

T cell interactions with microglia in immune-inflammatory processes of ischemic stroke

Yuxiao Zheng^{1, #}, Zilin Ren^{1, #}, Ying Liu^{1, #}, Juntang Yan², Congai Chen³, Yanhui He¹, Yuyu Shi¹, Fafeng Cheng¹, Qingguo Wang^{1, *}, Changxiang Li^{1, *}, Xueqian Wang^{1, *}

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Abstract

The primary mechanism of secondary injury after cerebral ischemia may be the brain inflammation that emerges after an ischemic stroke, which promotes neuronal death and inhibits nerve tissue regeneration. As the first immune cells to be activated after an ischemic stroke, microglia play an important immunomodulatory role in the progression of the condition. After an ischemic stroke, peripheral blood immune cells (mainly T cells) are recruited to the central nervous system by chemokines secreted by immune cells in the brain, where they interact with central nervous system cells (mainly microglia) to trigger a secondary neuroimmune response. This review summarizes the interactions between T cells and microglia in the immune-inflammatory processes of ischemic stroke. We found that, during ischemic stroke, T cells and microglia demonstrate a more pronounced synergistic effect. Th1, Th17, and M1 microglia can co-secrete pro-inflammatory factors, such as interferon- γ , tumor necrosis factor- α , and interleukin-1 β , to promote neuroinflammation and exacerbate brain injury. Th2, Treg, and M2 microglia jointly secrete anti-inflammatory factors, such as interleukin-4, interleukin-10, and transforming growth factor- β , to inhibit the progression of neuroinflammation, as well as growth factors such as brain-derived neurotrophic factor to promote nerve regeneration and repair brain injury. Immune interactions between microglia and T cells influence the direction of the subsequent neuroinflammation, which in turn determines the prognosis of ischemic stroke patients. Clinical trials have been conducted on the ways to modulate the interactions between T cells and microglia toward anti-inflammatory communication using the immunosuppressant fingolimod or overdosing with Treg cells to promote neural tissue repair and reduce the damage caused by ischemic stroke. However, such studies have been relatively infrequent, and clinical experience is still insufficient. In summary, in ischemic stroke, T cell subsets and activated microglia act synergistically to regulate inflammatory progression, mainly by secreting inflammatory factors. In the future, a key research direction for ischemic stroke treatment could be rooted in the enhancement of anti-inflammatory factor secretion by promoting the generation of Th2 and Treg cells, along with the activation of M2-type microglia. These approaches may alleviate neuroinflammation and facilitate the repair of neural tissues.

Key Words: brain; immune; inflammation; interaction; ischemic stroke; mechanism; microglia; neuron; secondary injury; T cells

Introduction

Stroke is a devastating cerebrovascular event that occurs due to an interruption of cerebral blood flow. Such events are classified as either ischemic stroke or hemorrhagic stroke. In the past, the brain was thought to be an immune-privileged organ protected by the blood–brain barrier (Halder et al., 2023; Wei et al., 2023; Yue and Hoi, 2023; Singh et al., 2024). Recent studies however have revealed that cerebral ischemia/reperfusion injury can compromise the integrity of the blood–brain barrier and

induce endogenous signaling and peripheral antigen activation (Goldmann and Prinz, 2013; Qiu et al., 2021; Candelario-Jalil et al., 2022). Subsequently, there is a massive infiltration of peripheral immune cells into the site of injury, in which both innate and adaptive immune responses are involved, leading to full-blown neuroinflammation. The inflammatory response in the brain after ischemic stroke may accelerate the formation of ischemic damage, affecting neuronal death and neural tissue regeneration and leading to secondary damage (Bayraktutan,

¹School of Traditional Chinese Medicine, Beijing University of Chinese Medicine, Beijing, China; ²Library, Beijing University of Chinese Medicine, Beijing, China;

³Dongzhimen Hospital, Beijing University of Chinese Medicine, Beijing, China

*Correspondence to: Qingguo Wang, PhD, wangqg8558@sina.com; Changxiang Li, PhD, changxiang1202@163.com; Xueqian Wang, PhD, wxqbcum@126.com.
<https://orcid.org/0000-0002-7682-3877> (Xueqian Wang); <https://orcid.org/0000-0001-8382-1717> (Changxiang Li);

<https://orcid.org/0000-0003-2752-1925> (Qingguo Wang)

These authors contributed equally to this paper.

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2019). The neuroinflammatory response after cerebral ischemia is characterized by the activation of microglia and astrocytes and an increase in inflammasomes. Microglia are the predominant immune cells in the central nervous system (CNS) and the first line of immune defense in the CNS (Lambertsen et al., 2019). As the first immune cells to be activated after ischemic stroke, microglia in the ischemic core and penumbra are induced to aggregate by Toll-like receptor (TLR)-mediated nuclear factor kappa-B (NF-κB) pathways within minutes. Activated microglia remove cellular debris and release inflammatory mediators while expressing major histocompatibility complex II (MHC II), which functions to present antigens to other cells (Abellanas et al., 2019).

After the onset of ischemic stroke, peripheral blood immune cells, including myeloid dendritic cells, lymphocytes, monocytes/macrophages, and neutrophils, are recruited by the chemokines secreted by immune cells in the brain; they migrate into the brain within 1 day after stroke, peaking at 3 days after stroke, and persist until day 7 after stroke. Peripheral T cells can re-edit their gene expression profiles, upregulate the expression of chemokine receptors and adhesion molecules, and promote their migration toward inflammatory tissues while enhancing their ability to cross the blood–brain barrier. They thus enter the CNS to interact with CNS cells and trigger a secondary neuroimmune response (Yoshida et al., 2022).

After ischemic stroke, microglia can promote the infiltration of T cells into the nervous system by means of antigen presentation. This interaction can induce T cell differentiation toward two distinct subpopulations: pro-inflammatory and anti-inflammatory cells. Additionally, peripheral T cells entering the CNS can regulate the function of resident microglia. It is worth exploring how these two immune cells influence the course of the inflammatory response through their mutual interactions. Therefore, the aim of

this paper is to explore the mechanisms of interaction between T cells and microglia and classify them into pro-inflammatory and anti-inflammatory directions. This understanding can guide future therapeutic approaches for ischemic stroke, enabling the development of targeted drugs to block the pro-inflammatory links between T cells and microglia while promoting their anti-inflammatory mechanisms. The glial cell line-derived neurotrophic factor (GDNF) secreted by microglia during inflammation suppression can promote the repair and regeneration of neurons and therefore play a neuroprotective role. The regeneration of brain tissue is an important manifestation of ischemic stroke repair. Therefore, the information covered in this review is closely connected to nerve regeneration and provides a theoretical support for future treatments of ischemic stroke.

Search Strategy and Selection Criteria

To conduct this narrative review, the literature was searched using a combination of the following terms: “microglia” AND “ischemic stroke,” “T cell” AND “ischemic stroke,” “Th1 cell” AND “microglia” OR “ischemic stroke,” “Th17 cell” AND “microglia” OR “ischemic stroke,” “Treg” AND “microglia” OR “ischemic stroke,” “immune-inflammatory” AND “microglia” AND “ischemic stroke,” and “immune-inflammatory” AND “T cell” AND “ischemic stroke.” The PubMed database was used to search for relevant articles published between January 2000 and April 2023. Articles that provided information on the classification of T cells and microglia, their physiological functions, interactions between T cells and microglia in ischemic stroke, and their role in immunoinflammation were included in this review to explore the interactions among T cells and microglia in immune inflammation in ischemic stroke. A timeline of findings on the roles of microglia and T cells in ischemic stroke provided in the literature is presented in **Figure 1**.

