

Inactivation of JNK1 enhances innate IL-10 production and dampens autoimmune inflammation in the brain

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Environmental insults such as microbial pathogens can contribute to the activation of autoreactive T cells, leading to inflammation of target organs and, ultimately, autoimmune disease. Various infections have been linked to multiple sclerosis and its animal counterpart, autoimmune encephalomyelitis. The molecular process by which innate immunity triggers autoreactivity is not currently understood. By using a mouse model of multiple sclerosis, we found that the genetic loss of the MAPK, c-Jun N-terminal kinase 1 (JNK1), enhances IL-10 production, rendering innate myeloid cells unresponsive to certain microbes and less capable of generating IL-17-producing, encephalitogenic T cells. Moreover, JNK1-deficient central nervous system myeloid cells are unable to respond to effector T cell inflammatory cytokines, preventing further progression to neuroinflammation. Thus, we have identified the JNK1 signal transduction pathway in myeloid cells to be a critical component of a regulatory circuit mediating inflammatory responses in autoimmune disease. Our findings provide further insights into the pivotal MAPK-regulated network of innate and adaptive cytokines in the progression to autoimmunity.

c-Jun N-terminal kinase | experimental autoimmune encephalomyelitis | innate immunity | macrophages

Although the precise etiology of many autoimmune diseases is still unclear, both environmental and genetic factors have been implicated (1). Experimental autoimmune encephalomyelitis (EAE) serves as a useful prototype to dissect the cellular and molecular mechanisms underlying autoimmunity in multiple sclerosis. In mice, microbial components from mycobacteria or *Bordetella pertussis* can substitute for environmental conditions in the development of EAE. For example, the administration of *B. pertussis* toxin or adjuvant that contains heat-killed *Mycobacterium tuberculosis* is sufficient to trigger EAE in anti-myelin T cell receptor transgenic mice (2, 3). Similarly, housing these transgenic mice in an environment where they are regularly exposed to microbes increases the incidence of spontaneous EAE (3). In nontransgenic mice, immunization with a myelin peptide in the presence of an adjuvant containing mycobacterial components is a common strategy to activate a pathogenic, T cell-mediated, anti-myelin autoimmune inflammation. The triggering of myeloid cell activation by the microbial products initiates the activation of myelin-specific T cells in the peripheral lymphoid organs, resulting in effector T cell invasion of the CNS, neuroinflammation, and limb paralysis, a clinical outcome in EAE.

The c-Jun N-terminal kinases (JNKs) are part of the evolutionarily conserved MAPK pathways that can be activated in a plethora of cell types by mitogenic or proinflammatory signals (4). Once activated, JNKs phosphorylate multiple transcription factors, such as c-Jun, ATF-2, Elk-1, and nuclear factor of activated T cells (NFAT), to impart genomic changes in a cell responding to the environmental stress. There are three different *JNK* genes, *JNK1*, *-2*, and *-3*; each member has multiple isoforms through alternative splicing. *JNK1* and *-2* are ubiquitously expressed, whereas *JNK3* is restricted to the brain, heart, and

testis. Studies using gene-targeting disruption of specific JNKs have revealed both overlapping and distinct biological roles in cellular processes such as regulation of adipose cell metabolism, formation of osteoclasts, proliferation, and apoptosis of fibroblasts or neurons (5–8). The functional specification of JNK members is thought to depend on the responding cell types and the nature of the stimuli.

When naïve T cells are stimulated through their T cell receptors and costimulatory molecule CD28, both *JNK1* and *-2* are activated and have divergent roles in T cell activation, differentiation, and apoptosis *in vitro* (9–12). An *in vivo* consequence of such defects points to the importance of *JNK2* in *CD8⁺* T cells to mount antiviral immunity (13) and in the development of autoimmune diabetes in nonobese diabetic (NOD) mice (14). *JNK1*, however, is implicated in controlling Th2 responses after *Leishmania major* infection (15).

