Absence of Monocyte Chemoattractant Protein 1 in Mice Leads to Decreased Local Macrophage Recruitment and Antigen-specific T Helper Cell Type 1 Immune Response in Experimental Autoimmune Encephalomyelitis

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Abstract

Monocyte chemoattractant protein (MCP)-1 plays a critical role in innate immunity by directing the migration of monocytes into inflammatory sites. Recent data indicated a function for this chemokine in adaptive immunity as a regulator of T cell commitment to T helper cell type 2 (Th2) effector function. Studies in a Th1-dependent animal model, experimental autoimmune encephalomyelitis (EAE), showed that MCP-1 was highly expressed in the central nervous system (CNS) of affected rodents, and MCP-1 antibodies could block relapses of the disease. Mice deficient for the major MCP-1 receptor, CC chemokine receptor (CCR)2, did not develop EAE after active immunization but generated effector cells that could transfer the disease to naive wild-type recipients. We analyzed EAE in mice deficient for MCP-1 to define the relevant ligand for CCR2, which responds to murine MCP-1, MCP-2, MCP-3, and MCP-5. We found that C57BL/6 MCP-1–null mice were markedly resistant to EAE after active immunization, with drastically impaired recruitment of macrophages to the CNS, yet able to generate effector T cells that transferred severe disease to naive wild-type recipients. By contrast, adoptive transfer of primed T cells from wild-type mice into naive MCP-1–null recipients did not mediate clinical EAE. On the SJL background, disruption of the MCP-1 gene produced a milder EAE phenotype with diminished relapses that mimicked previous findings using anti–MCP-1 antibodies. There was no compensatory upregulation of MCP-2, MCP-3, or MCP-5 in MCP-1–null mice with EAE. These results indicated that MCP-1 is the major CCR2 ligand in mice with EAE, and provided an opportunity to define the role of MCP-1 in EAE. Compared with wild-type littermates, MCP-1–/– mice exhibited reduced expression of interferon γ in draining lymph node and CNS and increased antigen-specific immunoglobulin G1 antibody production. Taken together, these data demonstrate that MCP-1 is crucial for Th1 immune responses in EAE induction and that macrophage recruitment to the inflamed CNS target organ is required for primed T cells to execute a Th1 effector program in EAE.

Key words: autoimmune disease • chemokine • chemokine receptor • macrophage • T helper cell type 1/T helper cell type 2

Introduction

Chemokines are small proteins (8–12 kD) divided into four subfamilies (CXC, CC, C, and CX3C) according to the organization of positionally conserved cysteine residues (1). Monocyte chemoattractant protein 1 (MCP-1)1 is a prototype CC chemokine, active towards monocytes, dendritic cells, and NK cells, thereby playing an important role in innate immunity by directing the migration of monocytes into inflammatory sites. Recent data indicated a function for this chemokine in adaptive immunity as a regulator of T cell commitment to Th2 effector function. Studies in a Th1-dependent animal model, experimental autoimmune encephalomyelitis (EAE), showed that MCP-1 was highly expressed in the central nervous system (CNS) of affected rodents, and MCP-1 antibodies could block relapses of the disease. Mice deficient for the major MCP-1 receptor, CC chemokine receptor (CCR)2, did not develop EAE after active immunization but generated effector cells that could transfer the disease to naive wild-type recipients. We analyzed EAE in mice deficient for MCP-1 to define the relevant ligand for CCR2, which responds to murine MCP-1, MCP-2, MCP-3, and MCP-5. We found that C57BL/6 MCP-1–null mice were markedly resistant to EAE after active immunization, with drastically impaired recruitment of macrophages to the CNS, yet able to generate effector T cells that transferred severe disease to naive wild-type recipients. By contrast, adoptive transfer of primed T cells from wild-type mice into naive MCP-1–null recipients did not mediate clinical EAE. On the SJL background, disruption of the MCP-1 gene produced a milder EAE phenotype with diminished relapses that mimicked previous findings using anti–MCP-1 antibodies. There was no compensatory upregulation of MCP-2, MCP-3, or MCP-5 in MCP-1–null mice with EAE. These results indicated that MCP-1 is the major CCR2 ligand in mice with EAE, and provided an opportunity to define the role of MCP-1 in EAE. Compared with wild-type littermates, MCP-1–/– mice exhibited reduced expression of interferon γ in draining lymph node and CNS and increased antigen-specific immunoglobulin G1 antibody production. Taken together, these data demonstrate that MCP-1 is crucial for Th1 immune responses in EAE induction and that macrophage recruitment to the inflamed CNS target organ is required for primed T cells to execute a Th1 effector program in EAE.

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1 Abbreviations used in this paper: CCR, CC chemokine receptor; CNS, central nervous system; EAE, experimental autoimmune encephalomyelitis; IP-10, IFN-γ-inducible 10-kD protein; MCP, monocyte chemoattractant protein; MIP, macrophage inflammatory protein; MNC, mononuclear cell; MOG, myelin oligodendroglial glycoprotein; MS, multiple sclerosis; pi, postimmunization; PILN, popliteal and inguinal lymph node; PLP, proteolipid protein; RANTES, regulated upon activation, normal T cell expressed and secreted; RPA, RNase protection assay; RT, reverse transcription; TCA, T cell activation gene.
nate immunity (2–7). However, MCP-1 is also a crucial factor for the development of adaptive Th2 responses. In this regard, MCP-1 directs the differentiation of Th0 cells to Th2 in vitro (8) by a mechanism dependent on IL-4. Administration of anti–MCP-1 Abs (9) or disruption of the MCP-1 gene (2) significantly reduced the size of schistosome egg antigen (SEA) secondary granulomata, a Th2-dominant disease model. Conversely, local overexpression of MCP-1 increased the size of SEA secondary granulomata (10). Immunization with trinitrophenol-derivatized ovalbumin plus IFA elicited a reduced Th2 and unaltered Th1 response in MCP-1–deficient (MCP-1−/−) mice. MCP-1−/− mice of Balb/c strain were relatively resistant to *Leishmania major* infection, indicating that lack of MCP-1 led to reduced Th2 immunity (11).