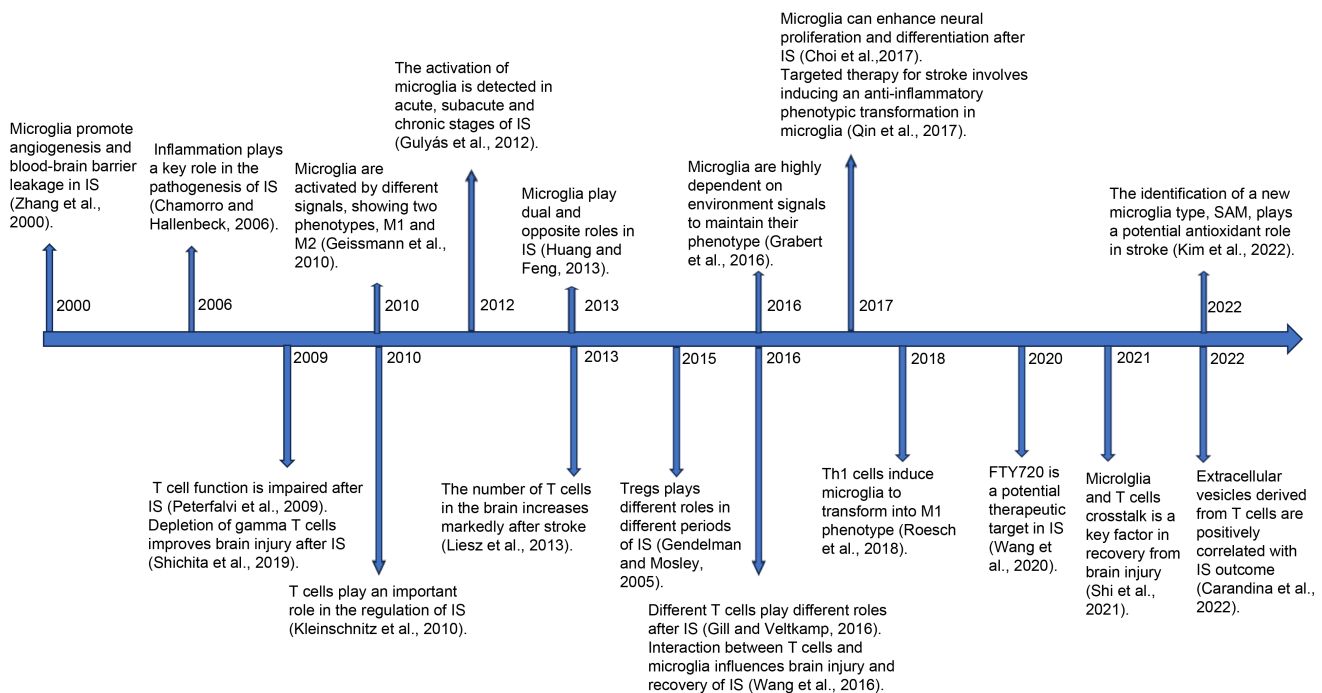


Figure 1 | Timeline showing the roles of microglia and T cells in ischemic stroke described in the literature. SAM: Stroke-associated microglia; IS: ischemic stroke.

Although microglia were discovered by Franz Nissl (Ziebell et al., 2012) and F. Robertson in the late 19th century through immunohistochemical experiments, they were first named “microglia” in the 1920s. In the century that followed their discovery, their important role in encephalopathy has been gradually explored. Huang and Feng (2013) reported that microglia play important roles in ischemic stroke. Microglia can promote angiogenesis and blood–brain barrier (BBB) leakage during (Zhang et al., 2000) and enhance neuronal proliferation and differentiation after ischemic stroke (Choi et al., 2017). Under various signal stimulations, the two phenotypes of M1 and M2 were shown to interact and regulate the treatment and prognosis of ischemic stroke patients (Geissmann et al., 2010). By promoting the transformation of microglia to an anti-inflammatory phenotype, they can be targeted in treatments (Qin et al., 2017). In 2022, a novel cell type, stroke-associated microglia, was identified and reported to potentially play an antioxidant role in stroke (Kim et al., 2022).

T cells, essential immune cells in the human body, are found as a wide range of cellular species and have complex functions. They play significant regulatory roles in ischemic stroke, with different T cell types playing distinct roles at different stages (Gill and Veltkamp, 2016). For instance, regulatory T cells (Tregs) are considered the primary neuroprotective regulators in ischemic stroke (Gendelman and Mosley, 2015). Th1 cells promote microglial phenotypic conversion (Roesch et al., 2018), while the depletion of γ T cells mitigates brain damage following ischemic stroke (Shichita et al., 2009).

Peripheral Immune Cells: T lymphocytes

Introduction to T cells

T lymphocytes, or T cells, originate from bone marrow and mature in the thymus (Adu-Berchie et al., 2023). Upon reaching maturity, they are distributed via the bloodstream and settle in thymus-dependent areas of peripheral immune organs. They can be recirculated via lymphatics, peripheral blood, and tissue fluids to execute functions such as cellular immunity and immunomodulation. T cells play pivotal roles in the adaptive immune system (Qin et al., 2020), and execute important roles in the crosstalk between the innate and adaptive immune systems (Ponsaerts et al., 2021). Within the body, T cells are constantly renewed, and various subpopulations at different developmental stages or functions can coexist simultaneously. The taxonomic nomenclature of T cells is therefore highly complex.

Classification of T cells

Based on differences in their T cell antigen receptors (TCRs), T lymphocytes can be classified into $\alpha\beta$ T lymphocytes and $\gamma\delta$ T lymphocytes. The TCR is a specific receptor that allows T cells to recognize and bind foreign antigens, serving as a unique identifier for T cells. The receptor is a heterodimer composed of two distinct peptide chains. The TCR of $\alpha\beta$ T lymphocytes consists of a heterodimer of α and β chains, while that of $\gamma\delta$ T lymphocytes comprises a heterodimer of γ and δ chains. In adult peripheral blood, 95% of T cells belong to the $\alpha\beta$ T lymphocyte category, with only a small fraction belonging to $\gamma\delta$ T lymphocytes. However, the role of the latter should not be overlooked. Furthermore, $\alpha\beta$ T lymphocytes can be further subclassified into CD4 T cells and CD8 T cells based on their surface expression of differentiation antibody population cluster of differentiation molecules. In peripheral lymphoid tissues, CD4 T cells account for approximately 65% of T cells. The CD4 T cell TCR recognition of antigens is MHC class II molecule-restricted, i.e., it is stimulated when CD4 T cells bind to antigenic MHC class II molecules expressed on the surface

of antigen-presenting cells (APCs) (Lu, 1995).

The primary role of $\alpha\beta$ T lymphocytes is to promote the proliferation and differentiation of B cells, T cells, and other immune cells. This assistance is crucial for both humoral immunity and cellular immunity. CD8 T cells, which account for approximately 35% of T cells, have TCR recognition that is MHC class I molecule-restricted. These CD8 T cells play significant roles in immune defense and immune surveillance. They eliminate intracellular infections and malignant cells by secreting cytokines and directly killing infected cells, and this process is crucial in providing long-term protective immunity. CD8 T cells are classified further into cytotoxic T cells (TC) and suppressor T cells (TS) based on their functions (Zhang et al., 2021). CD4 T cells, however, are categorized into conventional helper T cells (Th) and Tregs (Hedrick, 2002). Conventional Th cells are further divided into Th1 cells, Th2 cells, Th9 cells, Th17 cells, and T follicular helper cells (Tfh), depending on their cytokine secretion patterns (Stockinger et al., 2006; Zhu et al., 2010; Crotty, 2011; Lei et al., 2021), and these subsets possess distinct markers and functions.