In myeloid macrophages, recognition of pathogen-associated molecular patterns by Toll-like receptors (TLRs) activates NF- κ B and MAPKs, which are key molecules in the induction of proinflammatory cytokines, including TNF- α , IFN- γ , IL-6, IL-12, and reactive oxygen/nitrogen intermediates (16). These proinflammatory mediators are critical for killing the pathogens as well as controlling infection by promoting macrophage phagocytosis. Such innate immune response of the myeloid cells also determines the nature of adaptive immune response of *CD4⁺* T cells and the outcome of an inflammation. In addition to JNKs, two other major MAPK pathways (p38 kinases and ERKs) are activated after TLR stimulation. The relative contribution of individual MAPK pathway in TLR signaling remains controversial and appears to depend on the nature of the ligands. For example, p38 and ERKs, but not JNKs, have been shown to be required for macrophage response to mycotoxin (17), bacterial CpG, or endotoxin from Gram-negative bacteria (18), whereas p38 and JNK activation mediate macrophage immunity to Shiga toxin (19). It remains unclear how JNKs may influence macrophage innate response to pathogenic stimuli in the development of T cell-mediated autoimmune inflammation in the brain. Here we show that *JNK1* is a reciprocal regulator of myeloid pro- and anti-inflammatory cytokine balance and the encephalitogenicity of brain-reactive T cells.

Results and Discussion

To explore the role of *JNK1* in myeloid cell activation and EAE, we used C57BL/6 mice with a disrupted *JNK1* gene. These

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Abbreviations: CFA, complete Freund's adjuvant; DLN, draining lymph node; EAE, experimental autoimmune encephalomyelitis; MOG, myelin oligodendrocyte glycoprotein; Th, T helper; TLR, Toll-like receptor; Treg, T regulatory cell.

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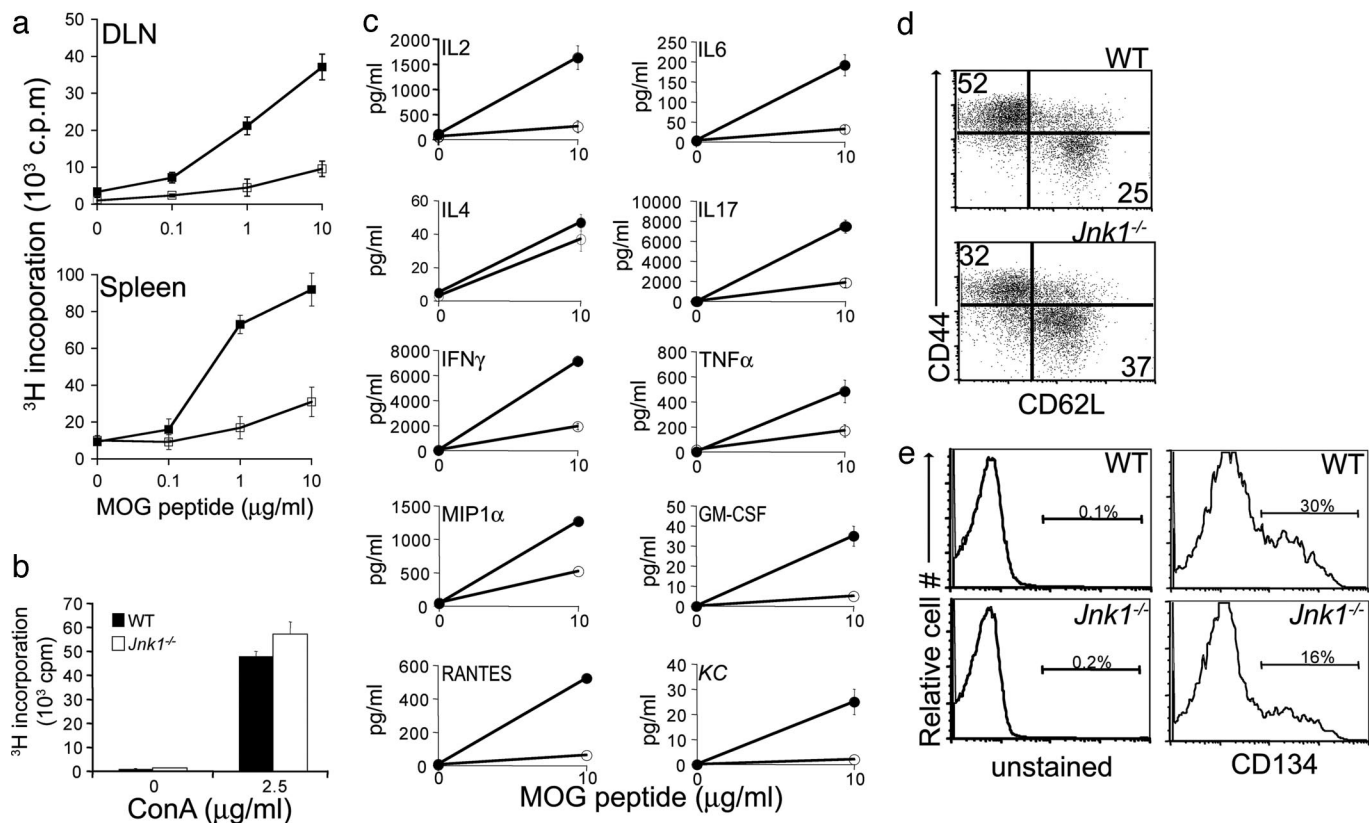