Experimental autoimmune encephalomyelitis (EAE), a model for autoimmune demyelination of the central nervous system (CNS), has been widely employed to explore pathogenic mechanisms underlying the human disease multiple sclerosis (MS [12, 13]). The generation of myelin protein–reactive T cells is an immunological hallmark of both EAE and MS and is required for disease expression in EAE. These autoreactive T cells traffic to the CNS, and initiate inflammation and destruction of CNS myelin with consequent neurological impairment (14, 15). Th1-type T cells, producing IFN-γ, IL-2, and TNF-β, are associated with cellular immune responses, delayed-type hypersensitivity, and macrophage activation, whereas Th2-type T cells, producing IL-4, IL-5, and IL-10, are important for humoral immune responses (16, 17). The dynamic interplay and reciprocal inhibition between Th1 and Th2 cytokines has been demonstrated in numerous research reports. IL-4 is a major factor that governs Th2 differentiation and inhibits the development of IFN-γ–secreting cells (18). The activation of macrophages and the production of Th1 cytokines such as IFN-γ can also be inhibited by IL-10 (19). Most encephalitogenic T cell clones examined are Th1 polarized (20–22), although exceptions have been reported (23). Th1 cytokines are markedly elevated in the CNS of animals during EAE attacks whereas Th2 cytokines are associated with disease recovery (24). IL-4–induced immune deviation is beneficial for recovery from EAE (25); EAE can be prevented and/or reversed by myelin antigen–specific T cells that are genetically transduced with either IL-4 or IL-10 genes (26, 27); anti–IL-4 treatment reverses the tolerance induced by an altered peptide ligand (28), and absence of IL-4 in gene-targeted mice increases the severity of EAE (29). In summary, Th1 immune responses are pathogenic and Th2 responses are protective in the initiation and evolution of EAE.

However, antigen-specific T cells constitute only a small proportion of infiltrating leukocytes in EAE or MS lesions (30). Secondarily recruited inflammatory cells account for the vast majority of infiltrating cells and play a pivotal role in CNS tissue damage (31). Although the detailed mechanisms by which inflammatory cells influx into the CNS compartment are not completely understood, increasing evidence suggests that chemokines, in concert with adhesive molecules, are essential for this process (32). In EAE, elevated expression of MCP-1 by CNS parenchymal cells, tightly linked to clinical disease, has been demonstrated repeatedly (33–35). Further, anti–MCP-1 Abs blocked relapses of adoptive transfer EAE in SJL mice (36). Additionally, mice that lacked CC chemokine receptor 2 (CCR2), the major receptor on monocytes for MCP-1, failed to develop EAE after active immunization and were resistant to induction of EAE by the adoptive transfer of primed T cells from syngeneic wild-type mice (37, 38). It was uncertain whether MCP-1 was the relevant ligand for CCR2 in these experiments, as this receptor also responds to MCP-2, MCP-3, and MCP-5. However, there is also support for the possibility that regulation of Th2 responses by MCP-1 could be important for the pathogenesis of EAE; in particular, MCP-1 was critical for the development of tolerance after oral administration of a proteolipid protein (PLP) peptide containing residues 139–151 (39).

Therefore, the phenotype of EAE in MCP-1–deficient mice could not readily be predicted. On one hand, defective MCP-1–dependent monocyte recruitment might lead to attenuated disease. Alternatively, if functional replacement of MCP-1 by another MCP mediated monocyte accumulation in the CNS of these mice, defective Th2 responses might lead to very severe, nonremitting disease. Finally, in view of redundancy in the immune/inflammatory system, it remained possible that MCP–1–null mice could be important for the pathogenesis of EAE; in particular, MCP-1–null mice could be critical for the development of tolerance after oral administration of a proteolipid protein (PLP) peptide containing residues 139–151 (39).

In this report, we describe the phenotype of EAE in MCP-1–null mice. These mice exhibited markedly reduced clinical and histological EAE after active immunization and did not develop clinical disease after receiving encephalitogenic T cells from wild-type animals. Expressions of MCP-2, MCP-3, and MCP-5 in the CNS of both wild-type and MCP-1–null mice with EAE were virtually identical. These findings indicated that MCP–1 was the major ligand for CCR2 in murine EAE. In this EAE model, we found that disruption of the MCP-1 gene led to an attenuated Th1 autoimmune response and complimentary increased Th2 response. These results indicated a crucial role for MCP-1 in generating CNS inflammatory reactions that mediate the effector phase of myelin–specific Th1 autoimmune responses. Therefore, the data suggested that primed encephalitogenic Th1 cells cannot manifest effector functions in the CNS without recruiting hematogenous macrophages.
Materials and Methods

Mice. The disruption of the MCP-1 gene has been described previously (2). MCP-1−/− mice were backcrossed onto the C57BL/6 (B6) strain for eight generations. One F8 MCP-1−/− mouse was further backcrossed to a B6 mouse (obtained from The Jackson Laboratory). The heterozygous offspring were intercrossed to produce F9 wild-type (+/+), heterozygous (−/+), and MCP-1−/− mice. F10 mice were generated in a similar manner. MCP-1+/+, MCP-1−/+, and MCP-1−/− F9 and F10 mice on the B6 background were used in this study.

MCP-1−/deficient mice on the B6/129 background were also backcrossed onto SJL for seven generations. MCP-1−/+ and MCP-1−/− F7 mice on SJL background were used in this study.