Functions of Th1 cells

The surface markers of Th1 cells include $\alpha\beta$ TCR, CD3, CD4, interleukin (IL)-12 receptor (R), interferon gamma (IFN- γ)R, and C-X-C motif chemokine receptor 3. The major transcription factors for Th1 cell differentiation are T-box expressed in T cells (T-bet), signal transducer and activator of transcription (STAT)4, and STAT1, which promote the expression of specific genes in Th1 cells (Watanabe et al., 2020). The major effector molecules secreted by Th1 cells include IFN- γ , IL-2, and lymphotoxin alpha. These molecules promote protective immunity against intracellular pathogens by secreting IFN- γ , which induces macrophage activation and the upregulation of inducible nitric oxide synthase (iNOS), leading to intracellular pathogen death (Gocher et al., 2022).

Functions of Th2 cells

The surface markers of Th2 cells include $\alpha\beta$ TCR, CD3, CD4, IL-4R, IL-33R, C-C motif chemokine receptor 4, IL-17R B, chemoattractant receptor-homologous molecule expressed on T-helper type 2 cells, and others. The major transcription factors for Th2 cell differentiation are GATA-binding protein 3, STAT6, differentiated embryo-chondrocyte expressed gene 2, and macrophage activation factor (Zhang et al., 2017). Additionally, IFN regulatory factor 4 is a transcription factor involved in Th2 cell differentiation (Williams et al., 2013). These transcription factors promote the expression of specific genes in Th2 cells. The major effector molecules secreted by Th2 cells are IL-4, IL-5, IL-13, and IL-10. These cytokines play crucial roles in promoting humoral immune responses and host defenses against extracellular parasites; however, they also contribute to allergic reactions and asthma (Altin et al., 2012).

Functions of Th9 cells

The surface markers of Th9 cells encompass $\alpha\beta$ TCR, CD3, and CD4. The key transcription factor driving Th9 cell differentiation is the Spi-1 proto-oncogene (PU.1), which facilitates the expression of specific genes in these cells. The predominant effector molecules secreted by Th9 cells are IL-9 and IL-10. These cytokines primarily contribute to the host's defense against extracellular parasites. Additionally, Th9 cells produce IL-10, which can exacerbate allergic inflammation. However, given that this Th9 subpopulation has been recently characterized, its involvement in other inflammatory conditions remains unclear (Khokhar and Purohit, 2024).

Functions of Th17 cells

The surface markers of Th17 cells include $\alpha\beta$ TCR, CD3, CD4, IL-23R, C–C motif chemokine receptor 6, IL-1R, and CD161 (humans only). The major transcription factors for Th17 cell differentiation are retinoic acid receptor-related orphan receptor γ t, STAT3, and related orphan receptor A, which promote the expression of Th17-cell-specific genes. The main effector molecules secreted by Th17 cells are IL-17A, IL-17F, IL-21, IL-22, and C–C motif chemokine 20. These cytokines are primarily distributed on mucosal surfaces and play crucial roles in promoting protective immunity against extracellular bacteria and fungi. Additionally, they contribute to the treatment of autoimmune and inflammatory diseases (Wang et al., 2014).

Functions of Tfh cells

The surface markers of Tfh cells include $\alpha\beta$ TCR, CD3, CD4, C–X–C motif chemokine receptor 5, signaling lymphocytic activation molecule, OX40L (CD252), CD40 ligand, inducible T cell costimulator (ICOS), IL-21R, and programmed cell death-1 (PD-1). The pivotal transcription factor for Tfh cell differentiation is B-cell lymphoma 6, facilitating the expression of specific genes in these cells. The primary effector molecule secreted by Tfh cells is IL-21, which plays a crucial role in information transfer during B-cell differentiation. Specifically, IL-21 facilitates the activation of B cells, promotes the formation of germinal centers and immunoglobulin class switching, and sustains prolonged humoral immune responses (Crotty, 2014; Jogdand et al., 2016).

Functions of cytotoxic T cells and suppressor T cells

TC cells function by recognizing target cells expressing specific antigens, establishing contact with them, and releasing cytotoxic agents such as perforins, granzymes, and granzyme hemolysins. These substances form transmembrane pore channels, ultimately leading to the lysis and death of the target cells. Alternatively, TC cells can induce apoptosis by manipulating the Fas–Fas ligand system within the target cells. Conversely, TS cells suppress the activity of Th cells, indirectly inhibiting B cell differentiation and the cytotoxic function of TC cells. This leads to negative regulation of both humoral and cellular immunity (Utzschneider et al., 2016; Prasad Singh et al., 2020).

Functions of $\gamma\delta$ T cells

$\gamma\delta$ T cells are the initial T cell population to emerge in the thymus. However, their relative abundance diminishes with the emergence and growth of $\alpha\beta$ T cells. In adulthood, $\gamma\delta$ T cells constitute 1%–5% of peripheral blood T cells (Kou et al., 2022), whereas in specific regions of the intestine, they can constitute up to 30% of all T cells (Mayassi and Jabri, 2018). Furthermore, $\gamma\delta$ T cells are extensively present in epithelial-dense tissues that form the inner and outer surfaces of the body, including the germinal tract, intestinal epithelium, and the epidermis and dermis of the skin (Alves de Lima et al., 2020). $\gamma\delta$ T cells express intracellular TLR2, TLR17, the central TLR junction molecule MyD54, as well as TLR55 (Schafer et al., 2012); they are characterized by the expression of $\gamma\delta$ TCR and CD3 molecules on their surface, and their main secreted effector molecules include IFN- γ , IL-17A, IL-17F, and IL-22. Cells producing IFN- γ and IL-17A are distinct populations that can be distinguished by the expression of CD27 and C–C motif chemokine receptor 6 on their surface. $\gamma\delta$ T cells possess both anti-inflammatory and pro-inflammatory functions and contribute to both innate and adaptive immunity. Their recognition of innate immunity is affected by the expression of TLRs. The surface phenotypes and functions of T cell subpopulations are summarized in **Table 1**.

Table 1 | Surface phenotypes of T cell subsets and their functions

T cell categorization	Surface phenotype	Function
$\alpha\beta$ T	CD3, $\alpha\beta$ TCR	–
CD4 ⁺ $\alpha\beta$ T	CD3, $\alpha\beta$ TCR, CD4	–
Th1	CD3, $\alpha\beta$ TCR, CD4, CXCR3, CCR5, IFN- γ , IL-12R	Th1 cells are involved in cellular immune responses as well as defenses against pathogenic microbial infections, including promotion of proliferation and differentiation of cytotoxic T cells and phagocytosis by macrophages.
Th2	CD3, $\alpha\beta$ TCR, CD4, CCR3, CCR4, IFN- γ , IL-4R, IL-17R, IL-33R	Th2 cells are involved in the humoral immune response and in defending against parasitic infections, including promoting the proliferation and differentiation of B cells into plasma cells that produce antibodies.
Th9	CD3, $\alpha\beta$ TCR, CD4	Th9 cells are associated with tumor progression, may exert both antitumor and pro-tumor effects, and promote the development and progression of certain inflammatory diseases.
Th17	CD3, $\alpha\beta$ TCR, CD4, CCR6, CD161, IL-1R, IL-23R	Th17 cells are involved in the inflammatory response and defense against extracellular pathogenic microorganisms, especially in the intestinal and respiratory mucosa.
Treg	CD3, $\alpha\beta$ TCR, CD4, CD25, CD39, CD73, CD103	Treg is a class of cells that can negatively regulate immune responses, and its most prominent function is to maintain self-tolerance and immune homeostasis.
Tfh	CD3, $\alpha\beta$ TCR, CD4, CD84, CD126, CXCR5, OX40L, CD40L, PD1	Tfh aids B-cell survival and proliferation in the germinal center and promotes B-cell differentiation to plasma cells, antibody class switching, and antibody affinity maturation.
CD8 ⁺ $\alpha\beta$ T	CD3, $\alpha\beta$ TCR, CD8, CCR7, IL-7R	CD8 ⁺ $\alpha\beta$ T cells specifically recognize antigenic peptide-MHC class I molecular complexes and thus kill target cells, including cells infected by intracellular pathogens and tumor cells.
$\gamma\delta$ T	CD3, $\gamma\delta$ TCR	$\gamma\delta$ T has both anti-inflammatory and pro-inflammatory functions, both intrinsic and adaptive immunity, and its recognition of intrinsic immunity is influenced by the expression of TLRs.