Fig. 2. Impaired generation of MOG-reactive T cells in $JNK1^{-/-}$ mice. Seven days after immunization with MOG/CFA, DLNs and splenocytes were isolated from WT (filled squares) or $JNK1^{-/-}$ (open squares) mice to measure their proliferative responses to MOG (a) or ConA (b). (c) The same number of splenocytes from MOG-immunized WT (filled circles) or $JNK1^{-/-}$ (open circles) mice were restimulated with MOG, and the production of cytokines and chemokines was measured. (d and e) Flow cytometric analysis of a gated splenic $CD44^{+}$ T cell population for a memory $CD44^{+}CD62L^{lo}$ phenotype (d) or for the expression of CD134 (e) 12 days after immunization with MOG/CFA. Results are representative of three experiments.

cytokine $IFN-\gamma$ is a strong activator of macrophages, we investigated whether the addition of $IFN-\gamma$ would restore the production of $TNF-\alpha$ in $JNK1^{-/-}$ myeloid cells. Although exogenous $IFN-\gamma$ increased the levels of $TNF-\alpha$ (from a mean of 0.42 to 3.9 ng/ml), $JNK1^{-/-}$ macrophages still produce less $TNF-\alpha$ and more IL-10 than WT cells (Fig. 3f). Hence, the altered development of encephalitogenic Th1 cells and IL-17-producing effector T cells in $JNK1^{-/-}$ mice is likely to be associated with the requirement of JNK1 activity in the myeloid compartment. The cytokine IL-10 is known to inhibit myeloid cell antigen-presenting and effector functions as well as counteract the activity of IL-12 (31). Thus, in the absence of JNK1 function in myeloid cells, immune suppression can result at several levels, including pathogenic autoreactive T cell generation, effector cytokine imbalance, and resistance to EAE induction.

