after LCMV-W-E infection of C57BL/6 mice, most CTL activity (~50–60%) is directed towards this epitope. Two other epitopes, defined by residues 276–286 of the viral glycoprotein (GP276) and residues 396–404 of the viral nucleoprotein (NP396), represent 10–20% and 20–30% of the total CTL activity, respectively (12–17).

The tetrameric class I–peptide complexes, which stained CTLs specifically, were used to follow the fate of GP33-specific CD8+ T cells in mice during the acute phase of LCMV infection. This study demonstrates the accumulation of stained virus-specific CTLs in the CSF and spleens of mice after intracranial infection with a substrain of LCMV-WE called LCMV-DOCILE, and in the spleens of mice infected intravenously with the same virus. In both cases, the accumulation of GP33-specific CTLs after both low- and high-dose infection with LCMV-DOCILE, was monitored in relation to the capacity of the cells to produce IFN-γ; to mediate cytotoxic activity, and to mediate virus clearance.

Materials and Methods

Mice. C57BL/6 (H-2b), 318 TCR transgenic mice (18), and perforin-deficient mice (PKOB; reference 10) were obtained from the Institut für Zuchthygiene (Tierspital Zürich, Switzerland). All mice were kept in a specific pathogen-free mouse housing facility.

Peptides. The LCMV peptides GP33–41, nucleoprotein (NP)396–404, and GP276–286 were purchased from Neosystem Laboratoire (Strasbourg, France).

Virus. The LCMV-DOCILE strain was a gift from C. Pfau (C. Pfau, Rensselaer Polytechnic Institute, Troy, NY) and was grown using Madin Darby canine kidney (M DCK) cells. The LCMV strain DOCILE was constructed by PCR of the gene fragment encoding the α1, α2, and α3 domains and the first two amino acids of the transmembrane domain (residues 1–276) using the primers 5'-CATGACATATGGCCACACTCGATGCGGTATTTC-3' and 5'-CATATGAAGCTTTTATCAAGTCTCCATCTCGAGT-3'. The resulting fragment was cut with the restriction enzymes NdeI and HindIII (New England Biolabs, Beverly, MA) and cloned into the expression vector pGM7 (a pET derivative; reference 27). The expression plasmid was transformed into the Escherichia coli strain BL21 (DE3) pLysS (Novagen, Madison, WI) and grown at 37°C in Luria-Bertani medium containing 100 μg/ml ampicillin (Sigma Chemical Co., St. Louis, MO). Protein expression was induced at midlog phase (A600 = 0.6) with 0.5 mM isopropyl-β-D-thiogalactosidase (IPTG). After 5 h, the cells were harvested and lysed by an overnight freeze thaw step (~8°C). Inclusion bodies were isolated as follows. The lysed pellet from 1 liter of culture was resuspended in 40 ml of 25% sucrose in 10 mM 2-mercaptoethanol, and 10% Con A supernatant. The cultures were restimulated at 14-d intervals using irradiated peptide-pulsed RMA-S cells as APCs at a responder/APC ratio of 10:1. These CTL lines were found to be of a single specificity after three rounds of restimulation.

Virus Titeration. LCMV titers in spleens were determined as previously described (24).

Isolation of Cerebrospinal Fluid. Cerebrospinal fluid was isolated from mice which had been infected after intracranial inoculation with LCMV-DOCILE as described previously (25, 26).