CCR: Chemokine receptor; CD: cluster of differentiation; CD40L: CD40 ligand; CDL: cluster of differentiation 126; CXCR: C–X–C motif chemokine receptor 3; CXCR5: C–X–C motif chemokine receptor 5; IFN- γ : interferon- γ ; IL-17R: interleukin 17 receptor; OX40L: OX40 ligand; PD1: programmed death-1; TCR: T cell antigen receptor; Tfh: T follicular helper cells; Th: helper T cell; Treg: regulatory cells.

T cell infiltration into the brain after ischemic stroke

After ischemic stroke, innate immune cells are activated, and T cells are induced to infiltrate the brain tissue. There are three main routes of infiltration: the blood-brain barrier, choroid plexus, and soft meningeal infiltration (Benakis et al., 2018). Following stroke, immune cells accumulate near the epithelial cells of the blood–brain barrier and secrete inflammatory factors that disrupt its integrity, allowing T cells to infiltrate the ischemic brain tissue. The choroid plexus is a structure that produces cerebrospinal fluid, and after ischemic stroke, there is a significant increase in chemokine expression in the injured brain tissue. T cells can migrate from the choroid plexus to the brain parenchyma along a chemokine gradient (Zhang et al., 2021b).

As for the soft meningeal infiltration pathway, fluorescent cell-tracking experiments have shown that labeled intestinal $\gamma\delta$ -T cells accumulate in the soft meninges in the early stages of stroke (Benakis et al., 2016), but the specific pathway of their migration to the meninges remains unclear and requires further investigation.

A study by Feng et al. (2017) demonstrated that T cells begin to infiltrate ischemic brain tissue as early as 24 hours after stroke and can remain in the brain tissue until post-ischemia week 7. Different types of T cells infiltrate the brain at different times and are simultaneously regulated by the varying effects of various cytokines and chemokines (Zhang et al., 2021b). CD4 T cells infiltrate the infarcted area on the first day and increase in numbers significantly thereafter, peaking on the seventh day. Some subsets of CD4 T cells continue to infiltrate the infarcted area for 7 days after acute ischemic stroke (Stubbe et al., 2013; Miró-Mur et al., 2020). Among the various subtypes of CD4 T cells, Th1 and Th17 cells have been demonstrated to increase in numbers within the first 3 days and decrease on day 7, whereas the opposite trend is observed for Th2 cells (Yu et al., 2022). CD8 T cells infiltrate the brain tissue within a few hours of stroke, and as the first invading lymphocyte, they significantly increase within the first 3 days after a stroke (Miró-Mur et al., 2020). In addition, only a small amount of Treg infiltration occurs during the acute phase of ischemic stroke (Shi et al., 2021), but Treg cell infiltration increases significantly from day 7 onward and can persist for more than 5 weeks (Stubbe et al., 2013; Xu et al., 2013). Studies have also shown that $\gamma\delta$ -T cells can be observed in ischemic brain tissue within 24 hours of stroke, and their secretion of the major effector molecule IL-17 peaks on the third day (Shichita et al., 2009; Dong et al., 2022).

Distinct functions of various T cells after ischemic stroke

It is being increasingly recognized that the interactions between ischemic stroke and T cells are complex and multifaceted, with different types of T cells playing distinct roles after ischemic stroke. These interactions have important implications for effective brain function during the disease and recovery process (Selvaraj and Stowe, 2017).

Th1 cells

A key feature of the Th1 cell response is the production of IFN- γ , which has been associated with a delayed immune response to stroke. Blockade of the IFN- γ signaling pathway has been shown to reduce neurodegeneration following middle cerebral artery occlusion and decrease the expression of the inflammatory chemokine IFN-inducible protein 10 (Seifert et al., 2014). IFN- γ is also a crucial activator of microglia, significantly enhancing their number and morphology (Kaya et al., 2022). As CD4 T cells are depleted over time after a stroke, IFN- γ levels decrease, ultimately leading to improved behavioral outcomes (Harris et al., 2020).

Th2 cells

The Th2 cell response is characterized by the release of various IL cytokines, such as IL-10. IL-10 exerts anti-inflammatory effects by downregulating various pathways, including NF- κ B in activated B cells. IL-10 has also been shown to reduce infarct volume and exhibit immunomodulatory effects in the CNS. Furthermore, evidence showed that BBB integrity improved in a cerebral ischemia/reperfusion model using hydrogen sulfide donors, and this was accompanied by enhanced IL-10 expression and reduced NF- κ B nuclear translocation (Wang et al., 2011).

Tregs

Tregs are considered major neuroprotective modulators in the nervous system (Zhang et al., 2021b). During the early phase of ischemic stroke, Tregs curb the systemic inflammatory immune response and the activation of peripheral effector T cells (Teffs). They also restore the integrity of the BBB and promote neuronal recovery. Additionally, Tregs influence the trafficking of peripheral immune cells, primarily by inhibiting the elevation of IL-6 and tumor necrosis factor- α (TNF- α) in the blood (Asseman et al., 1999; Qin et al., 2020).

During the later stages after ischemic stroke, Tregs trigger apoptosis in Teffs by promoting the secretion of granzymes and perforin (Gondek et al., 2005). Simultaneously, Tregs can reduce microglia/monocyte activation through the release of anti-inflammatory cytokines, such as IL-4, IL-10, and transforming growth factor- β (TGF- β), in the post-stroke environment (Liesz et al., 2009; Liu et al., 2020). Furthermore, Tregs can inhibit reactive oxygen species (ROS) synthesis and release from microglia and regulate IL-6 and STAT3 pathways in microglia and astrocytes via amphiphysin, thereby exerting neuroprotective effects (Ito et al., 2019). Crucially, Tregs are involved in immunomodulatory processes during ischemic stroke, expressing a range of immunosuppressive molecules, such as CD47 (also known as integrin-associated protein), CD39 (also called ectonucleoside triphosphate diphosphohydrolase-1), PD-1, and signal-regulated protein-alpha (Huang et al., 2017).