Mice were genotyped using a PCR-based analysis of genomic DNA extracted from tail clips. Primers MCP-1F, 5′-GGA GCA TCC AGC TGT TGG C-3′ and MCP-1R, 5′-ACA GCT TCT TTT GTC GGG GAT GC-3′ amplified a fragment in the neomycin resistance gene insert. PCR reactions were performed in a PerkinElmer 9700 cycler (annealing temperature, 50°C) and products were visualized by electrophoresis on ethidium bromide–stained NuSieve GTE™ agarose gel.

Rat Myelin Oligodendroglial Glycoprotein and Mouse Proteolipid Protein Peptides. Rat myelin oligodendroglial glycoprotein (MOG)35–55 and mouse PLP139–151 peptides were obtained from BIO-SYNTHESIS and purified by HPLC with a purity of 98%. The sequence of MOG35–55 was MEVGYWRRSPS-RVVHLYRNGK and that of PLP139–151 was HSLGKWLGHPDKF.

Active Induction of EAE with MOG and PLP Peptides and Clinical Evaluation. Mice of 8–9 wk of age were subcutaneously injected with 300 μg MOG35–55 emulsified in CFA (Difco) containing 400 μg Mycobacterium tuberculosis. Mice were intravenously injected withpertussis toxin (Sigma-Aldrich) as indicated in the figure legends on day 0 and 2 postimmunization (pi). The immunization in SJL mice was carried out as described previously (43). All mice were weighed, examined, and graded daily for neurological signs in a double blind manner by one of us (J. Wang) as follows: 0, no disease; 1, decreased tail tone or slightly clumsy gait; 2, tail atony and moderately clumsy gait and/or poor righting ability; 3, limb weakness; 4, limb paralysis; and 5, moribund state. Disease relapse was determined when an increase of one EAE score unit was observed. Signs of neurological impairment were typically accompanied by an abrupt, substantial weight loss (>7%). The average day of EAE onset was calculated by adding the first day of clinical signs for individual mice and divided by the number of mice in the group. Day of EAE onset in mice that showed no clinical EAE was deliberately regarded as 1 d after the experiment was terminated (44). The EAE index was calculated by adding all the daily EAE scores to obtain cumulative score and dividing by day of EAE onset. Active immunization with MOG35–55 induced monophasic EAE in B6 mice and was followed for 65 d. Chronic relapsing EAE induced by PLP139–151 was monitored for 90 d.

T Cell Proliferation Assay. Mice were killed and draining lymph nodes (popliteal and inguinal lymph nodes [PILNs]) were dissected on day 10 pi. Single cell suspensions (5 × 10^6/ml) were prepared and cultured in triplicate in 96-well flat-bottomed plates (Falcon; Becton Dickinson) in 200 μl/well in the presence or absence of MOG35–55, PLP139–151, LPS, or anti-CD3 (R&D Systems) in RPMI 1640 (GIBCO BRL) supplemented with 2 mM l-glutamine, 100 U/ml penicillin, 100 μg/ml streptomycin, 10% FCS, and 5.5 × 10^−5 M 2-mercaptoethanol. Cells were pulsed with [3H]thymidine (Amersham Pharmacia Biotech), 0.5 μCi/well 72 h after culture initiation, and incubated 10 h further. Plates were harvested using a harvester (INOTECH). Incorporated radioactivity was measured in a MicroBeta PLUS liquid scintillation counter with software v3.3 (Wallac Co.).

Cell Cultures and Cytokine Assay. Mononuclear cell (MNC) suspensions (5 × 10^6/ml) were prepared from PLNIs of mice that had been immunized with MOG35–55 plus CFA for 10 d. Cells were cultured at 2 × 10^6/ml in RPMI 1640 supplemented with 10% FCS and 1% nonessential amino acids (NEA) and fed daily. Protected fragments were visualized and quantified by autoradiography with PhosphorImager (Molecular Dynamics).

Reversal Transcription PCR Detection for Levels of MCPs, Chemokines, and Cytokines. Mice were anesthetized with sodium pentobarbital and intracardially perfused through the left ventricle with ice-cold PBS. Spinal cords were extruded by flushing the vertebral canal with PBS, rinsed in PBS, and kept at −80°C. Total cellular RNA was prepared from spinal cord tissue by TRIzol (Life Technologies). Quantification of CC chemokines and chemokine receptors in CNS tissue was done by RNase protection assay (RPA) with Template Sets and In vitro Transcription Kit (BD PharMingen) according to the manufacturer’s instructions. Protected fragments were visualized and quantified by autoradiography with PhosphorImager (Molecular Dynamics).
and washed in PBS. rIL-12 (R&D Systems). After 4 d incubation, cells were collected in the presence of 300 μg/ml MOG35–55 plus 500 ng pertussis toxin intravenously injection on day 0 and 2 pi. One MCP-1–/– mouse died of immunization on day 4 pi. The remainder of the mice, regardless of genotype, developed clinical EAE. MCP-1–/– mice showed EAE signs around day 10 pi, consistent with previously reported results in C57BL/6 mice (44, 48). MCP-1–/– manifested significantly delayed EAE with an average onset on day 21 pi (Fig. 1). Three out of nine wild-type mice died of EAE, and another three had to be killed because of moribund state. None of the MCP-1–/– mice died of EAE or had to be killed throughout the experiment and they recovered from the disease significantly faster and more completely than wild-type littermate controls. Heterozygote MCP-1–/+ mice developed EAE with an intermediate kinetics and severity. None of the MCP-1–/– mice died of EAE, whereas three were killed in a moribund state due to severe EAE. Analyses of CNS tissue histology revealed massive inflammatory infiltrates in wild-type control mice (+++ to ++++ in regions of lumbar and sacral spinal cord, n = 4; Fig. 2 A) but markedly reduced inflammatory reaction in MCP-1–null mice (+ to ++ in regions of corresponding levels of affected spinal cord, n = 4; Fig. 2 B; reference 49). Examination of demyelination using Luxol Fast Blue stain-

