

ticipate in the autoimmune process. Potentially, neuronal OPN secretion could modulate inflammation and demyelination and could influence the clinical severity of the disease. Consistent with this idea, a role for neurons in the pathophysiology of MS and EAE has recently been described (21, 22), and neurons are known to be capable of cytokine production (35, 36). OPN inhibits cell lysis (6), and thus, neuronal OPN might even protect the axon from degeneration during autoimmune demyelination.

CD44 is a known ligand of OPN, mediating a decrease of IL-10 production (10). As shown here, OPN^{-/-} mice produced elevated IL-10 during the course of EAE. We recently demonstrated that antibodies against CD44 prevented EAE (37), suggesting that the proinflammatory effect of OPN in MS and EAE might be mediated by CD44. The binding of OPN to its integrin fibronectin receptor $\alpha_V \beta_3$ through the arginine-glycine-aspartate tripeptide motif may also perpetuate T_H1 inflammation (10). In active MS lesions, the α_V subunit of this receptor is overexpressed in macrophages and endothelial cells, and the β_3 subunit is expressed on endothelial cell luminal surfaces (23). By means of its tripeptide-binding motif, OPN inhibits inducible nitric oxide synthetase (iNOs) (38), which is known to participate in autoimmune demyelination (7). Thus, in conclusion, OPN is situated at a number of checkpoints that would allow diverse activities in the course of autoimmune-mediated demyelination.

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detection of rare RNA species, we produced nonnormalized libraries, where manipulation of clones is avoided. White matter brain tissue from the plaques of three MS patients was collected and frozen within 2 hours after death. Patient history on the specimen used for the first library (herein MS1) included clinically definite MS and the presence of active inflammatory lesions. Material for the second MS library (herein MS2) came from a pool of tissues from two patients, one with acute, active lesions and widespread inflammatory involvement in the white matter, and the other with chronic, "silent" lesions, with gliosis, but without evidence of a lymphocytic infiltrate. The control library (CTRL) was constructed using pooled mRNA isolated from midbrain white matter, inferior temporal cortex, medulla, and posterior parietal cortex tissue removed from a 35-year-old Caucasian male who died from cardiac failure and who had no neuropathological changes. Details of construction of the libraries can be found as supplementary material on Science Online, particularly in the legends to Web fig. 1 and Web table 1 (14).

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the cellular staining was evaluated by an observer blinded to the experimental design according to a semiquantitative scale (three grades). MPIIB10₁ stains OPN in immunohistochemical sections from mice, although it does not recognize OPN on Western blots (26). The successful use of MPIIB10₁ in mouse sections has been reported (27, 28).

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30 May 2001; accepted 15 October 2001

Priming of Memory But Not Effector CD8 T Cells by a Killed Bacterial Vaccine

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Killed or inactivated vaccines targeting intracellular bacterial and protozoal pathogens are notoriously ineffective at generating protective immunity. For example, vaccination with heat-killed *Listeria monocytogenes* (HKLM) is not protective, although infection with live *L. monocytogenes* induces long-lived, CD8 T cell-mediated immunity. We demonstrate that HKLM immunization primes memory CD8 T lymphocyte populations that, although substantial in size, are ineffective at providing protection from subsequent *L. monocytogenes* infection. In contrast to live infection, which elicits large numbers of effector CD8 T cells, HKLM immunization primes T lymphocytes that do not acquire effector functions. Our studies show that it is possible to dissociate T cell-dependent protective immunity from memory T cell expansion, and that generation of effector T cells may be necessary for long-term protective immunity.

CD8 T lymphocytes mediate immunity to a broad range of viral, bacterial, and protozoal pathogens (1), and increasing evidence suggests that effector T cells primed during infection evolve into long-lived memory cells

(2–4). Memory T cells can be subdivided into two categories on the basis of activation markers, homing receptor expression, and effector function (5). Central memory T cells, which express high levels of the chemokine

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receptor CCR7 and the adhesion molecule CD62L and do not express effector functions, may differentiate into effector mem-

ory T cells, which express low levels of CCR7 and CD62L but produce cytokines (6). Whether these memory T cell subsets

differ in their capacity to mediate protective immunity is unknown. Furthermore, the stimuli that generate central versus ef-