Lastly, it is important to note that, in Treg-cell-depleted stroke mice, the surface area and volume of microglia were reported to decrease, indicating that the presence of Treg cells can alter the activation status of microglia in the post-stroke brain. Experimental data have shown that genes associated with the anti-inflammatory phenotype, such as arginase-1, fibrinogen-like 2, mannose receptor 1, IL-1R antagonist, and advanced glycation end-product receptor 3, are significantly up-regulated in microglia co-cultured with activated Treg cells. Similarly, genes encoding proteins involved in brain repair, such as chemokine receptor 5, protein-glutamine gamma-glutamyltransferase 2, vascular endothelial growth factor A, fibroblast growth factor 1, and Gdnf, are also up-regulated. This suggests that Treg cells play a role in polarizing microglia toward an anti-inflammatory and repair phenotype, which can have potential therapeutic implications. Additionally, Treg cells and microglia create an osteopontin-rich microenvironment through their interactions. This environment can positively impact microglia responses, oligodendrocyte regeneration, white matter repair in the chronic phase of ischemic stroke, and long-term neurological function and outcomes after cerebral ischemic stroke (Shi et al., 2021).

$\gamma\delta$ T cells

The crucial role of $\gamma\delta$ T cells in brain tissue injury after ischemic stroke is not performed in isolation, but rather is primarily a result of their release of cytokines, including IL-17A, IL-21, IL-22, and IFN- γ . The primary source of IL-17A in stroke is $\gamma\delta$ T cells (Dong et al., 2022). IL-17a downregulates tight junctions (TJs) by inducing the production of ROS, leading to a decrease in TJ molecule expression. IL-17a also elevates matrix metalloproteinase-2, matrix metalloproteinase-3, and matrix metalloproteinase-9 levels in brain microvascular endothelial cells, hydrolyzing TJs and increasing BBB permeability. IL-17A promotes neutrophil infiltration, and the enhanced infiltration of neutrophils further disrupts the integrity of the BBB and promotes neuronal lysis and apoptosis, thereby exacerbating brain injury (Gu et al., 2015; Ruhnau et al., 2017). Additionally, IL-17A promotes the expression of apoptosis-associated proteins, including caspase-3,

caspase-9, and BCL2-associated X, and increases the BCL2-associated X/B-cell lymphoma-2 ratio after brain injury, leading to neuronal apoptosis (Li et al., 2017).

IL-21 is strongly up-regulated in the injured brain, and it significantly affects brain injury by upregulating autophagy-related genes in neuronal cells expressing IL-21R. This receptor has been found to exert neuroprotective effects through the Janus tyrosine kinase (JAK)/STAT signaling pathway and the upregulation of caspase-3 (Lee et al., 2016; Weiner and Ducruet, 2016).

IL-22 has protective effects against ischemic stroke, and experiments have shown that the exogenous administration of IL-22 reduces oxidative stress and neuronal apoptosis in affected brain tissue. A small subset of $\gamma\delta$ T cells can produce IFN- γ (Wang et al., 2023), which can act on both microglia and the CNS.

$\gamma\delta$ T cells play roles in the pathological process of ischemic stroke through their secreted cytokines, which have significant effects on brain tissue injury, neurological deficits, and functional improvement in later stages (Wang et al., 2022). In conclusion, T cells have critical functions in various aspects of ischemic stroke pathology, and the interactions between T cells and microglia, as well as the underlying mechanisms involved in stroke, require further exploration and investigation.

Central Immune Cells–Microglia

Introduction to microglia

Microglia are macrophages that reside within the brain and serve as the immune cells of the CNS. These cells remain in the brain for the duration of an individual's life, maintaining their numbers and spatial distribution through self-renewal (Huang et al., 2021). Microglia are recognized as the first non-neuronal cells to respond to various acute brain traumas and are the principal component of the brain parenchyma's defense network (Woodburn et al., 2021). Microglia are analogous to brain sentinels and are very dynamic APCs in the healthy brain (Woodburn et al., 2021). They vigilantly guard the brain, continuously detecting and tracking any disturbances in homeostasis (Borst et al., 2021). To quickly react to changes in the extracellular environment; to do this, mature microglia in the brain utilize a diverse range of surface molecules, allowing them to respond to hormones, neurotransmitters, chemokines, purines, and cytokines. Like other tissue-resident macrophages, microglia express common markers such as CD11b, the surface glycoproteins F4/80 and CD68, the fractalkine receptor CX3CR1, colony-stimulating factor 1 receptor, ionized calcium-binding adapter molecule 1, and the pan-hematopoietic cell marker CD45. However, at steady state, their expression levels are lower than those of perivascular macrophages and blood monocytes (Greter and Merad, 2013). Furthermore, the presence of any deleterious conditions leads to microglia activation, which results in distinct activation phenotypes. There are two main activation phenotypes of microglia: the classical M1 phenotype and the alternative M2 phenotype (Gao et al., 2021). The M2 phenotype primarily secretes anti-inflammatory factors that function to suppress inflammatory responses and promote tissue repair and functional remodeling. Conversely, the M1 phenotype plays a pro-inflammatory role by producing high levels of oxidative metabolites and pro-inflammatory cytokines to initiate a series of inflammatory responses (Lanza et al., 2021). These two phenotypes are not mutually exclusive; they can coexist side by side and transition between one another (Song and Suk, 2017).

Physiologic functions of microglia

Microglia, as specialized phagocytes of the brain (Mallat et al., 2005), play a pivotal phagocytic role in the normal development of the brain. They are responsible for the elimination of surplus synapses, dendrites, axons, myelin, neurons, and neuronal precursors (Schafer et al., 2012; Butler et al., 2021), and thus contribute to the execution of functions needed for a healthy brain and positively impact memory and learning. Furthermore, microglia have the ability to support neurons. Microglia are crucial components of the effective myelin regeneration response and can sustain and nourish neurons by releasing neurotrophic factors (Sen et al., 2022). In addition, they are involved in programmed neuronal death apoptosis, and the clearance of neurons during development (Long et al., 2023), and are crucial for the health and survival of neurons. Under physiological conditions, microglia can secrete trophic factors such as insulin-like growth factor 1, brain-derived neurotrophic factor, and nerve growth factor to provide nutrients to neurons and other cells.

Recent studies have demonstrated the critical roles of microglia in the formation and maturation of neural circuits, particularly during postnatal development (Wake et al., 2011; Paolicelli and Ferretti, 2017; Sun et al., 2023). In the healthy CNS, microglia are located in close proximity to neuronal synapses, maintaining constant contact with neurons and astrocytes. They can engage in direct interactions with presynaptic and postsynaptic components. During brain development, the establishment of neuronal synaptic structures is integral to the maturation of the CNS. Using transmission electron microscopy and two-photon microscopic imaging, Tremblay et al. (2019) revealed that microglia engage in contact with various sites during learning and perception, including dendritic spines, axon terminals, astrocyte protrusions, and synaptic clefts. Microglia phagocytose and prune synaptic components, contributing to synaptic remodeling during sensory processing (Gaire, 2022).

Functions of microglia in pathological conditions

Under pathological conditions, microglia become activated and stimulated to express various surface receptors, mainly scavenger receptors (SRs), e.g., SR class A types A (SRA-A), SR class B types B (SRB-B), CD36, CD68, and other SRs. These receptors are primarily involved in the phagocytosis of heterogeneous molecules on the surface of apoptotic cells, as well as the endocytosis of ligands associated with intercellular adhesion and cell debris. Neuronal injury triggers microglia to migrate to the site of injury by releasing adenosine triphosphate (ATP). The SR binds to ectopically translated phosphatidylserine, and phagocytosis of cellular debris by SRA-A promotes the phagocytosis of A β oligomers by microglia as well as astrocytes (Yang et al., 2011). When SRA-A is able to bind to amyloid- β (A β) and is activated, CD36 can mediate the phagocytosis of A β by inhibiting the prostaglandin E2 pathway (Silverstein and Febbraio, 2009).