**Results**

C57BL/6 MCP-1–deficient Mice Are Relatively Resistant to Active EAE Induction with MOG35–55. MCP-1+/+, MCP-1+/–, and MCP-1–/– F9 mice (n = 9 in each group) were immunized with 300 μg MOG35–55 plus 500 ng pertussis toxin intravenously injection on day 0 and 2 pi. One MCP-1–/– mouse died of immunization on day 4 pi. The remainder of the mice, regardless of genotype, developed clinical EAE. MCP-1–/– mice showed EAE signs around day 10 pi, consistent with previously reported results in C57BL/6 mice (44, 48). MCP-1–/– manifested significantly delayed EAE with an average onset on day 21 pi (Fig. 1). Three out of nine wild-type mice died of EAE, and another three had to be killed because of moribund state. None of the MCP-1–/– mice died of EAE or had to be killed throughout the experiment and they recovered from the disease significantly faster and more completely than wild-type littermate controls. Heterozygote MCP-1–/+ mice developed EAE with an intermediate kinetics and severity. None of the MCP-1–/– mice died of EAE, whereas three were killed in a moribund state due to severe EAE. Analyses of CNS tissue histology revealed massive inflammatory infiltrates in wild-type control mice (+++ to ++++ in regions of lumbar and sacral spinal cord, n = 4; Fig. 2 A) but markedly reduced inflammatory reaction in MCP-1–null mice (+ to ++ in regions of corresponding levels of affected spinal cord, n = 4; Fig. 2 B; reference 49). Examination of demyelination using Luxol Fast Blue stain-

**Figure 1.** Effect of MCP-1 gene disruption on MOG35–55-induced EAE. F9 MCP-1+/+, MCP-1+/–, and MCP-1–/– mice were immunized with MOG35–55 emulsified in CFA and intravenously injected with pertussis toxin (PT) on the day of immunization and 48 h later (500 ng/injection). Three wild-type mice died of EAE and another three had to be killed due to severe EAE attack. None of the MCP-1–/– mice died of EAE, but three were killed. None of the MCP-1–/– mice died of EAE or had to be killed. Shown are EAE score (mean ± SD) in each group of mice that had been followed throughout the experiment (n = 3, 6, and 8 for MCP-1+/+, MCP-1+/–, and MCP-1–/– group, respectively). This graph is representative of three experiments with similar results.

**Statistical Analyses.** The Instat 2.02 software was used for the analyses of the difference between the MCP-1+/+, MCP-1+/–, and MCP-1–/– mice. The Mann-Whitney U-test was used for the comparisons of disease severity and cytokine and chemokine gene expressions. A chi-square test was used for the comparisons of disease incidence in groups of MCP-1+/+, MCP-1+/–, and MCP-1–/– mice. A P value <0.05 was considered as significant.
ing revealed significant reduction in MCP-1−/− mice compared with their littermates controls 65 d after immunization (data not shown).

F10 MCP-1−/− mice and wild-type littermate controls were analyzed in a subsequent experiment. As pertussis toxin has been shown to increase the permeability of the blood-brain barrier (BBB [50, 51]), enhance delayed type hypersensitivity (DTH) responses and the production of IFN-γ (52–54), augment expression of CD80, CD86 on antigen-presenting cells and CD28 on T cells (55), and T cell immune responses (55, 56), we reduced the amount of pertussis toxin in the immunization to 200 ng per injection in an attempt to reduce the high death rate observed in wild-type mice in the F9 experiments. In this experiment, all MCP-1+/+ and MCP-1−/− mice showed signs of clinical EAE; none died of EAE or required killing. In contrast to the MCP-1+/+ mice that developed full-blown EAE, the disease was largely suppressed with significantly delayed onset and milder neurological impairment (Fig. 3) and significantly less weight loss (data not shown) in MCP-1−/− mice. These data demonstrated that MCP-1−null mice and CCR2-deficient mice exhibited strikingly similar EAE phenotypes (37, 38) and suggested that MCP-1 may be the relevant ligand for CCR2 in this model.

Disruption of MCP-1 Gene Attenuates the Severity of PLP-induced EAE and Reduces the Number of Relapses in SJL Mice. 11 MCP-1+/+ and 10 MCP-1−/− mice on SJL background were immunized with PLP139−151 peptide emulsified in CFA. One MCP-1+/+ and two MCP-1−/− mice died of immunization within the first week after immunization. The reminder of MCP-1+/+ mice (10/10) and 7 out of 8 MCP-1−/− mice showed clinical EAE. Compared with the MCP-1−/− group, a significantly higher percentage of MCP-1+/+ mice died of EAE during the first attack and relapses thereafter. Among the mice that survived the first attack, six MCP-1+/+ mice had eight relapses (four mice had one attack each and two had two attacks each), whereas three out of seven MCP-1−/− mice had single attacks. At day 90 pi, the average EAE index was significantly higher in MCP-1+/+ group (n = 4) than that in MCP-1−/− group (n = 7) (Fig. 4). These results implicated a role for MCP-1 in eliciting relapses of EAE in this model. These findings were consistent with previous reports showing reduction of relapses using anti-MCP-1 Abs in a passive EAE model (36) and in mice receiving vaccine containing naked DNA encoding for MCP-1 (57).

Figure 2. Spinal cord histology of MCP-1+/+ and MCP-1−/− mice with EAE score 4.0. Hematoxylin and eosin staining of longitudinal cryosections. Note the numerous perivascular cuffs and subpial infiltrates (arrows) as well as leukocytes disseminated in the white matter of MCP-1+/+ mice (A), whereas merely fewer inflammatory infiltrates were found in MCP-1−/− mice (B).