Protein Expression and Refolding. The H-2Db expression vector was constructed by PCR of the gene fragment encoding the α1, α2, and α3 domains and the first two amino acids of the transmembrane domain (residues 1–276) using the primers 5'-CATGACATATGGCCACACTCGATGCGGTATTTC-3' and 5'-CATATGAAGCTTTTATCAAGTCTCCATCTCGAGT-3'. The resulting fragment was cut with the restriction enzymes NdeI and HindIII (New England Biolabs, Beverly, MA) and cloned into the expression vector pGM7 (a pET derivative; reference 27). The expression plasmid was transformed into the Escherichia coli strain BL21 (DE3) pLysS (Novagen, Madison, WI) and grown at 37°C in Luria-Bertani medium containing 100 μg/ml ampicillin (Sigma Chemical Co., St. Louis, MO). Protein expression was induced at midlog phase (A600 = 0.6) with 0.5 mM isopropyl-β-D-thiogalactosidase (IPTG). After 5 h, the cells were harvested and lysed by an overnight freeze thaw step (~8°C). Inclusion bodies were isolated as follows. The lysed pellet from 1 liter of culture was resuspended in 40 ml of 25% sucrose in 10 mM Tris (pH 8.0), 1 mM EDTA, 1 mM PM SF, and 10 mM dithiothreitol (DTT) and sonicated until no longer viscous. The solution was then centrifuged at 18,000 rpm for 30 min before the supernatant was discarded and the pellet resuspended in 50 ml of 50 mM Tris, pH 8.0, 25% sucrose, 1% N oni d P-40, 0.5% sodium deoxycholate, 5 mM EDTA, and 2 mM DTT. The insoluble material was again recovered by centrifugation at 18,000 rpm for 30 min and resuspended in 50 ml of 25 mM Tris (pH 8.4), 2 mM DTT, 2 mM NaCl, and 2 mM urea. After centrifugation as above, the washed inclusion body pellet was resuspended in 5 ml of 20 mM Tris (pH 7.5), 150 mM NaCl, 0.5 mM PM SF, and stored at -20°C. The inclusion bodies were solubilized in 8 M urea at 4°C for 6 h immediately before refolding. β2M was pro-
duced as described above from the vector pHNβ2M (28). After solubilization of the β2M inclusion bodies in 6 M guanidinium HCl (pH 8.2), the protein was biotinylated using a fivefold molar excess of N-hydroxysuccinimide biotin (Sigma Chemical Co.). After a 30-min incubation at room temperature and a further 1-h incubation on ice, the biotinylation reaction was stopped by adding NH₄Cl to a final concentration of 10 mM. The biotinylated β2M was dialyzed against 6 M guanidinium HCl to remove any free biotin and was subsequently used in refolding reactions. A dilution method of refolding was used to produce specific H-2D<sup>b</sup>-peptide complexes (29). In brief, denatured biotinylated human β2M and H-2D<sup>b</sup> heavy chain (>1 mg/ml in 8 M urea) and peptide were diluted into refolding buffer (0.4 M l-arginine, 0.1 M Tris, pH 7.5, 2 mM EDTA, 5 mM reduced glutathione, 0.5 mM oxidized glutathione, and 0.5 mM PMSF) to a final concentration of 0.5 mM PMSF) to a final concentration of 10 mM. The biotinylated β2M was dialyzed against 6 M guanidinium HCl to remove any free biotin and was subsequently used in refolding reactions. A dilution method of refolding was used to produce specific H-2D<sup>b</sup>-peptide complexes (29). In brief, denatured biotinylated human β2M and H-2D<sup>b</sup> heavy chain (>1 mg/ml in 8 M urea) and peptide were diluted into refolding buffer (0.4 M l-arginine, 0.1 M Tris, pH 7.5, 2 mM EDTA, 5 mM reduced glutathione, 0.5 mM oxidized glutathione, and 0.5 mM PMSF) to a final concentration of 0.76 μM heavy chain, 1.15 μM β2M, and 7.6 μM peptide. Refolding was carried out with stirring at 4°C for 36-48 h. The refolding solution was then concentrated using an Amicon (Millipore, Waltham, MA) stirred cell and centrifprep (MW cutoff: 10,000) and purified by gel filtration using a Superdex-75 column (Pharmacia, Piscataway, NJ) in 20 mM Tris-HCl (pH 7.5) and 150 mM NaCl on an FPLC system.