Fig. 1. HKLM immunization primes antigen-specific memory CD8⁺ T cell responses. **(A)** CB6 mice (6 to 8 weeks old) were immunized intravenously with PBS, 10⁴ live *L. monocytogenes*, or two consecutive daily doses of 10⁹ HKLM (13). Mice were infected 21 days later with 10⁴ (naïve and HKLM-immunized mice) or 10⁵ (immune mice) live *L. monocytogenes*. Splenocytes, 7 days after infection in PBS-treated mice and 5 days after infection in HKLM and live *L. monocytogenes*-immunized mice, were stained for CD8 α and CD62L and with H2-K^d tetramers complexed with three *L. monocytogenes*-derived epitopes (LLO₉₁₋₉₉, p60₂₁₇₋₂₂₅, and p60₄₄₉₋₄₅₇). Dot plots are gated on live CD8 T lymphocytes and show CD62L and H2-K^d tetramer staining. The percentage of activated tetramer-positive CD8 T cells is shown in the upper left quadrant of each panel. **(B)** CB6 mice were immunized with two intravenous doses of 10⁹ HKLM, 10⁴ live *L. monocytogenes*, or PBS and were challenged 21 days later with 10⁴ live *L. monocytogenes*. Protective immunity was measured by dissociating spleens 72 hours after infection and quantifying viable bacteria. Mean numbers of colony-forming units (CFU) from six mice per group are shown (error bars, SD). **(C)** BALB/c mice were immunized with 2000 live *L. monocytogenes* and, 4 weeks later, left untreated or depleted of CD8 T cells by intravenous administration of three consecutive daily doses of 100 μ g of CD8 α -specific mAb (anti Lyt 2, 53-6.72, American Type Culture Collection) before rechallenge with 10⁵ live *L. monocytogenes*. Mice received an additional dose of CD8 mAb 3 days after infection. CD8 T cell depletion was greater than 90%. Mean numbers of bacteria in spleens (three mice per group) 72 hours after infection are shown (error bars, SD).

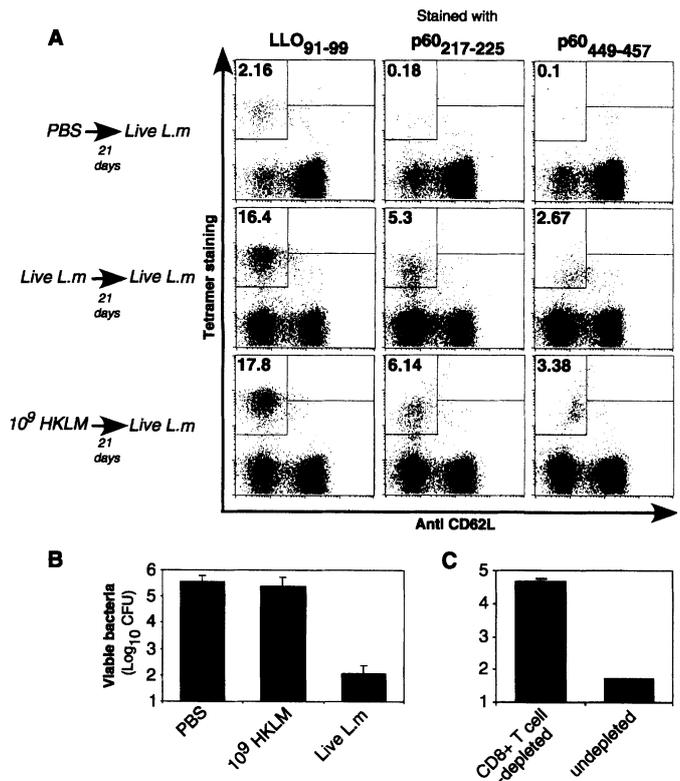
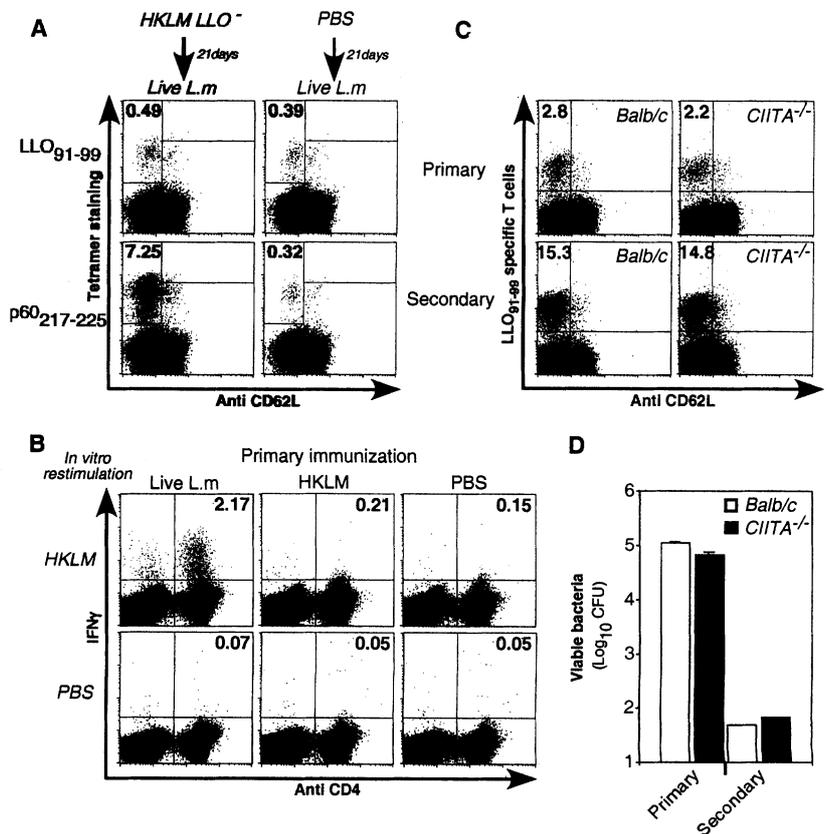