Figure 3. Attenuated MOG35−55−induced EAE in MCP-1−null mice. F10 MCP-1−/− mice (n = 6) and their littermate wild-type controls (n = 6) were immunized with MOG35−55 in CFA and intravenously injected with pertussis toxin (PT; 200 ng/injection). Shown are EAE score (mean ± SD) of individuals in each group.
creased expression of MCP-3 in MCP-1–null SJL mice and their littermate wild-type controls were immunized with PLP139–151 in CFA plus intravenous injection of pertussis toxin as described in Materials and Methods. Mice were monitored for 90 d after immunization. MCP-1–null SJL mice showed significantly decreased EAE index, reduced number of relapses, and nonsignificantly delayed disease onset.

Methods. Mice were monitored for 90 d after immunization. MCP-1 expression in wild-type controls was immunized with PLP139–151 in CFA plus intravenous injection of pertussis toxin as described in Materials and Methods. Mice were monitored for 90 d after immunization. MCP-1–null SJL mice showed significantly decreased EAE index, reduced number of relapses, and nonsignificantly delayed disease onset.

There Is No Compensatory Upregulation of MCP-2, MCP-3, or MCP-5 in the CNS of MCP-1–null Mice with EAE. Because CCR2 is shared in common among all MCPs (58–60), we analyzed MCP-2, MCP-3, and MCP-5 mRNA expression in the CNS of C57BL/6 mice with MOG-induced EAE, using quantitative real-time reverse transcription (RT)-PCR. Levels of MCP-2 and MCP-3 but not MCP-5 were elevated in CNS tissue from EAE mice. There was no significant difference in MCP-2 expression in CNS tissue between MCP-1+/+ and MCP-1–null group (23.8 ± 0.4 vs. 23.7 ± 0.5, mean ± SD; n = 5 in each group). No significant difference of MCP-3 expression in CNS tissue from MCP-1+/+ (31.0 ± 0.6, n = 5) and MCP-1–null (33.1 ± 2.6, n = 5, P = 0.4) mice was found, whereas CNS MCP-5 expression in mice with EAE was essentially undetectable (data not shown). The unaltered levels of MCP-2 and MCP-5 and a trend towards decreased expression of MCP-3 in MCP-1–null mice (higher PCR cycle number) uncovered no compensatory expression of other MCPs in this system, consistent with a previous report in autoimmune kidney disease model in MCP-1–null MRL-Faslpr mice (61). These findings supported the hypothesis that the similarity of EAE phenotype in MCP-1– and CCR2–deficient mice is caused by absence of the ligand, i.e., MCP-1 signaling pathway through its major receptor CCR2.

T Cell Proliferation to MOG35–55 Peptide. To examine the afferent limb of the immune response to MOG peptide in wild-type and MCP-1–null mice, MNC suspensions were prepared from PILNs primed with 200 µg MOG35–55 peptide in CFA and rechallenged with MOG35–55 in vitro. MCP-1–null mice showed a nonsignificantly higher recall T cell response than wild-type controls. No T cell recall response was induced by stimulation with PLP139–151 in vitro, a specificity control. No difference in anti-CD3e–induced T cell proliferation was found between wild-type and MCP-1–deficient mice (data not shown). These data indicated that the CD3 pathway was intact in MCP-1–null mice, and that MOG35–55–specific T cells can be generated in MCP-1–null mice.

MCP-1–null Mice Do Not Develop Clinical EAE in Passive Transfer Model. An adoptive transfer EAE model was used to further address whether MCP-1–null mice could develop pathogenic autoimmune responses to MOG35–55 peptide. MCP-1+/+ and MCP-1–null B6 mice were immunized with MOG35–55 peptide in CFA, and MNCs isolated from draining lymph nodes were cultured in the presence of MOG35–55 peptide and IL-12 before transfer into MCP-1–null mice or littermate controls. As expected, MCP-1+/+ mice receiving MCP-1+/+ T cells developed clinical EAE (Fig. 5). MCP-1–null T cells showed approximately the same encephalitogenic capacity in this adoptive transfer model, resulting in a comparable incidence, severity, and clinical course of EAE in MCP-1–null recipients. In contrast, MCP-1–null mice that received MCP-1+/+ T cells failed to develop clinical EAE (Fig. 5). This result indicated that the attenuated EAE in MCP-1–null mice was not caused by impaired generation of encephalitogenic T cells. The data also demonstrated that absence of MCP-1 expression in the recipient rendered the mice unable to respond to encephalitogenic signals produced by wild-type T cells.

To dissect the mechanisms underlying the relatively resistance to EAE induction in MCP-1–deficient mice, the MOG35–55–induced EAE model was used in the following mechanistic studies. To obtain samples from MCP-1–null mice with full-blown EAE, F10 mice were immunized with MOG35–55 in CFA plus 500 ng pertussis toxin per injection. Reduced CD11b+/CD4+/CD44+ Ratio but Unchanged Levels of CD3e and CD8 Transcripts in CNS Tissue from MCP-1–null Mice with EAE. Results described above suggested that MCP-1–null mice were deficient in recruiting mono-
cytes to the CNS during EAE. To address this issue, leukocytes were isolated from CNS tissue of wild-type and MCP-1--null mice with comparable severity of EAE, and analyzed with flow cytometry. Cell numbers in preparations isolated from MCP-1--/-- mice during EAE attacks (score 4) were ~1/3 of those from MCP-1+/+ littermate controls with comparable EAE severity. Compared with MCP-1+/+ littermate controls, MCP-1--/-- mice showed a sharply reduced percentage of CD11b+CD4+ cells in the CNS during EAE attacks (71.8 ± 4.6% vs. 44.0 ± 4.2%, mean ± SD, n = 5 and 4, respectively; P < 0.0001). In contrast, percentages of CD4+ cells were relatively increased in MCP-1--/-- mice (20.2 ± 5.9% vs. 39.8 ± 2.7%, mean ± SD, n = 6 and 4, respectively; P < 0.001; Fig. 6). The CD4+ infiltrating T cells expressed high levels of CD11b (αβ2 integrin) both in MCP-1+/+ (79.0 ± 5.6, n = 5) and MCP-1--/-- (78.1 ± 6.8, n = 5) mice, indicating that most infiltrating CD4+ T cells are activated.