Construction of Tetrameric Class I–Peptide Complexes. Phycoerythrin-labeled neutravidin (Molecular Probes Inc., Eugene, OR) was mixed stepwise with biotinylated H-2D<sup>b</sup> complexes containing peptide GP33–41 at a 1:4 molar ratio. The tetrameric complexes were subsequently concentrated using a centrifprep (MW cutoff: 100,000) to 1 mg MHC class I complex/ml. Aliquots of 2 × 10⁶ cells were stained using 50 μl of Tetramer. Approximately 200,000 cells were analyzed by flow cytometry.

Flow Cytometry. Single cell suspensions were prepared from the spleens of LCMV-infected or uninfected control mice. 2 × 10⁶ cells were stained for 30 min with tricolor-conjugated anti-CD8 (Caltag Labs., South San Francisco, CA). After washing in PBS containing 2% FCS and 0.5 mM EDTA (PBS/FCS/EDTA), the cells were stained using 50 μl of FACS buffer, the cells were stained as described above using tricolor-conjugated anti-CD8 antibodies and tetrameric class I–peptide complexes. Subsequently, the cells were stained with FITC-conjugated anti–IFN-γ antibodies (Pharmacia, St. Albans, UK). The stained cells were washed twice in permeabilization buffer, resuspended in FACS buffer, and analyzed by flow cytometry. In all cases, staining and washing of the cells was carried out at 4°C.

Results and Discussion

T.CR Staining Using Tetrameric Class I–Peptide Complexes. GP33 tetramer staining of GP33-specific CD8<sup>+</sup> T cells was examined in spleen cells from a 318 TCR transgenic mouse. Approximately 40–60% of T cells in the 318 mouse express the transgenic TCR designated P14 (V<sub>a</sub>2V<sub>b</sub>8.1), which recognizes H-2D<sup>b</sup> in association with peptide GP33. 318 spleen cells were stained with anti-CD8 antibodies and with GP33 tetramers or antibodies specific for V<sub>a</sub>. As shown in Fig. 1, B and D, the percentage of CD8<sup>+</sup> cells that stained positive with GP33 tetramers correlated well with the percentages obtained using the transgenic TCR-specific antibody. In contrast, only 3.7% of CD8<sup>+</sup> cells recovered from a naive B6 mouse stained positive with GP33 tetramer (Fig. 1 C). GP33 tetramers were also used to stain polyclonal CD8<sup>+</sup> CTL lines that had been generated after peptide restimulation of spleen cells from LCMV-immune mice. These CTL lines had been maintained in culture for 2 mo

![Figure 1](image-url)  

**Figure 1.** Staining of GP33-specific H-2D<sup>b</sup>-restricted TCR transgenic T cells and polyclonal LCMV-specific CTL lines with GP33 tetramers. Spleen cells from a naive B6 mouse and a 318 TCR transgenic mouse were stained with monoclonal antibodies specific for CD8 and the TCR gene segment V<sub>a</sub>2 (A and B) or with anti-CD8 and GP33 tetramers (C and D). CD8<sup>+</sup> CTL lines specific for H-2D<sup>b</sup>-restricted LCMV peptide epitopes, GP33-41, NP396–404, and GP276–286 were stained using GP33 tetramers 2 wk after in vitro restimulation with peptide-pulsed spleen cells (E).
The results, shown in Fig. 2, indicate that ~24% of CD8+ cells recovered from CSF on day 7 after infection with LCMV-DO C I L E were positive for peptide GP33. As a negative control, B6 mice were infected intracranially with 2 × 10^5 PFU of rVVINDG. Although similar numbers of cells were recovered from the CSF of these mice, no significant staining of the cells was observed using the GP33 tetramers. These experiments demonstrate the specificity and potential usefulness of tetrameric class I–peptide complexes in the identification of antigen-specific T cells. Tetrameric class I complexes were used to stain GP33-specific T cells in the CSF of mice infected intracranially 7 d before with 30 PFU of LCMV-DO C I L E. Intracerebral infection with LCMV induces a lethal CD8+ T cell–dependent and perforin-mediated immunopathology (10). In B6 mice, the specific inflammatory response begins by days 5–6 after infection and reaches ~10^6 infiltrating cells/µl of CSF on days 6–7. The results shown in Fig. 2 B, indicate that ~24% of CD8+ cells recovered from CSF on day 7 after infection with LCMV-DO C I L E were positive for peptide GP33. As a negative control, B6 mice were infected intracranially with 2 × 10^5 PFU of rVVINDG. Although similar numbers of cells were recovered from the CSF of these mice, no significant staining of the cells was observed using the GP33 tetramers (Fig. 2 A).