To verify that lymphoid JNK1 activity is dispensable for the generation of pathogenic CD4 T cells and inflammation in the brain, we examined the ability of naïve $JNK1^{-/-}$ lymphoid cells to become encephalitogenic in a host with normal JNK1 function. We isolated naïve CD4 T cells from $JNK1^{-/-}$ or WT mice and transferred them into *Rag1*-deficient recipients, which otherwise lack T cells. All host cells, including myeloid, therefore had an intact JNK1 gene, whereas only CD4 T cells were either defective or normal for their JNK1 function. After immunization with MOG/CFA, all recipients exhibited a comparable incidence, severity, and clinical course of EAE (Fig. 4a), indicating that both $JNK1^{-/-}$ and WT T cells had comparable encephalitogenic capacity. Histological examination of the CNS also revealed a comparable degree of leukocyte infiltration and cellular composition (data not shown). Thus, the absence of

JNK1 in T cells does not alter their ability to mount a pathogenic autoimmune response to myelin. This finding is surprising in light of previous studies, in which purified $JNK1^{-/-}$ CD4 T cells showed defects in T cell activation (9) or differentiation *in vitro* (11). Although the exact reasons for such differences are unclear, possible explanations include the mouse genetic background, the stimulation method, and the complexity of an *in vivo* environment. In addition, IL-17 is key to EAE and was not measured in those studies.

To further support the importance of nonlymphoid JNK1 activity in neuroinflammation, we performed adoptive transfer of MOG-reactive, effector WT T cells into either WT or $JNK1^{-/-}$ hosts. According to the prevailing concept, the clinical signs of EAE are manifested in this model after the transferred encephalitogenic effector T cells infiltrate the CNS and stimulate local myeloid cells (e.g., microglia, macrophages, or dendritic cells), which present myelin to them for their reactivation *in situ* (32, 33). As expected, WT mice receiving MOG-reactive effector WT T cells developed clinical signs of EAE (Fig. 4b). In sharp contrast, $JNK1^{-/-}$ mice that received WT effector T cells failed to develop disease (Fig. 4b). Furthermore, these pathogenic WT T cells failed to stimulate $JNK1^{-/-}$ macrophage or microglial activation as judged by minimal CD11b/Mac-1 immunoreactivity on $JNK1^{-/-}$ spinal cord sections (Fig. 4c). Because $CD25^{+}CD4^{+}$ T regulatory cells (Tregs) have been implicated in suppressing autoimmune T cell activation and EAE (34), we investigated whether the resistance to the adoptive EAE in $JNK1^{-/-}$ mice could also be attributed to a higher number of Tregs or an enhanced suppressive function. Treg number in naïve $JNK1^{-/-}$ mice was comparable with that