Microglia play crucial roles in CNS injury responses. Recombinant purinergic receptor P2Y G-protein-coupled 12 (P2RY12) on microglia is involved in ATP sensing and regulates microglia recruitment following cell injury. Purinergic receptor P2Y13 may contribute to microglia activation and the response to adenosine diphosphate (ADP) sensing during inflammation (Fekete et al., 2018; Jairaman et al., 2022).

Additionally, microglia expressing MHC II molecules can cross-present antigens to regulate T cell responses within the CNS, dysregulation of both interactions may lead to altered phagocytosis of neuronal synapses by microglia, resulting in neurocognitive deficits (Chhatbar and Prinz, 2021).

Various roles of microglia after ischemic stroke

When an ischemic stroke occurs, microglia are activated and quickly migrate to the site of injury, where ischemia, hypoxia, and energy deprivation cause the branching of microglia populations into cells with a hypertrophic amoeba-like morphology. It is increasingly recognized that microglia play a diversity of roles depending on the stage of injury, interactions with other cells, pro- and anti-inflammatory conditions, and the composition of the surrounding cellular environment (Hu et al., 2014; Ma et al., 2017; Qin et al., 2019). In the acute phase, activated microglia secrete cytokines such as sphingosine-1-phosphate (S1P), high-mobility group box 1 protein, S-100 protein beta chain, and chemokine like factor-1. These cytokines bind to membrane receptors such as the cytokine receptors S1P receptor, advanced glycosylation end product-specific receptor, TLR4, IL-4R, and recombinant chemokine C-C motif receptor 4. This triggers pro-inflammatory cell signaling pathway activity via JAK2/STAT1/NF-κB, JAK2/STAT3/NF-κB, and myeloid differentiation factor 88/NF-κB. These signaling cascades initiate an intense inflammatory response (Jiang et al., 2020) and can also lead to endothelial necrosis and disruption of the BBB (Qin et al., 2019). Following the acute phase, the inflammatory response gradually subsides, and the infarcted area of ischemic stroke enters a new phase of

tissue healing (Hiu et al., 2016). Microglia contribute to the repair of infarcted tissue, the promotion of cerebral revascularization, and improvements in neurological functions in this later stage (Zhang et al., 2000; Liu et al., 2016), and the mechanisms primarily involve the phagocytosis of cellular debris and secretion of growth factors (De Sousa, 2022). **Figure 2** illustrates the effects of distinct subpopulations of T cells and microglia on neurons.

Interaction of T Cells with Microglia

Antigen presentation

Antigen processing and presentation play crucial roles in regulating the immune response. These mechanisms process foreign or intracellular antigens and present them to T cells for recognition, ultimately leading to their activation (Kawasaki et al., 2022).

Studies by Almolda et al. (2010, 2011) have shown that two key signals are involved in the antigen presentation mechanism. The first signal is provided by the binding of MHC molecules carrying antigens to TCRs on the surface of T lymphocytes. This interaction confers specificity to the mechanism, as MHC class I molecules are specifically recognized by CD8⁺ T lymphocytes, while MHC class II molecules bind exclusively to CD4⁺ T cells (Joffre et al.,

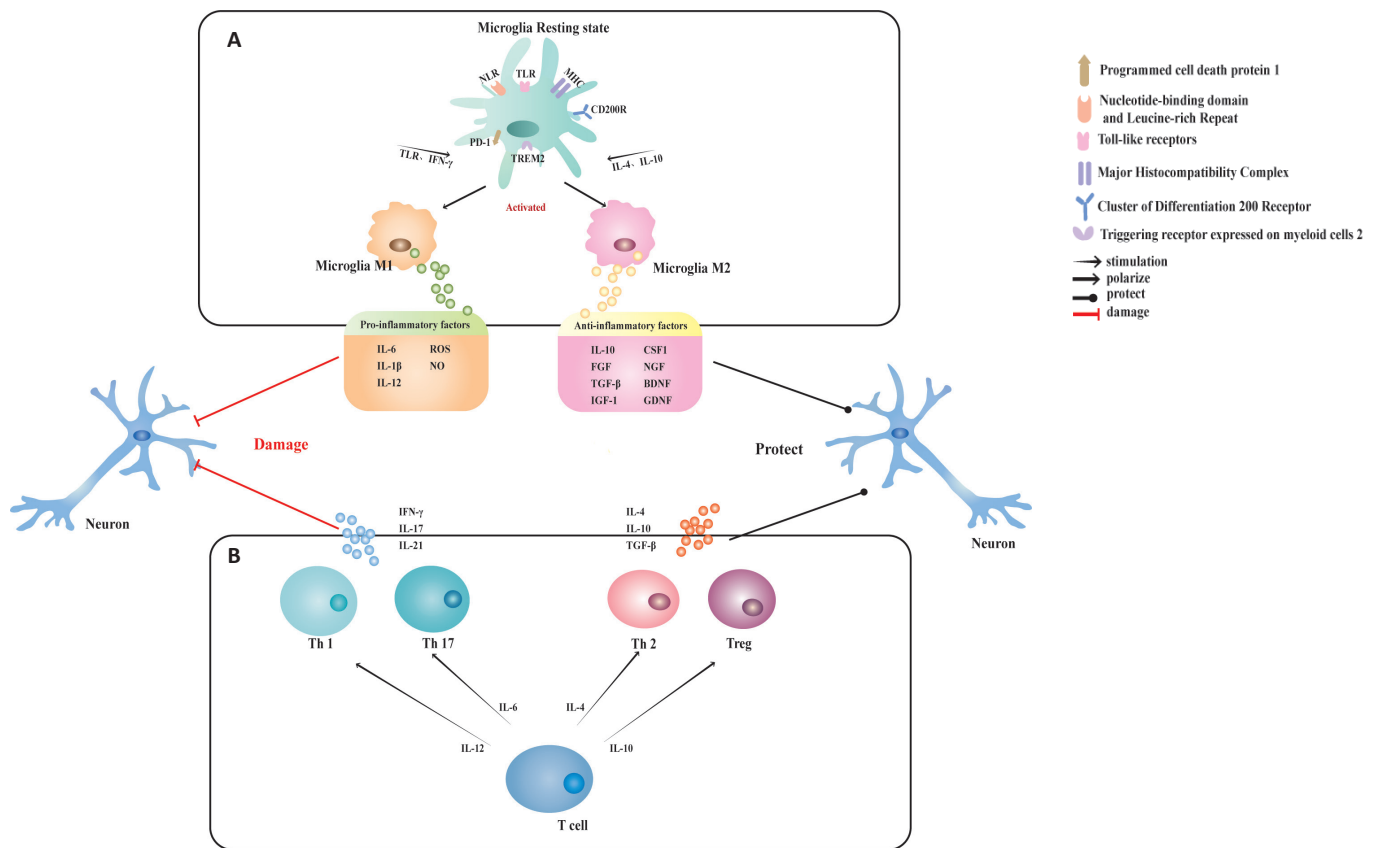


Figure 2 | Effects of different T cell and microglia subpopulations on neurons.