Fig. 2. HKLM directly primes antigen-specific CD8⁺ T cells, but not *L. monocytogenes*-specific CD4⁺ T cells. **(A)** CB6 mice (three per group) were injected intravenously with PBS or HKLM derived from a mutant *L. monocytogenes* strain lacking the LLO₉₁₋₉₉ epitope (see text) and were rechallenged 21 days later with live *L. monocytogenes*. The frequencies of CD8 T cells specific for LLO₉₁₋₉₉ and p60₂₁₇₋₂₂₅ were determined by tetramer staining 5 days after infection. **(B)** Mice were immunized with live *L. monocytogenes*, HKLM, or PBS; 6 days later, splenocytes were restimulated in vitro with HKLM (top row) or PBS (bottom row) in the presence of brefeldin A (BFA) for 6 hours. IFN- γ production was measured by intracellular cytokine staining using manufacturer's protocols (Cytotfix/Cytoperm, Pharmingen). Dot plots show CD4 and IFN- γ staining; percentages of CD4 T cells producing IFN- γ are shown. **(C)** BALB/c or CIITA^{-/-} mice (8 to 10 weeks old) were infected intravenously with live *L. monocytogenes* and analyzed 7 days later for the frequency of LLO₉₁₋₉₉-specific T cells by tetramer staining (top row). Mice that had received live *L. monocytogenes* 21 days earlier were rechallenged with 10⁵ live bacteria, and the LLO₉₁₋₉₉-specific CD8 T cell response was measured 5 days later. **(D)** The number of live bacteria in the spleens of mice 72 hours after primary or secondary infection was determined as described in Fig. 1. Means of two mice per group are shown (error bars, SD). In (A) and (C), the percentage of activated, tetramer-positive CD8 cells is shown in the upper left quadrant of each panel.



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factor memory T cells remain undefined.

Listeria monocytogenes is a Gram-positive, facultative intracellular bacterium that causes severe disease in immunocompromised patients (7). Studies in a mouse model of listeriosis have demonstrated that CD8 T cells mediate protective immunity after immunization with live bacteria (8–10). HKLM immunization, on the other hand, does not induce protective immunity (11). Infection of

mice with live *L. monocytogenes* induces antigen-specific CD8 T cell responses, with peak frequencies 7 to 9 days after infection of naïve mice and 5 days after reinfection of immune mice (12). To determine whether HKLM immunization primes *L. monocytogenes*-specific CD8 T cells, we immunized CB6 (C57BL/6 × BALB/c F₁) mice with live *L. monocytogenes* or HKLM (13) and, 21 days later, intravenously infected these mice with live *L. monocytogenes*. The magnitude of the *L. monocytogenes*-specific CD8 T cell response was measured by H2-K^d tetramer staining of splenocytes (Fig. 1A). CD8 T cells specific for the immunodominant listeriolysin epitope LLO₉₁₋₉₉ were detectable 7 days after infection of naïve mice (Fig. 1A, upper left panel), whereas responses to the subdominant p60₂₁₇₋₂₂₅ and p60₄₄₉₋₄₅₇ epitopes were smaller. As expected, reinfection of mice immunized with live *L. monocytogenes* induced markedly larger CD8 T cell memory responses (Fig. 1A, middle row), and these mice were resistant to rein-