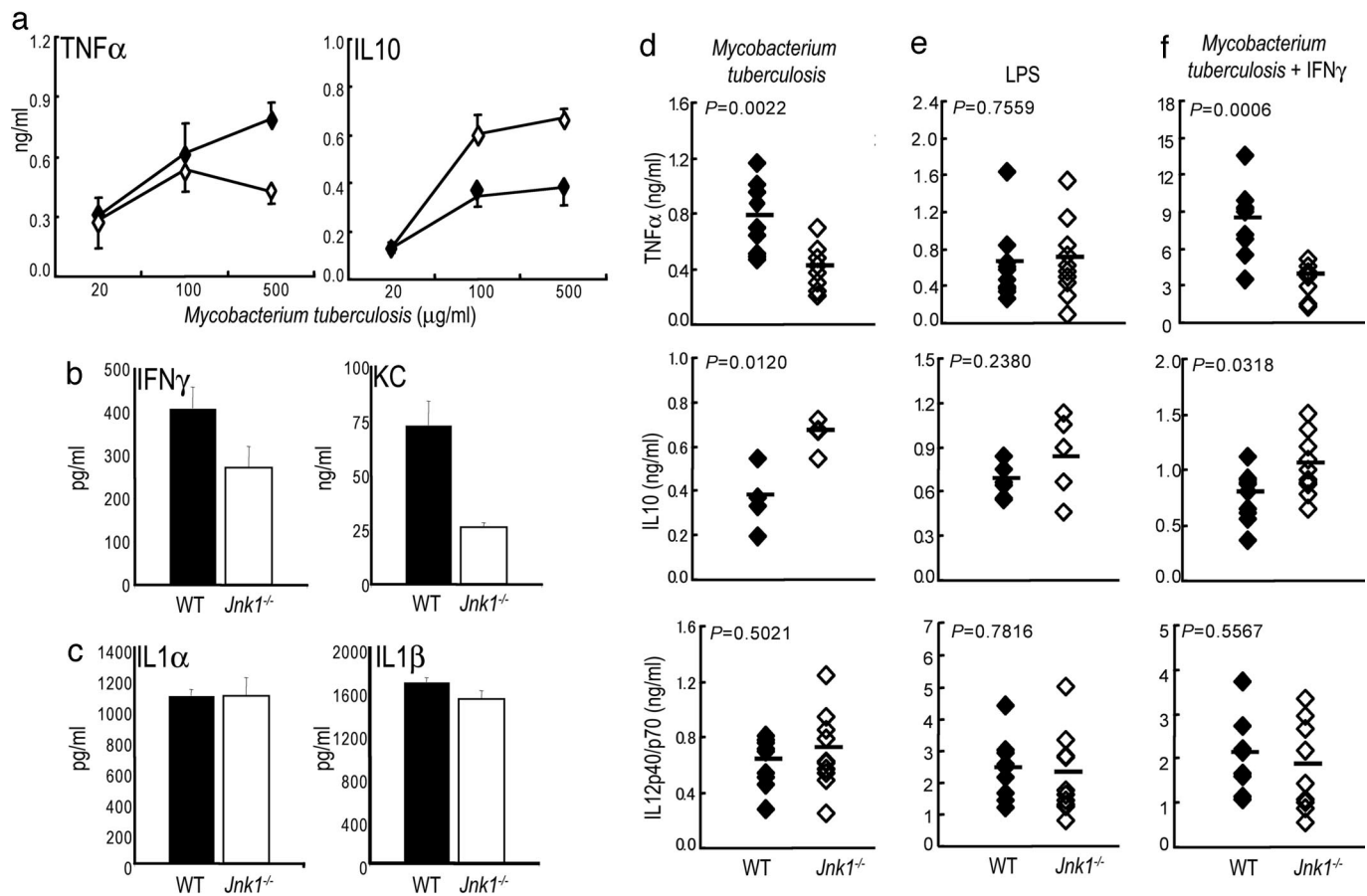


Fig. 3. JNK1^{-/-} myeloid cells produce less TNF- α and KC but more IL-10. (a) Production of TNF- α and IL-10 by WT (filled diamonds) and JNK1^{-/-} (open diamonds) bone marrow-derived macrophages after stimulation with titrated amounts of heat-killed *M. tuberculosis*. Data represent the mean of 12 and 7 mice per group for TNF- α and IL-10, respectively. Error bars indicate SE. (b and c) Secretion of IL-1 α , IL-1 β , IFN- γ , and the chemokine KC by macrophages after stimulation with 500 $\mu\text{g/ml}$ *M. tuberculosis*. Each bar represents the mean of three mice per group. (d–f) Production of the cytokine TNF- α , IL-10, and IL-12 by macrophages when cultured with LPS (10 ng/ml) or *M. tuberculosis* (500 $\mu\text{g/ml}$) in the presence or absence of IFN- γ (20 units/ml) for 24 h. Symbols represent individual mice, and the mean is indicated by a solid line. *P* values were calculated with Student's *t* test.