(A) Resting microglia are activated to diversify into two types by different stimuli. Stimulation with TLR and IFN-γ activates resting microglia to adopt the M1 type, while stimulation with IL-4 and IL-10 activates resting microglia to form the M2 type. M1-type microglia release various pro-inflammatory cytokines that can damage neural tissue, while M2-type microglia release anti-inflammatory cytokines and trophic factors that promote neural tissue repair. (B) T cells also differentiate into different subpopulations in response to different stimuli. Th1 and Th17 cells secrete pro-inflammatory factors to trigger severe inflammation, while Th2 and Treg cells secrete anti-inflammatory factors to suppress the onset of inflammation. BDNF: Brain-derived neurotrophic factor; CD200R: cluster of differentiation 200 receptor; CSF1: colony-stimulating factor 1; FGF: fibroblast growth factor; GDNF: glial cell line-derived neurotrophic factor; IFN-γ: interferon gamma; IGF-1: insulin-like growth factor 1; IL: interleukin; LRR: leucine-rich repeat; MHC: major histocompatibility complex; NGF: nerve growth factor; NLR: nucleotide-binding domain and leucine-rich repeat; NO: nitric oxide; PD1: programmed cell death protein 1; ROS: reactive oxygen species; TGF-β: transforming growth factor-β; Th: helper T cell; TLR: Toll-like receptors; Treg: regulatory T cells; TREM2: triggering receptor expressed on myeloid cells 2.

2012). Typically, MHC class I molecules present intracellular antigens, such as those produced by viruses, faulty self-molecules, or tumor-associated antigens. In contrast, MHC class II molecules are used to present antigens from extracellular pathogens like bacteria, parasites, and poisons (Elmer and McAllister, 2012). While MHC class I molecules are expressed by all nucleated cells, APCs, such as dendritic cells and macrophages, also possess the necessary processing and delivery mechanisms for presenting external antigens via MHC class II molecules (Almolada et al., 2011). The second signal, known as the co-stimulatory signal, is generated by the binding of different receptors expressed on the surface of APCs and lymphocytes to their respective co-receptors and is antigen-independent. The presence of these co-stimulatory signals is necessary for adequate T cell activation, as TCR binding to MHC without co-stimulation can lead to T cell apoptosis or incompetence (Lanzavecchia, 1997; Kishimoto and Sprent, 1999). Common co-stimulatory molecules include ICOS, CD154, and OX40 (also known as CD134), which bind to their corresponding ligands, B7h3 (also known as CD276), CD40, and OX40L, respectively, delivering stimulatory signals to lymphocytes and induce their activation and proliferation. In contrast, other co-stimulatory molecules, such as PD1, which binds to its receptor PD-L1, trigger inhibitory signals that lead to lymphocyte apoptosis (Nurieva et al., 2009).

Throughout the years, numerous investigations have established the fundamental roles of microglia in the CNS. When the brain is healthy, microglia exist in a dormant state, yet these resting cells are far from inert. Instead, they undergo significant and continuous structural changes, displaying remarkable motility. Microglia use the high motility conferred by cell protrusions to constantly observe and monitor the CNS microenvironment (Catalano et al., 2022). Microglia sense subtle changes in their environment through a variety of surface receptors. When the microenvironment undergoes molecular transformations, such as the appearance of lesions, neurotoxicity, or infection, microglia respond swiftly, transitioning from irregular patrol movements to purposeful movements toward the site of the lesion (Dheen et al., 2007; Hanisch and Kettenmann, 2007).

Infected and damaged cells secrete nucleotides, which microglia sense through their purinergic receptors, causing them to migrate toward the affected cells. Microglia's perception of damage-associated molecular patterns triggers their activation, enhancing their ability to detect pathogen-associated molecular patterns. At this point, microglia transform into fully functional APCs, significantly increasing MHC II expression and presenting antigens to T cells (Goddery et al., 2021). Microglia possess the unique ability to cross-present antigens, allowing MHC class I molecules to present extracellular antigens, thereby accelerating the immune response process (Moseman et al., 2020). Quantitative analysis of cells using flow cytometry revealed that T cells in the spleen exhibit a strong negative correlation with cells in the brain during the first 3 days of ischemic stroke, indicating there is a large influx of T cells from the spleen into the brain following the event (Liu and Sorooshyari, 2021). This finding also confirms the antigen-presenting role of microglia at the experimental level.

The regulation of antigen presentation by microglia is influenced by multiple factors. In the intact CNS, microglia are maintained in a quiescent state through local interactions between the microglial cell receptor CD200 and its ligand. Healthy neurons can suppress the expression of MHC molecules in microglia as a means to reduce inflammation (Neumann, 2001). In neurodegenerative diseases, microglia can promote the recruitment of CD8⁺ memory T cells to the aging CNS by releasing

C-C motif chemokine ligand 3 and upregulating the expression of vascular cell adhesion molecule and intercellular cell adhesion molecule in brain endothelial cells (Zhang et al., 2022). Both IL-6 and IFN- α contribute to the formation of reactive microglia in the core area and promote antigen presentation in animal models of neurodegenerative and neuroinflammatory diseases (West et al., 2022). During the early stages of neurological injury, activated microglia with a pro-inflammatory phenotype predominate and upregulate the expression of molecules involved in phagocytosis, oxidative damage, antigen presentation, and T cell stimulation. In later stages, microglia in active lesions shift to a phenotypic intermediate between pro-inflammatory and anti-inflammatory activation, and antigen presentation is greatly diminished (Zrzavy et al., 2017).

Secondary neurodegeneration is a chronic neuroinflammatory response to a traumatic brain injury event that occurs in areas distant from, but synaptically connected to, the primary site of damage (Stuckey et al., 2021). Current findings suggest that the inflammatory population at the site of secondary neurodegeneration is a mixture of cells, consisting of resident immune cells (microglia) and cells recruited from the circulation (Jones et al., 2018). Microglia play a key role in the stimulation of the leukocyte adhesion cascade response, which triggers the migration of T cells to the infarct zone (Jones et al., 2018). The secondary neurodegeneration site, like the primary infarcted area, is characterized by pronounced microglia activation. Activated microglia assist in the extravasation of immune cells to thalamic sites in secondary neurodegeneration at a later stage. However, unlike the primary infarct site, the secondary neurodegeneration site is not characterized by extensive and rapid BBB catabolism during stroke (Kluge et al., 2017). Therefore, infiltrating immune cells are not present at the secondary neurodegeneration site in the early stages after stroke. Fourteen days after ischemic stroke, there is a period of intense neuroinflammation at the site of secondary neurodegeneration (Jones et al., 2015). At this point, T cells infiltrate the secondary neurodegeneration site and become the main resident immune cells (Jones et al., 2018).

T cells interact with microglia to promote inflammation

The role of microglia as APCs is to promote the infiltration of T cells into the CNS and hence trigger inflammation. In addition to presenting antigens to T cells, microglia become activated after brain tissue injury and worsen neurological inflammation by secreting various cytokines that interact with subsets of T cells that release pro-inflammatory factors. As the main intrinsic immune cells in the brain, microglia are swiftly activated to take part in the immune response after ischemic stroke, when neurovascular units are damaged. Microglia in the ischemic brain can differentiate into two subpopulations according to the microenvironment: the classically activated microglia (M1 phenotype) and the alternatively activated microglia (M2 phenotype) (Zeng et al., 2022). Within 24 hours of cerebral ischemia, the activated M1 microglia develop in the ischemic brain tissue and commence secreting chemokines and inflammatory cytokines (Schilling et al., 2009; Gombault et al., 2012). These cytokines interact with subsets of Th1 and Th17 cells that infiltrate the CNS and, together, they produce high levels of pro-inflammatory cytokines, exacerbating the inflammatory response. Following ischemic injury, neurons in the hippocampus release zinc ions, which promote the M1-type polarization of microglia and the release of pro-inflammatory factors through the activation of P2X7 receptors (Higashi et al., 2017).