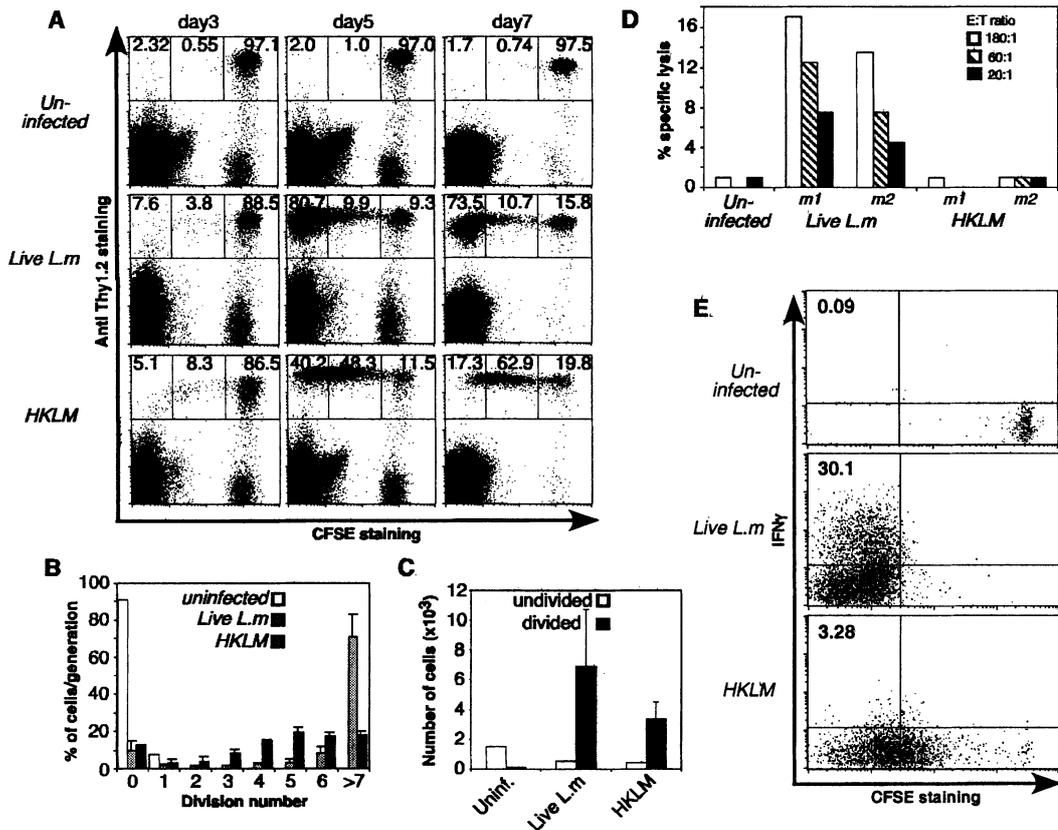
fection (Fig. 1B). In contrast, HKLM immunization was not protective (Fig. 1B). Surprisingly, HKLM-immunized mice mounted memory CD8 T cell responses that were indistinguishable in size from those detected in immune mice (Fig. 1A, bottom row). Mice immunized with HKLM derived from an avirulent strain of *L. monocytogenes* lacking LLO also primed CD8 T cell responses to p60 (14), indicating that residual LLO associated with HKLM is not allowing access to the cytosol of APCs.