in WT mice (7% vs. 8%, respectively). In addition, we found no difference between JNK1^{-/-} and WT Treg in their ability to suppress the activation of CD25⁻CD4⁺ T cells by using the standard *in vitro* coculture assay (Fig. 4d). Hence, EAE resistance in JNK1^{-/-} recipients is not due to enhanced Treg function on the transferred encephalitogenic effector T cells. Rather, one possible scenario is that microglial activation, which is crucial for EAE progression (35), requires JNK1 activity to mediate signals from effector T cell proinflammatory cytokines, such as IL-17. Of interest, inhibitors of the JNK pathways were shown to block IL-17 responsiveness in human macrophages (36). Also, JNK1^{-/-} macrophages did not seem to enable them to the effector T cell cytokine IFN- γ (Fig. 3f). Because CNS microglia are bone-marrow derived, our data suggest that JNK1 may positively regulate T cell inflammatory signaling in microglia and macrophages for EAE progression in mice. Alternatively, JNK1 may modulate TLR and myeloid differentiation factor 88 (MyD88) signaling in microglia and macrophages within the CNS for the progression of EAE. In support of this hypothesis, stimulation of TLR on microglia has been shown to enable them to efficiently present myelin antigens to CD4⁺ T cells (37). Moreover, mice deficient in either MyD88 or TLR9 did not develop EAE after the adoptive transfer of WT encephalitogenic T cells (38). Our observation that WT encephalitogenic T cells failed to induce EAE in JNK1^{-/-} recipient mice upon transfer points to a potential role for JNK1 in the regulation of TLR

signaling in CNS myeloid cells. The use of an adoptive transfer system is particularly useful to examine microglial activation and EAE progression without the complication of the adjuvant effect, which persists in the immunization model. In fact, in addition to its activation of the innate immunity in lymphoid organs, *M. tuberculosis* in CFA can cause other systemic changes (e.g., myelopoeisis, blood–brain barrier damage, and activation of CNS resident cells) that result in lowering the threshold for the induction of neuroinflammation (39–41). As illustrated in Fig. 3, the absence of JNK1 selectively affects the production of cytokines in response to heat-killed *M. tuberculosis*. The production of IL-1 was normal, suggesting that certain systemic changes induced by CFA likely remain intact in JNK1^{-/-} mice. Hence, these systemic factors may contribute to the mild EAE in MOG/CFA-immunized JNK1^{-/-} mice (Fig. 1), in which the initial immunization failed to produce a critical quantity of encephalitogenic T cells (Fig. 2). Because CFA was not given to recipient mice in the adoptive transfer model, the CFA influence that facilitates neuroinflammation is absent, thereby increasing the threshold for the activation of CNS microglia and progression to EAE. Therefore, the CFA effect likely contributes to the mild disease in the immunization model and its absence to the lack of disease in the adoptive transfer model.

Collectively, our studies point to a critical and nonredundant role for JNK1 activity in nonlymphoid cells as being an important regulator of inflammation in the brain and possibly other organs.