Staining of GP33-specific CD8+ cells after intravenous inoculation of B6 mice with LCMV-DO C I L E. Spleen cells from mice infected intravenously with either a low dose (2 × 10^5 PFU) or a high dose (2 × 10^6 PFU) of LCMV-DO C I L E were stained using anti-CD8 antibodies and GP33 tetramers at days 3, 6, 9, and 15 after infection. After low-dose infection with this virus, GP33-specific CD8+ cells were barely detectable at day 6 after infection (Fig. 3 A), but had increased significantly in number by day 9 after infection to comprise ~1 in 3 CD8+ T cells. On day 9, viral titers were no longer detectable in the spleens of the infected mice (Fig. 4). After infection with a high dose of LCMV-DO C I L E, GP33-specific CD8+ cells comprised almost 1 in 2 CD8+ T cells at day 3 after infection, but had already decreased in number by day 6 after infection and were barely detectable at day 9 after infection (Fig. 3 B). In contrast to the situation observed after low-dose infection and despite the rapid induction of GP33-specific cells, these cells were not able to control the infection since virus titers continued to increase in the spleens of these mice after the time point at which GP33-specific cells were no longer detectable (Fig. 4; reference 30).

GP33-specific CTL responses in B6 mice infected intravenously with LCMV-DO C I L E. To establish if a correlation exists between the expansion of GP33-specific cells observed by flow cytometry and GP33-specific cytotoxicity, direct CTL assays were carried out using spleen cells from the mice described above. CTL activity mediated by antigen-specific CD8+ cells was usually difficult to measure at day 3 after infection since LCMV induces, at this early time point, strong NK cell activity that lyses both antigen-pulsed and unpulsed target cells. To circumvent this problem, direct CTL assays were carried out by adding a twofold excess of unlabeled NK-sensitive target cells (YAC-1 cells) in the presence of cold YAC-1 cells abrogated lysis of chromium-labeled MC57 target cells by NK cells from the spleens of mice that had been injected 48 h previously with poly-IC (data not shown). No inhibition of antigen-specific CTLs was observed when YAC-1 cells were included as cold-target competitors in a direct CTL assay using spleen cells from mice that had been infected 8 d before with LCMV (data not shown). Thus, antigen-specific CTL activity in spleen cells from mice infected 3 d previously with LCMV was measured in the presence of cold YAC-1 cells. Although only low levels of specific killing were observed using
spleen cells from mice infected with a low dose of LCMV-DOCILE, strong cytotoxic activity was evident in the spleens of mice that had been infected intravenously with a low dose (E–H) of LCMV-DOCILE. Spleen cells were tested for lysis of normal MC57 cells (○) or MC57 cells that had either been pulsed with peptide GP33 (▲) or which had been infected with LCMV-DOCILE (●). Virus titers, measured as PFU in the spleen (Spl) and thymus (Thy) of each mouse are shown in the upper right-hand corner of each panel. 200 PFU represents the detection limit of the assay.