Although CD8 T cells can provide protective immunity to *L. monocytogenes* infection, CD4 T cells and antibodies have also been implicated in immunity (10, 15). To measure their contribution to protective immunity, we depleted CD8 T cells from immune mice and challenged them with *L. monocytogenes*. Depleted animals were markedly more susceptible than control immune mice to *L. monocytogenes* infection (Fig. 1C), demonstrating the essential role that CD8 T cells play in protective immunity. Depletion of CD8 T cells with a monoclonal antibody (mAb) spe-

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Fig. 3. HKLM-primed CD8⁺ T cells undergo limited primary expansion and do not acquire effector functions. (A) TCR-transgenic mice specific for p60₂₁₇₋₂₂₅ were generated with the TCR α and β chain genes from CTL clone L9.6 and backcrossed to the BALB/c (Thy1.2) background (19). L9.6 TCR-transgenic splenocytes (2.2×10^7) were labeled with 5 μ M CFSE and transferred into BALB/c Thy1.1 mice. One day later, mice were intravenously injected with PBS, live *L. monocytogenes*, or HKLM. Splenocytes were recovered 3, 5, and 7 days after immunization and stained with antibodies specific for CD8 α and Thy1.2. CFSE staining intensity and staining for Thy1.2, which identifies transferred T cells, are plotted for live CD8 T cells. Percentages of Thy1.2⁺ cells that are CFSE low, intermediate, and high are shown in each plot. (B) Percentages of cells that have divided from 0 to >7 times (as determined by the intensity of CFSE fluorescence) are plotted for uninfected, live *L. monocytogenes*-infected, and HKLM-immunized mice on the fifth day after inoculation. (C) Absolute numbers of transferred Thy1.2⁺ T cells that remained undivided or that proliferated in uninfected, live *L. monocytogenes*-infected, or HKLM-immunized mice were determined on the fifth day after inoculation. (D) For determination of the cytolytic activity of transferred L9.6 TCR-transgenic T cells, CD8 T cells were enriched with MACS anti-CD8 α beads (Ly-2, Miltenyi Biotech) from mice immunized 7 days previously with PBS, live *L. monocytogenes*, or HKLM. Direct cytolytic activity of these CD8 T cells was measured in a 6-hour ⁵¹Cr release assay using p60₂₁₇₋₂₂₅-pulsed P815 target cells. Percent specific lysis is plotted for three different effector-to-target ratios for an uninfected



cific for CD8 β yielded similar results (16).

The robust CD8 T cell memory responses in HKLM-immunized mice could have resulted from direct priming of antigen-specific CD8 T cells or, indirectly, through priming of *L. monocytogenes*-specific CD4 T cells that would act to accelerate naïve CD8 T cell responses upon challenge with live bacteria. To distinguish between these two mechanisms, we immunized mice with HKLM lacking the immunodominant LLO₉₁₋₉₉ CD8 T cell epitope (HKLM^{LLO-}). This strain of *L. monocytogenes* was generated by point mutation of an essential anchor residue of LLO₉₁₋₉₉ and expresses functional LLO (17).

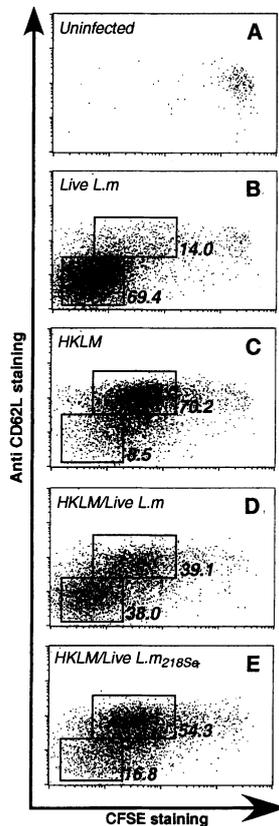


Fig. 4. Concurrent live infection with *L. monocytogenes* does not alter the phenotype of HKLM-primed CD8 T cells. L9.6 TCR-transgenic splenocytes (Thy1.1; 10⁶) were labeled with CFSE and transferred into Thy1.2 mice. One day later, mice were either left uninfected (A), infected with live wild-type *L. monocytogenes* (B), or immunized with HKLM alone (C) or together with live wild-type *L. monocytogenes* (D). One group of mice was immunized with HKLM and concurrently infected with live *L. monocytogenes* 218Ser (E), which contains a point mutation in p60, eliminating the p60₂₁₇₋₂₂₅ epitope. After 4.5 days, splenocytes were stained with mAbs specific for CD8, Thy1.1, and CD62L. Dot plots show CFSE and CD62L staining of gated Thy1.1 CD8 T cells. Each plot shows results for a single animal and is representative of two mice per experimental group. The percentage of transferred CD8 T cells in the CD62L^{hi} and CD62L^{lo} quadrants is shown.