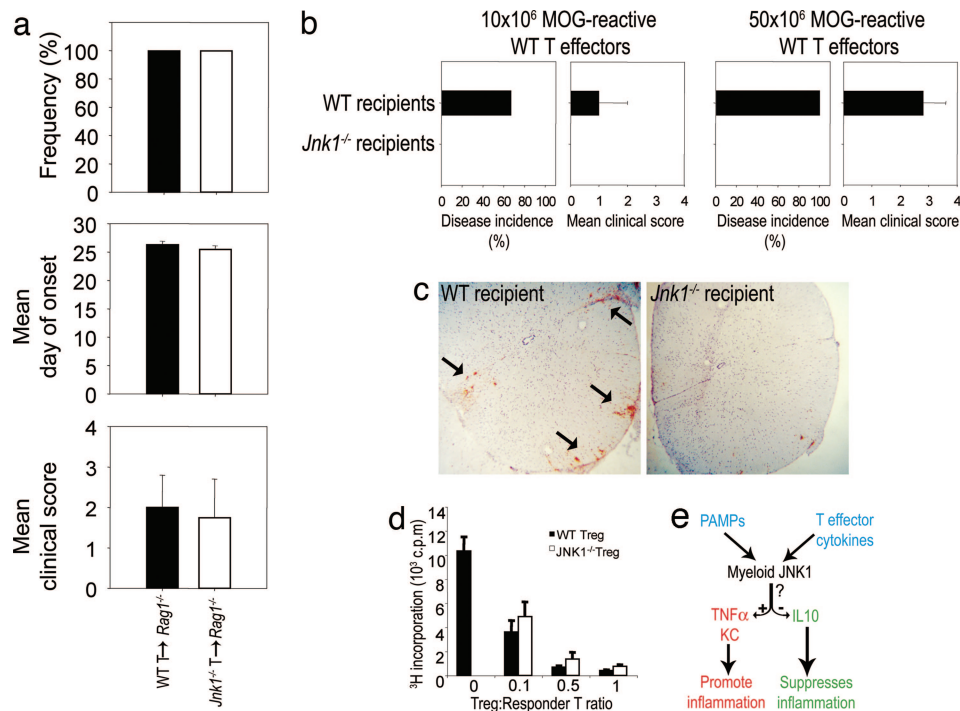


Fig. 4. Adoptive transfers of naïve or effector T cells. (a) WT or JNK1^{-/-} naïve CD4⁺ T cells equivalently provoked EAE in Rag1^{-/-} hosts ($n = 4$ per group) in response to immunization with MOG/CFA. (b) MOG-reactive WT effector T cells failed to induce EAE in JNK1^{-/-} recipient mice ($n = 4$). This procedure was repeated in a separate experiment with another four recipients per group. (c) Spinal cord sections from a WT recipient with EAE and an asymptomatic JNK1^{-/-} recipient were immunostained for CD11b/Mac-1⁺ (arrows) activated microglia and macrophages. (d) WT or JNK1^{-/-} Tregs (CD25⁺CD4⁺) had the same potency in suppressing the activation of WT responder effector T cells. (e) A proposed model implicating myeloid JNK1-mediated regulation of counteracting immune cytokines and responses.

We propose that JNK1 activity in myeloid cells is induced in response to microbial pathogens and functions as a reciprocal regulator of counteracting cytokines that shape the outcome of the innate and acquired immune response (Fig. 4e). Invading microbial pathogens are detected by TLRs on myeloid cells. Although TLR signaling depends heavily on myeloid differentiation factor 88 (MyD88) (16), JNK1 activity may further fine-tune the innate myeloid cell response. Of note, it may do so selectively depending on the types of pathogens, TLRs, or inflammatory milieu. Our findings that JNK1 contributes to both positive and negative regulation of counteracting myeloid-derived cytokines suggest that this pathway is a viable target for therapeutic intervention in autoimmune and infectious diseases.

Methods

Mice and Immunization. JNK1^{-/-} mice (11) were backcrossed onto the C57BL/6 genetic background for at least nine generations, and WT C57BL/6 control mice were purchased from The Jackson Laboratory (Bar Harbor, ME). RAG1^{-/-} mice were bred and maintained in our animal facilities. EAE was elicited in female or male mice at 8–12 weeks of age by s.c. tail-base injection of 50 μ g of MOG_{35–55} peptide (35-MEVGWYRSPF-SRVVHLYRNGK-55) in CFA containing 500 μ g of heat-inactivated *M. tuberculosis* (H37RA; DIFCO Labs, Sparks, MD), and 200 ng of *B. pertussis* toxin (LIST Biological Labs, Inc., San Jose, CA) was injected i.p. on days 0 and 2. Mice were monitored daily for clinical signs of EAE and scored as follows: 1, flaccid tail; 2, inability to right and one hind limb paralysis; 3, paralysis of both hind limbs with or without incontinence; 4, paralysis of all limbs; and 5, moribund. Supplementary food and water were provided on the cage floor for disabled animals. Animal maintenance and all experimental protocols were in accordance with Yale Animal Research Committee.