Figure 4. Cytotoxic activity and virus titers in mice infected intravenously with LCMV-DOCILE. LCMV-specific CTL activity was measured using spleen cells from mice that had been infected intravenously with a low (A–D) or high dose (E–H) of LCMV-DOCILE. Spleen cells were tested for lysis of normal MC57 cells (○) or MC57 cells that had either been pulsed with peptide GP33 (▲) or which had been infected with LCMV-DOCILE (●). Virus titers, measured as PFU in the spleen (Spl) and thymus (Thy) of each mouse are shown in the upper right-hand corner of each panel. 200 PFU represents the detection limit of the assay.
IFN-γ. Spleen cells from naïve B6 mice and mice that had been infected intravenously 8 d previously with a low dose of LCMV-DOCILE were stimulated with PMA and ionomycin, permeabilized, and stained with anti–IFN-γ antibodies. Expression of IFN-γ was analyzed by flow cytometry. Fig. 5 shows that after LCMV infection, CD8+ T cells (C), including those that are GP33 specific (D), exhibit an elevated capacity to produce IFN-γ after stimulation with PMA and ionomycin when compared to stimulated (B) spleen cells recovered from naïve mice. Subsequently, CD8+ cells recovered from the spleens of mice that had been infected 3, 6, 9, or 15 d previously with either a low- or high-dose of LCMV-DOCILE were stained with GP33 tetramers and examined for their capacity to produce IFN-γ. The results, shown in Fig. 6, indicate that following both low- and high-dose infection with LCMV-DOCILE, the capacity of tetramer+/CD8+ cells to produce IFN-γ exhibits the same kinetics as cytotoxic activity (compare to Fig. 4).

Virus-specific CTL activity after intracranial inoculation with LCMV. Intracranial inoculation of immunocompetent wild-type mice with a low dose (30 PFU) of LCMV results in the induction of lethal choriomeningitis (31). The immunopathology observed in these mice correlates directly
showed that although relatively few CD8+ cells were detected in the spleen after intracranial infection with a high dose of LCMV-DOCILE. Intracranial inoculation of mice with low (30 PFU) and high (3 x 10^4 PFU) doses of LCMV-DOCILE provides an example of how the extent and kinetics of virus spread influences the kinetics of the immune response and therefore the outcome of infection. Immunopathologic disease appears to be limited in those mice infected with a high dose of LCMV-DOCILE because the overwhelming antigen burden causes functional exhaustion of effector CTLs. Functional exhaustion and subsequent deletion of effector CTLs proceeded more slowly after intracranial infection than intravenous infection. Although this correlates with the lower doses of LCMV that were used to infect mice intracranially rather than intravenously, it may also reflect the delayed induction of CTLs after infection via a peripheral (intracranial; reference 35) rather than a systemic (intravenous) route.

Production of IFN-γ by GP33-specific T cells after Intracranial Infection of B6 Mice with LCMV-DOCILE. IFN-γ has been shown to have a direct antiviral effect after intracranial infection of mice with VV (4). In addition, exposure to IFN-γ has been shown to increase expression of MHC class I antigens on the surface of brain cells such as astrocytes, oligodendrocytes, microglia, and neurons (36). As shown in Fig. 9, CD8+ cells identified in the spleens (9, E and F) and CSF (9, A and B) of mice infected intracranially with a low dose (30 PFU) of LCMV-DOCILE, including those that are peptide GP33 specific, exhibited a high capacity to produce IFN-γ. A smaller proportion of cells recovered from both the CSF (9, C and D) and spleens (9, G and H) of mice that had been infected with a higher dose (3 x 10^4 PFU) of LCMV-DOCILE-produced IFN-γ. This correlated with the increasingly anergic status of the cells after exposure to excessive antigen.