If CD4 T cell responses accelerate CD8 T cell responses upon live challenge of HKLM-immunized mice, LLO₉₁₋₉₉-specific responses would be expected to still be enhanced in HKLM^{LLO-}-immunized mice. Infection of mice previously immunized with HKLM^{LLO-} resulted in a primary-like CD8 T cell response to LLO₉₁₋₉₉, similar to that obtained upon primary infection of naïve mice with *L. monocytogenes* (Fig. 2A, top row), but a memory-like CD8 T cell response to p60₂₁₇₋₂₂₅ (Fig. 2A, bottom row). This experiment demonstrates that HKLM immunization primed CD8 T cell responses independently of CD4 T cell priming. Indeed, CD4 T cells from HKLM-immunized and naïve mice, in contrast to mice infected with live bacteria, do not respond to bacterial antigen (Fig. 2B), which suggests that HKLM immunization does not efficiently prime CD4 T cells. Furthermore, infection of HKLM-immunized mice did not result in memory CD4 T cell responses (14).

One explanation for a lack of protective immunity in HKLM-immunized mice may be that deficient CD4 T cell responses limit *L. monocytogenes*-specific CD8 T cell function. To determine whether CD4 T cells influence CD8 T cell responses to *L. monocytogenes* infection, we measured the LLO₉₁₋₉₉-specific CD8 T cell response in mice lacking the class II transactivator gene (CIITA^{-/-}). These mice have a 95% reduction in peripheral CD4 T cells and do not mount peripheral CD4 T cell responses because B cells, dendritic cells, and macrophages lack major histocompatibility complex class II molecules (18). The primary CD8 T cell response to *L. monocytogenes* infection was identical in CIITA-deficient and control mice (Fig. 2C). Similarly, reinfection of previously immunized CIITA^{-/-} and control mice elicited indistinguishable memory LLO₉₁₋₉₉-specific CD8 T cell responses. CIITA-deficient mice cleared bacterial infection and developed similar protective immunity (Fig. 2D). Thus, CD4 T cell responses do not account for the difference in protective immunity after live and HKLM immunization.

To further characterize CD8 T cell priming by in vivo HKLM immunization, we generated a transgenic mouse line expressing T cells with specificity for p60₂₁₇₋₂₂₅ (19). Naïve, antigen-specific T cells from these mice were labeled with carboxyfluorescein diacetate succinimidyl ester (CFSE) and transferred into naïve recipient mice, allowing us to trace their fate after live infection or HKLM immunization. Donor CD8 T cells can be distinguished from recipient cells by a congenic Thy1 disparity. In the absence of infection, transferred p60₂₁₇₋₂₂₅-specific CD8 T cells remained CFSE positive (Fig. 3A, top row), indicating that they had not undergone in vivo divisions after transfer. In contrast, infection with live *L. monocytogenes* resulted in a large population of CFSE-low donor CD8 T cells (Fig. 3A, middle row).

Immunization with HKLM also induced the expansion of p60₂₁₇₋₂₂₅-specific T cells, with the majority of transferred T cells entering division (Fig. 3A, bottom row). However, in contrast to priming by live infection, p60₂₁₇₋₂₂₅-specific T cells primed with HKLM underwent fewer divisions (Fig. 3B). The proportions of naïve p60₂₁₇₋₂₂₅-specific T cells that proliferated after infection with live bacteria or immunization with HKLM were not significantly different (Fig. 3C), which suggests that transferred T cells encounter similar levels of processed antigen. It is unlikely that decreased T cell proliferation reflects insufficient antigen presentation, because mice immunized with 10⁸, 10⁹, and 10¹⁰ HKLM yielded similar results. Furthermore, we and others have demonstrated that naïve CD8 T cells require only brief encounter with antigen to undergo prolonged division (20–22). Thus, it is unlikely that a shorter duration of in vivo antigen presentation after HKLM immunization accounts for decreased proliferation.

To determine whether *L. monocytogenes*-specific CD8 T cells activated by HKLM immunization express cytolytic activity, we transferred p60₂₁₇₋₂₂₅-specific T cells and infected recipients with live *L. monocytogenes* or immunized with HKLM. Seven days later, CD8 T cells were assayed for cytolytic activity. CD8 T cells from infected recipients were cytolytic, whereas those from HKLM-immunized recipients displayed no lytic activity (Fig. 3D). Because the ratios of antigen-specific T cells to target cells were similar, these results demonstrate that CD8 T cells induced to proliferate by HKLM do not acquire cytolytic activity. Similarly, the proportion of p60₂₁₇₋₂₂₅-specific T cells producing interferon- γ (IFN- γ) after HKLM immunization is reduced relative to live infection (Fig. 3E). These experiments show that HKLM immunization supports CD8 T cell expansion, but not differentiation into effector CD8 T cells.