Adoptive Transfer of MOG-Reactive Effector T Cells. C57BL/6 WT mice were immunized as described above. Ten days later, a single cell suspension was prepared from DLNs, and cells (3×10^6 per ml) were cultured in the presence of 10 μ g of MOG and 5 units/ml recombinant human IL-2 in RPMI 1640 (Invitrogen, Carlsbad, CA) with 10% FCS, 50 μ M 2-mercaptoethanol, 100 units/ml penicillin, and 100 μ g/ml streptomycin. After 4 days of culture, nonadherent T cells were collected by centrifugation on Ficoll, and $10\text{--}50 \times 10^6$ T cells were injected i.v. into naïve WT or JNK1^{-/-} recipient mice. Each mouse also received 200 ng of pertussis toxin on days 0 and 2 after transfer.

Adoptive Transfer of Naïve T Cells. Naïve T cells (CD4⁺CD62L^{hi}CD44^{lo}) from WT or JNK1^{-/-} mice were isolated on a FACS Vantage sorter, and 10×10^6 were injected in the tail vein of RAG1^{-/-} recipient mice. After 24 h, all recipients were immunized with MOG as described above.

In Vitro Recall Response Assay. Cells from spleens or DLNs were isolated 6 days after immunization and were cultured for 4 days in the presence of indicated amounts of MOG peptide or Con A (Sigma, St. Louis, MO). To assess proliferation, cultures were pulsed with 1 μ Ci (1 Ci = 37 GBq) of [³H]thymidine during the last 18 h of incubation. The mean [³H]thymidine uptake in triplicate wells was measured as cpm. To measure production of cytokines and chemokines, supernatants were collected after 2–4 days of culture with MOG and subjected to analysis by specific enzyme-linked immunosorbent assays ELISA (for IL-2, IL-4, and IFN- γ), FlowCytomix (for IL-6 and TNF- α), and Bio-Plex [for MIP-1 α , RANTES, KC, granulocyte/macrophage colony-stimulating factor (GM-CSF), and IL-17] according to the manufacturer's instructions. Paired Abs against cytokines in ELISA were purchased from PharMingen (San Diego, CA).

FlowCytomix multiplex bead kits were obtained from Bender-MedSystems (Burlingame, CA), and Bio-Plex multiplex bead kits from Bio-Rad Laboratories (Hercules, CA).

Flow Cytometry. Cells were stained with fluorochrome-conjugated primary Abs for 30 min at 4°C in PBS containing 1% FCS and 0.01% sodium azide, washed twice, acquired on FACSCalibur flowcytometer (BD Biosciences, San Diego, CA), and analyzed with FlowJo software (Tree Star, Ashland, OR). All Abs against CD4, CD25, CD44, CD62L, and CD134 were purchased from PharMingen.

Histology. Spinal cords were removed after intracardial perfusion with ice-cold PBS and were fixed in 10% formalin. Paraffin-embedded sections were stained with hematoxylin and eosin. Immunohistochemistry on 10- μ m frozen sections was performed as described (42). The primary rat Ab against mouse CD11b/Mac-1 was obtained from PharMingen, and the biotinylated secondary Ab was from Vector Laboratories (Burlingame, CA).

Cytokine Production by Bone Marrow-Derived Macrophages. Bone marrow cells were cultured in DMEM (Invitrogen) with 20% FBS and 30% L929 supernatant containing macrophage-

stimulating factor for 6 days. Macrophages (1×10^6 cells per ml) were then cultured in DMEM supplemented with 5% FBS in the presence of heat-killed *M. tuberculosis*, LPS, CpG, or medium alone for 24 h. The concentration of cytokines and chemokines was determined in culture supernatants by ELISA (PharMingen and eBioscience, San Diego, CA) and Bio-Plex analysis (Bio-Rad), according to the manufacturer's guidelines.

Treg Suppression Assay. CD25⁻CD4⁺ responder T cells (5×10^4 cells per well) and CD25⁺CD4⁺ Tregs were separated on a MoFlo cytometer and cultured at the indicated ratio in 96-well round-bottom plates with T cell-depleted spleen cells as accessory cells (3,000 R irradiated) and 0.5 μ g/ml anti-CD3 Ab (clone 2C11) for 72 h. Cultures were pulsed with [³H]thymidine for the last 6 h of culture.

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