To normalize percentage of CD4+ T cells in the CNS infiltrates of wild-type and MCP-1--null mice with EAE, total CNS T cells were analyzed by determining levels of CD3ε mRNA levels in wild-type (0.789, n = 5; CD3ε/GAPDH) and MCP-1--/-- (0.791, n = 5; CD3ε/GAPDH) mice with clinical EAE scores of 3.5–4.0, suggesting that total T cell numbers in the CNS of wild-type and MCP-1--null mice were equivalent. Therefore, the increased proportion of CD4+ cells in the CNS leukocyte infiltrates of MCP-1--null mice was caused by a marked reduction in the number of CD11b+CD4+ cells in the CNS of MCP-1--deficient mice with EAE.

CD8+ T cells play an important downregulatory role in the pathogenesis of EAE (62). The possibility that the milder clinical EAE phenotype observed in MCP-1--/-- mice might be due to an increased number of CD8+ T cells in the CNS infiltrates was unlikely based on the fact that levels of CD8-specific mRNA were virtually identical in MCP-1--/-- (24.08 ± 0.3, n = 5) and MCP-1+/+ (24.21 ± 0.2, n = 5) CNS tissue during EAE attack. The passive transfer EAE model was used to examine further if CD8+ T cells can be preferentially recruited into CNS in the absence of MCP-1. MOG35–55–reactive MCP-1+/+ T cells were incubated in the presence of MOG35–55 and IL-12, and were injected intravenously into MCP-1+/+ and MCP-1--/-- mice. To reduce the influence of secondarily recruited macrophages in the CNS, recipient mice were killed at day 3 and 4 after T cell transfer, before the onset of clinical EAE. CNS-infiltrating T cells were recovered and analyzed using flow cytometry. No difference was found between MCP-1+/+ and MCP-1--/-- mice (data not shown). These results demonstrated that disruption of MCP-1 gene exerts no significant impact on the recruitment of adoptively transferred T cells into inflammatory CNS tissue. No significant difference in Mb-1 mRNA levels was found in EAE-affected CNS tissue from MCP-1+/+ and MCP-1--/-- mice (data not shown).

We also examined the leukocyte infiltrates in MCP-1--/-- mice at day 14 pi when the MCP-1--/-- controls were undergoing EAE attacks while the MCP-1--/-- mice were still free of EAE signs. Shown in Fig. 6 C are numerous CD45hiCD11b+ (mainly containing macrophages/activated microglia and activated T cells) isolated from MCP-1+/+ mice during EAE attack on day 14 pi. In contrast, the majority of cells isolated from MCP-1--/-- mice were CD45lowCD11b+ microglia, and the components of the infiltrates (Fig. 6 D) were virtually the same as those from healthy unimmunized mice (data not shown).

**Figure 6.** Altered pattern of CNS cell infiltrates in MCP-1--null mice. MCP-1--/-- and MCP-1+/+ littermate controls were immunized with MOG35–55 and pertussis toxin (500 ng/injection) and killed at the peak of EAE (score 4). Cells were isolated from the CNS and stained with anti-CD4-FITC, anti-CD11b-PE, and anti-CD45-Cy mAbs. Compared with wild-type controls (A), the percentages of CD11b+CD4+ cells were significantly decreased whereas the percentages of CD4+ T cells increased in MCP-1--/-- mice (B). In contrast to MCP-1+/+ mice that developed full-blown EAE with numerous CNS CD11b+CD45hi microglia/macrophages/activated microglia and T cells (C), the majority of cells isolated from CNS tissues of EAE symmetric-free MCP-1--/-- mice on day 14 pi were CD11b+CD45lo microglia (D).

**Diminished MOG35–55–specific Th1 Cytokine Responses in MCP-1--null Mice.** Significant changes in cytokine production have been described in the Th1 immune response that typifies MOG35–55 peptide–induced EAE in B6 mice (35). Serum concentrations of IFN-γ, IL-4, and IL-10 were determined by ELISA from mice immunized with MOG35–55 on days 8 and 10 pi and at the peak of EAE (day 14 pi in wild-type controls and 25 pi in MCP-1--deficient mice). At day 8 pi, concentrations of IFN-γ were slightly but significantly higher in wild-type than in MCP-1--deficient mice. This difference between wild-type and MCP-1--deficient mice became strikingly evident on day 10 pi near the onset of EAE in wild-type controls. The onset of EAE was also associated with increased serum levels...
of IFN-γ in MCP-1−/− mice, but the magnitude of increase was significantly less than in MCP-1+/+ mice (Fig. 7 A). Circulating IL-10 was detected at low levels in both MCP-1+/+ and MCP-1−/− mice before and after the onset of EAE. Before EAE onset at day 8 pi (Fig. 7 B), serum IL-10 was slightly but significantly higher in MCP-1−/− mice than in wild-type controls. Serum IL-4 remained below the limits of detection in both MCP-1+/+ and MCP-1−/− mice at all time points.