Consistent with their disparate effector functions, p60₂₁₇₋₂₂₅-specific CD8 T cells after live immunization became CD62L^{lo}, whereas HKLM-primed T cells remained CD62L^{hi} (Fig. 4, B and C). To determine the possible impact of CD4 T cell help and/or inflammation on CD8 T cell priming, we coimmunized mice with HKLM and live bacteria. Concurrent immunization of mice with HKLM and live wild-type bacteria resulted in two distinct populations, one CD62L^{hi} and the other CD62L^{lo}, suggesting that some T cells are primed by live infection while others follow the HKLM route (Fig. 4D). Concurrent immunization with wild-type HKLM and live bacteria lacking the p60₂₁₇₋₂₂₅ epitope primed T cells that underwent fewer divisions and remained CD62L^{hi} (Fig. 4E), demonstrating that CD4 T cell responses and in-

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flammation do not promote differentiation of HKLM-primed CD8 T cells. These results suggest that different antigen-presenting cells, perhaps in different sites, prime naïve CD8 T cells after HKLM and live infection.

Innate inflammatory responses to lipopolysaccharide enhance proliferation and survival of antigen-stimulated CD4 T cells (23), an effect that may result from the induction of antiapoptotic cellular factors (24). Although inflammatory responses to live infection and HKLM immunization differ, HKLM consists of a remarkable constellation of adjuvants: lipoteichoic acid, peptidoglycan, flagellin, lipoproteins, and bacterial DNA. Nevertheless, stimulation of CD40 during HKLM immunization augments the magnitude of *L. monocytogenes*-specific immune responses (25). It is possible that HKLM-derived molecules do not access relevant innate immune receptors, perhaps because killed and live bacteria are cleared by different cells. For example, HKLM uptake may be restricted to macrophages, whereas live infection may provide greater access of antigen to dendritic cells.

Recent studies have demonstrated that memory T cells reside predominantly in peripheral tissues (26, 27) and that central and effector memory cells differ with respect to trafficking (6). Previous studies have also shown that adoptively transferred CD8 T cells only limit bacterial growth if administered during the first day of infection (28). It is possible, therefore, that HKLM-primed memory CD8 T cells traffic and/or acquire effector functions too sluggishly to mediate protective immunity. Our findings suggest that differentiation of CD8 T cells into effec-

tor cells during primary immune responses has important consequences for the development of protective immunity. We believe this has important implications for pathogen- and cancer-specific vaccine development.

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AACTATGCTGACTCTCTCTCTGCC-3' and the antisense J β 2.4 primer 5'-TCTTATCGATACAGC-CCCAGAACCAAC-3'. The resulting fragment was digested with Xho I and Cla I and cloned into the TCR β cassette (30). The TCR α chain VJ region was amplified using the sense V α 4.2 primer 5'-CGCCCGGGCAGAAAATGAACCTTTGTCCTGAACTGGGG-3' and the antisense J α 4 primer 5'-CGCGCGGC-CGCGCTACCACCAAGCCTGTGG-3', which include Xma I and Not I restriction sites, respectively. This fragment was digested with Xma I and Not I and cloned into the TCR α cassette (30). Plasmids were screened by restriction mapping for presence of the complete TCR β or α gene. The TCR β gene insert was separated from the prokaryotic fragment of the vector by digestion with Kpn I and the TCR α gene insert by digestion with Sal I. These DNA fragments were purified and comicroinjected into fertilized oocytes of C57BL6 \times SJL F₁ mice at the Yale transgenic mouse facility. Founders were identified by PCR screening of DNA purified from mouse tails, and were bred for 10 generations onto the BALB/c background.

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31. We thank R. Flavell for providing the CIITA knockout mouse, C. Viret and D. Sant Angelo for excellent technical advice and discussions on the generation of TCR-transgenic mice, and C. Surh (Scripps Research Institute) for providing us with Thy1.1-congenic BALB/c mice. Supported by NIH grants AI 39031 and AI42135, a Human Frontier Science Program postdoctoral fellowship (G.L.), and National Research Service Award individual fellowship F32 AI09629-02 (S.V.).

19 July 2001; accepted 17 October 2001

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