Staining of GP33-specific Cells after Intravenous Infection of Perforin-deficient Mice with LCMV-DOCILE. A recent study by Sad et al. showed that GP33-specific T cell lines established by in vitro restimulation of spleen cells isolated from LCMV-infected perforin-deficient mice (PKOB) produced more IFN-γ than similar cell lines established from infected B6 mice (37). IFN-γ was measured in the supernatants of the cultures by ELISA. In this study, spleen cells from PKOB mice were isolated on days 9 and 15 after intravenous infection with a low dose (200 PFU) of LCMV-DOCILE. The cells were stained with anti-CD8 antibodies, GP33 tetramers, and anti-IFN-γ antibodies as described above. The results, shown in Fig. 10, A–C, indicate that the expansion of GP33-specific T cells appears to be similar at day 9 to that observed in B6 mice infected in the same way. Both the GP33-specific cell population and the remainder of the CD8+ cells isolated from PKOB mice did not, however, exhibit an elevated capacity to produce IFN-γ when compared to B6 mice (Fig. 11), despite the finding that virus titers were very high in PKOB mice (10^5 PFU LCMV/spleen) at a time point (day 9) when virus had already been cleared from the spleens of wild-type mice (Fig. 4). A similar analysis performed at day 15 revealed that the proportion of GP33-specific CD8+ cells had further increased in

with the cytotoxic action of virus-specific CD8+ T cells since although mice lacking the perforin gene remain persistently infected with LCMV, they do not die of disease (10). As shown in Fig. 7, A, B, E, and F, GP33-specific CD8+ T cells were readily detectable in the spleen and CSF of mice 7 days after infection with a low dose of LCMV-DOCILE. Direct CTL assays carried out using spleen cells from these mice show that although these cells exhibit strong cytotoxic activity (Fig. 8 A), the infection is not cleared; the mice develop encephalomyelitis and usually die within the next 48 h.

Mice infected intracranially with a 1,000 times higher dose (3 x 10^4 PFU) of LCMV-DOCILE also show signs of disease between days 7 and 9 after infection. They, however, recover from transient disease and survive the infection, even though they remain persistently infected with LCMV. This observation has been attributed to the exhaustion of virus-specific CTLs caused by the presence of excessive antigen. In keeping with this hypothesis, staining of lymphocytes isolated from the spleen and CSF of mice 7 days after intracranial infection with a high dose of LCMV showed that although relatively few CD8+ cells stained GP33 tetramer positive (Fig. 7, D and H), TCR expression was reduced when compared to cells isolated from the CSF of mice infected with a low dose of the same virus (compare mean fluorescence, parentheses, Fig. 7, F and H). Downregulation of TCR expression, which has been shown to occur after sustained contact between T cells and their corresponding antigen (32–34), correlated with very little cytotoxic activity in the spleen cells of mice either at day 7 (Fig. 8 B) or day 12 (data not shown) after intracranial infection with a high dose (3 x 10^4 PFU) of LCMV-DOCILE. Intracranial inoculation of mice with low (30 PFU) and high (3 x 10^4 PFU) doses of LCMV-DOCILE provides an example of how the extent and kinetics of virus spread influences the kinetics of the immune response and therefore the outcome of infection. Immunopathologic disease appears to be limited in those mice infected with a high dose of LCMV-DOCILE because the overwhelming antigen burden causes functional exhaustion of effector CTLs. Functional exhaustion and subsequent deletion of effector CTLs proceeded more slowly after intracranial infection than intravenous infection. Although this correlates with the lower doses of LCMV that were used to infect mice intracranially rather than intravenously, it may also reflect the delayed induction of CTLs after infection via a peripheral (intracranial; reference 35) rather than a systemic (intravenous) route.
PKOB mice (Fig. 10 E). These cells showed a reduced capacity to produce IFN-γ in response to stimulation with PMA and ionomycin (Fig. 11, compare A to C and B to D); the mice remained persistently infected with virus (3 x 10^6 PFU LCMV/spleen). In a similar fashion to CD8+ cells recovered from mice that had been infected intracranially with a high dose of LCMV-DOCILE, virus-specific CD8+ cells recovered from PKOB mice 15 d after LCMV infec-