These results were supported by data from in vitro restimulation experiments, using draining lymph node (PILN) cells from mice immunized with MOG35–55 in CFA. Upon rechallenge with MOG35–55 peptide in vitro, at all examined occasions during the cell culture, PILN cells from MCP-1−/− mice secreted ~50% less IFN-γ than cells from MCP-1+/+ mice (Fig. 8, left). IL-4 and IL-10, signature Th2 cytokines in EAE (26, 29, 63, 64), were also measured. Although low levels of IL-10 were found in the culture supernatants of restimulated PILN cells from both MCP-1+/+ and MCP-1−/− mice, significantly higher levels of IL-10 were detected in cultures of cells from MCP-1−/− mice (Fig. 8, right). IL-4 was undetectable in all cell culture supernatants.

CNS Cytokine mRNA Accumulation in MCP-1−/− and Wild-type Mice with EAE. Local expression of IFN-γ in the CNS was analyzed by real-time RT-PCR. Significantly higher geometric mean levels of IFN-γ were found in spinal cord tissue from MCP-1+/+ mice at the peak of EAE attacks (MCP-1+/+: 27.7 ± 0.3, mean ± SD, n = 4; MCP-1−/−: 29.9 ± 0.8, n = 4, P < 0.05). This result indicated approximately a fourfold difference in the CNS expression of IFN-γ between mice with intact and disrupted MCP-1 genes, despite equal numbers of CNS-infiltrating T cells (see above). There was no difference in IL-10 expression in CNS tissue between MCP-1+/+ and MCP-1−/− mice with full-blown EAE. IL-4 gene expression was undetectable both in MCP-1+/+ and MCP-1−/− mice (data not shown).

Decreased Expression of IFN-γ-inducible 10-kD Protein, Macrophage Inflammatory Protein 1α, and RANTES in CNS Tissue from MCP-1−/− Mice with EAE. CNS chemokine expression was quantified using RPA, in tissues from MCP-1+/+ and MCP-1−/− mice equally affected by EAE (score 3.5–4.0). MCP-1−/− mice had significantly lower levels of IFN-γ-inducible 10-kD protein (IP-10), macrophage inflammatory protein (MIP)-1α, and regulated upon activation, normal T cell expressed and secreted (RANTES) transcripts compared with wild-type littermate controls (Fig. 9). Expression of MCP-3 was low in MCP-1+/+ and MCP-1−/− mice at the peak of EAE attack without significant differences between MCP-1−/− mice and littermate controls, supporting the results obtained using real-time RT-PCR. T cell activation gene (TCA)-3 expression in the CNS of MCP-1−/− mice with EAE was near the lower limits of detection. No significant difference was found in CCR gene expression (data not shown). However, there was a nonsignificant trend towards decreased levels of CCR2 expression in MCP-1−/− EAE CNS tissue compared with wild-type controls. In the ab-
sence of MCP-1, the presence of CCR2 in the affected tissue might indicate the action of other CCR2 ligand(s). Alternatively, the migration of CCR2-bearing cells into CNS might be a bystander phenomenon.

Compelling evidence has shown that IP-10, MIP-1α, and RANTES are potent factors that attract Th1 T cells into sites of inflammation (65–70). Further, Th1/Th2 T cells have been recently reported to differentially secrete RANTES, lymphotactin, and TCA-3, respectively (71). Enhanced expression of IP-10, MIP-1α, and RANTES in MCP-1+/+ EAE CNS tissue and undetectable TCA-3 expression in either MCP-1+/+ or MCP-1−/− support the notion that immune reactions within the CNS during EAE attacks are Th1 biased and such responses are more pronounced in mice with an intact MCP-1 gene.

**Anti-MOG Ig Isotypes in MCP-1−/− and Wild-type Mice.** Sera from MCP-1+/+ and MCP-1−/− mice with EAE were analyzed for total IgG, IgG1, and IgG2a Abs against the immunizing MOG35–55 peptide. Wild-type and MCP-1−/− deficient mice produced similar amounts of total MOG35–55 specific IgG (Fig. 10). Despite the disparity in clinical severity between wild-type and MCP-1−/− null mice, this finding was not unexpected, given the results of experiments using B cell–deficient mice, which showed that Ig does not play an important pathogenic role in MOG35–55 peptide-induced EAE in B6 mice (48).

However, Ig isotype analyses differentiated the wild-type and MCP-1−/− deficient mice. Levels of anti-MOG35–55 IgG1 Abs in wild-type controls remained low from day 14 pi through day 60 pi (Fig. 10) and were not elevated at intermediate time points (data not shown). In contrast, significantly higher levels of anti-MOG35–55 IgG1 Abs were evident in MCP-1−/− mice (Fig. 10). Levels of anti-MOG35–55 IgG2a Abs showed a trend towards elevated levels in wild-type controls on day 14 pi, whereas no difference between wild-type and MCP-1−/− null mice was found on day 60 pi (Fig. 10).

Taken together, these results suggest that a polarized MOG35–55–induced Th1 immune response in wild-type mouse leads to a suppressed Th2 response, characterized by undetectable IL-4, lower levels of IL-10, lower levels of anti-MOG35–55 IgG1, and higher levels of IgG2a. In this model, the absence of MCP-1 results in a shift towards a Th2-biased response, with reduced production of IFN-γ, enhanced secretion of IL-10, and higher levels of IgG1.

**Discussion**

We and others have previously shown that MCP-1 was markedly elevated in the CNS of SJL and B6 mice with EAE (33, 35) and levels of MCP-1 expression correlated with the severity of relapsing EAE (72). Anti-MCP-1 Abs blocked relapses of EAE (36). CNS MCP-1 is largely produced by parenchymal astrocytes (34). MCP-1 expression by astrocytes in MS brain lesions has also been convincingly documented (40–42).