**Figure 9.** Intracellular cytokine staining of CD8+ cells isolated from the spleens and CSF of mice infected intracranially with a low or high dose of LCMV-DOCILE. Spleen cells and CSF recovered from mice that had been infected intracranially with either low (30 PFU) or high (3 x 10^4 PFU) dose LCMV-DOCILE were stimulated with PMA and ionomycin and subsequently stained with anti-CD8 antibodies, GP33 tetramers, and anti-IFN-γ antibodies. Subsequent FACS analysis was carried out as described for Fig. 5. (Left) CD8+ cells that express IFN-γ; (right) CD8+ GP33-specific T cells that express IFN-γ. The percentages of CD8+ and CD8+/tetramer+ cells (from the experiment shown in Fig. 7) that express IFN-γ are shown in each panel.

**Figure 10.** Staining of GP33-specific D8-restricted CD8+ cells isolated from the spleens of B6 and PKOB mice infected intravenously with LCMV-DOCILE. Cells isolated from the spleens of either B6 mice infected with a low dose of LCMV-DOCILE (A and D), PKOB mice infected with a low dose of LCMV-DOCILE (B and E), or B6 mice infected with a high dose of the same virus (C and F) were stained with anti-CD8 antibodies and GP33 tetramers (histograms). The gated CD8+ cell population stained with GP33 tetramers. The results are representative of two independent experiments carried out using groups of two mice.
tion also showed a reduced capacity to produce IFN-\(\gamma\). Virus-specific PKOB CD8\(^+\) T cells were not, however, deleted, as was observed previously in B6 mice infected intravenously with a high dose of the same virus (Fig. 10, compare B to C and E to F).

This study shows that high frequencies of CTLs are stimulated in B6 mice during the acute phase of infection with LCMV. Frequencies of virus-specific CTLs correlate directly with the extent of early infection. Despite the induction of higher frequencies of CTLs after intravenous infection with high-dose rather than low-dose infection with LCMV-DOCLE, these CTLs fail to control the virus and are subsequently deleted (30). Studies on mice infected intracranially with a high dose of LCMV-DOCLE, where the induction of virus-specific CTLs occurs in a staggered fashion and is less rapid than after intravenous infection, indicate that an anergic phase exists between CTL induction and deletion. During this phase, virus-specific CD8\(^+\) cells may be described as functionally exhausted since they are characterized by a lack of cytotoxic activity and a reduced capacity to produce IFN-\(\gamma\). Although the PKOB virus-specific CD8\(^+\) cells also showed a reduced capacity to produce IFN-\(\gamma\) after prolonged exposure to antigen, these cells were not deleted. These data indicate that disappearance of virus-specific CD8\(^+\) T cells correlates with sustained perforin-mediated cytotoxic activity. CTL in perforin-competent mice may die as a result of interleukin starvation after CTL-mediated destruction of LCMV-infected, cytokine-producing APCs. Alternatively, deletion of CTLs in perforin-competent mice may result directly from perforin-dependent activation-induced apoptosis. Both possibilities remain to be evaluated.

This study further demonstrates that tetrameric class I–peptide complexes provide novel opportunities for the detection of antigen-specific T cells. The technique used in this study differs from that described by Altman et al. (11) in that it uses, instead of enzymatic biotinylation to the COOH terminus of the class I heavy chain, chemical biotinylation of the \(\beta_2M\) subunit. This modification renders the technique versatile since the final product, biotinylated \(\beta_2M\), can be used to refold any mouse or human class I heavy chain. Use of tetrameric class I–peptide complexes has an advantage over the use of anti-TCR antibodies in that they allow phenotypic characterization of all T cell clones of a given peptide-specificity. In addition, they provide the opportunity to study the phenotype of antigen-specific T cells without prior in vitro manipulation and without the need for transgenic animals.

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