However, these studies did not establish a primary role for MCP-1 in disease pathogenesis. In recent definitive studies of EAE using CCR2-deficient mice (37, 38), the relevant ligand for the deleted receptor was not defined. Moreover, MCP-1 exhibits attributes that argue for a role in restraining autoimmune demyelination. In particular, MCP-1 exerts a direct or indirect (via IL-4) impact on Th2 T cell development (11). Further, the presence of MCP-1 in vitro cell culture systems decreased the encephalitogenic potential of T cells directed to PLP139–151 (36). NK cells that inhibited the encephalitogenic potential of autoreactive T cells in DA rats produced high levels of MCP-1 in vitro (73). The role(s) of MCP-1 in the pathogenesis and development of EAE (MS) has therefore been uncertain. Using gene-targeted mice, we demonstrate that lack of MCP-1 delays the onset of EAE and ameliorates its severity, by reducing the accumulation of inflammatory leukocytes within CNS. This phenotype was associated with impaired MOG35–55–specific Th1 immune responses.

Impaired macrophage recruitment into the CNS, as indicated by reduced total number of cells and percentage of CD11b+/CD45+CD4−CD8− cells recovered from CNS in MCP-1−/− deficient mice in our study, is consistent with the reduction of macrophages in MCP-1−/− mice in contact hypersensitivity responses (2), in kidney and lung lesions of MCP-1−/− MRL–Fas−/+ mice (74), in aortic walls of MCP-1−/− and low density lipoprotein receptor double-deficient mice (75), and in atherosclerosis plaques from MCP-1−/− mice that overexpress apolipoprotein B (76). Compelling evidence suggests that macrophages and their products can be detrimental in EAE and human MS. Expression of MHC class II is markedly elevated in EAE and MS (77), costimulatory molecules such as CD80, CD86 expressed mainly by macrophages have been demonstrated in MS lesions (78), and blockade of CD28/CD80, CD86 pathway prevents epitope spreading and

![Figure 10. Serum concentrations of anti-MOG35–55 IgG, IgG1, and IgG2a Abs in MCP-1−/− and MCP-1+/+ littermate controls after immunization with MOG35–55 plus pertussis toxin (500 ng/injection). Indicated are days pi when sera were collected. Data are expressed as mean ± SD, n = 6, in each group. *P = 0.08, **P < 0.01. WT, wild-type; KO, knockout.](image-url)
clinical relapses of EAE (79). B7-1/B7-2<sup>-/-</sup> mice are resistant to EAE induction (47). Similarly, blocking of interactions between CD40 on macrophages and CD40L on T cells has been shown to effectively prevent EAE (80). Products of macrophages like TNF-α, IFN-γ, and nitric oxide have also been demonstrated to be critical in the effector phase of EAE (81, 82). Macrophage depletion inhibits the induction of EAE (83). The absence of clinical EAE in MCP-1<sup>-/-</sup> recipients of wild-type encephalitogenic T cells further indicates the importance of CNS MCP-1 expression in recruiting macrophages to the CNS. We propose that it is the failure to recruit significant number of macrophages into CNS that constitutes the principal mechanism for resistance to EAE induction in MCP-1–deficient mice. Based on these studies, we cannot exclude the possibility that MCP-1 may directly alter trafficking pattern of dendritic cells in periphery or CNS, expression of costimulators, inflammatory cytokines, and adhesion molecules. Further studies are underway to address these issues.

As several studies have shown that MCP-1 is a critical factor for T cell commitment to the Th2 phenotype, we did not anticipate that MCP-1 gene disruption would result in reduced MOG35–55–specific Th1 immune response in these EAE experiments. Our results show that MOG35–55–specific MCP-1<sup>-/-</sup> T cells secreted a large amount of IFN-γ, although less than MCP-1<sup>+/-</sup> T cells, but undetectable levels of IL-4. In view of the reciprocal regulation between IFN-γ and IL-10, the enhanced in vitro secretion of IL-10 by MCP-1<sup>-/-</sup> T cells might be secondary to reduced levels of IFN-γ. However, increased expression of IL-10 was not observed in vivo as demonstrated by the equal amount of IL-10 transcripts in MCP-1<sup>+/-</sup> and MCP-1<sup>-/-</sup> EAE CNS tissue. The fact that MOG35–55–reactive MCP-1<sup>-/-</sup> T cells mediated severe EAE in wild-type recipient mice in the passive transfer EAE model further suggests that they were Th1 polarized. An explanation for the equal encephalitogenic capacity of MCP-1<sup>-/-</sup> mice is the failure to recruit significant number of MCP-1–null macrophages into CNS that constitutes the principal mechanism for resistance to EAE induction in MCP-1–deficient mice. Based on these studies, we cannot exclude the possibility that MCP-1 may directly alter trafficking pattern of dendritic cells in periphery or CNS, expression of costimulators, inflammatory cytokines, and adhesion molecules. Further studies are underway to address these issues.

We propose that the role of MCP-1 in EAE became manifest because of the extreme Th1 polarization implicated in this model. The impaired ability to mount Th2 responses was not relevant in these experiments because the disease was severely attenuated by the reduction of macrophage recruitment to the CNS. Such reduction of macrophage reaction might subsequently result in reduced Th1 immune responses. Remarkably, in the absence of recruited macrophages, highly polarized Th1 cells became unable to express the Th1 effector program, most clearly demonstrated by decreased circulating and CNS IFN-γ and failure to elicit EAE in MCP-1–deficient mice by MOG-primed MCP-1<sup>+/-</sup> encephalitogenic T cells. Taken in the context of recent reports (37, 38), our results indicate that the MCP-1/CCR2 ligand/receptor pair is critical for the expression of EAE in mice. In turn, these findings motivate a continuing effort to characterize the function of this multipotential chemokine in human disease.

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