The activation and functional expression of microglia is associated with the involvement of their surface receptors, particularly TLRs,

which play pivotal roles in microglia-mediated inflammation. During focal cerebral ischemia, TLR2 and TLR4 engagement triggers microglia activation, leading to the production of pro-inflammatory cytokines and subsequent brain damage (Yenari et al., 2010). Numerous studies have consistently demonstrated an association between TLR4 levels and the severity of ischemia-induced neuronal injury (Yang et al., 2008; Hamanaka and Hara, 2011; Ye et al., 2019). Furthermore, TLR4 serves as a regulator of the interactions between M1 and Th1 cells and influences the immune response following ischemic stroke (Wang et al., 2011). TLR2 is also expressed in T cells, where it promotes Th1 cell function and participates in Th1-mediated responses (Imanishi et al., 2007). M1 microglia themselves produce a variety of pro-inflammatory cytokines and chemokines, including IL-1 β , IL-6, TNF- α , CCL2, and C-X-C motif chemokine ligand 10. These pro-inflammatory cytokines can, in turn, further promote the activation of M1 microglia (Yenari et al., 2010). Thus, Th1 cells can also facilitate the activation of M1 microglia by secreting pro-inflammatory factors. Notably, NF- κ B is an essential pathway for microglia polarization (Kou et al., 2022).

Pyroptosis, a form of neuroinflammation, is a programmed necrotic cell death process driven by inflammatory vesicles. This process is characterized by the disruption of cell membrane integrity, cell swelling, and the release of intracellular inflammatory contents, leading to an inflammatory response. The signaling pathway of cellular pyroptosis can be divided into the caspase-1-dependent classical pyroptosis pathway and the caspase-4/5/11-dependent nonclassical pyroptosis pathway. In the former, the high expression of marker molecules, such as Nod-like receptor protein 3 (NLRP3) and caspase-1 and its downstream response factor gasdermin D, is a key feature (Kovacs and Miao, 2017). Once activated, caspase-1 exhibits a protein-shearing function that processes inactive IL-1 β and IL-18 precursors into their mature forms. Additionally, activated caspase-1 cleaves the gasdermin D protein, leading to cell membrane perforation and the release of cellular contents, ultimately contributing to cellular pyroptosis (Shi et al., 2015). The NLRP3 inflammasome is activated by various factors in response to pathogen invasion, including ROS, K⁺ efflux, Ca²⁺ signaling, and mitochondrial malfunction. Additionally, the NF- κ B pathway, which is mediated by Toll-like receptors, can initiate the activation process (Wang et al., 2019). In a mouse model of cerebral ischemia, immunofluorescence co-localization experiments revealed that the NLRP3 inflammasome is predominantly expressed in microglia (Yang et al., 2014). Silencing of the NLRP3 gene reduced the release of pro-inflammatory cytokines, attenuating the inflammatory response and tissue damage. Furthermore, NLRP3 modulates the microglia's polarization state, promoting M1-type microglia activation and enhancing their pro-inflammatory capacity (Yan et al., 2023). An additional subset of human TH17 cells expresses gasdermin E, a membrane-pore-forming molecule. With the aid of the NLRP3 inflammasome–caspase-8–caspase-3 axis, IL-1 α is released through the cell membrane pore, triggering a series of pro-inflammatory responses (Chao et al., 2023).

The JAK/STAT signaling pathway is a crucial signal-transduction mechanism that can be influenced by various cytokines and plays a pivotal role in various biological processes, including cell proliferation, differentiation, apoptosis, immunity, and metabolic regulation (Dodington et al., 2018). It has been established that the JAK2/STAT3 signaling pathway is activated during cerebral ischemia, leading to microglia activation and neuronal apoptosis. Cellular experiments have confirmed that the secretion of TNF- α and IL-1 β by lipopolysaccharide-induced BV2 microglia

can be reduced by downregulating the phosphorylation of the JAK2/STAT3 pathway (Shrivastava et al., 2013). Additionally, downregulating the phosphorylation of the JAK2/STAT3 pathway promotes the polarization of microglia to the anti-inflammatory M2 phenotype, mitigates neuronal apoptosis and degeneration, and alleviates inflammatory responses and brain damage (Hristova et al., 2016). Previous studies have shown that ischemic stroke patients exhibit significantly increased Th17 cells and significantly decreased peripheral regulatory T (Treg) cells (Dolati et al., 2018; Shan et al., 2023), suggesting that an imbalance between peripheral Th17 and Treg cells may underlie the pathogenesis of ischemic stroke (Dolati et al., 2018). IL-17, secreted by Th17 cells, is the primary mediator of the delayed inflammatory response in ischemic stroke (Roy et al., 2005). Upon cerebral ischemia, microglia activate the TLR17/IL-2/IL-23 signaling pathway and secrete IL-17, leading to neuronal injury (Lv et al., 2011). Once Th17 cells enter the CNS, the local APCs, i.e., microglia, stimulate them again, and this causes microglia activation and IL-1, TNF- α , and IL-6 release (Murphy et al., 2010). IL-1 β , TNF- α , and IL-17 all activate the JAK2/STAT3 signaling pathway (Boscia et al., 2013; Liddelow et al., 2017; Li et al., 2018). More interestingly, the JAK/STAT pathway not only promotes the transcription and translation of inflammatory cytokines but also regulates the differentiation of T cells into Th17 cells in the spleen, thereby influencing the Th17/Treg balance in the peripheral blood and further inducing inflammatory injury (Chang et al., 2018).

The primary effector cytokine produced by Th1 cells with a pro-inflammatory phenotype is IFN- γ (Dardalhon et al., 2008), which is a precise promoter of M1 microglia activation. These two types of cells and the cytokines they secrete interact to form a closed loop that together promote the progressive development of the inflammatory response.

In a clinical trial, Fu et al. (2014) treated 11 patients with acute ischemic stroke using standard management in combination with fingolimod. The results of contrast-enhanced T1-weighted imaging (CET1-WI) showed a significant reduction in infarct zones and a significant decrease in microvascular permeability in those who received fingolimod compared with the 11 patients who only received standard management. This clinical study demonstrated that fingolimod, acting as an immunosuppressant, can prevent the infiltration of T cells into the brain parenchyma, thereby mitigating the inflammatory response of the nervous system and slowing down the progression of secondary injury after ischemic stroke (Fu et al., 2014). Therefore, fingolimod may be a promising alternative drug for the acute phase of ischemic stroke.

T cells interact with microglia to suppress inflammation

Early research on antigen presentation by microglia revealed that these cells are inefficient T cell activators, thus leading to tolerance or anergy (Ford et al., 1996). Secondary signals that effectively suppress T cell responses, often referred to as T cell "exhaustion," are elicited by PD-1:PD-L1 interactions. The characteristics of these exhausted phenotypes include decreased proliferative capability, cytokine production, and cytotoxic effector functions. In uninfected mice, PD-L1 was expressed at basal levels on approximately 20% of microglia; however, a week following viral brain infection, increased expression was seen on over 90% of cells (Schachtele et al., 2014). During neuroinflammation, microglia upregulate PD-L1, which binds to PD-1 molecules on the surface of T cells, thereby inhibiting the production of T cell pro-inflammatory factors. This suggests resident neuroglia have a role in limiting CNS pathology (Lokensgard et al., 2015).

Evidence has shown that microglia expressing PD-L1 after middle cerebral artery occlusion can inhibit the pro-inflammatory capacity of T cells and limit the extent of infarction, peripheral inflammatory cell migration, and experimental neurologic injury (Ren et al., 2011). This contrasts with the fundamental role of antigen presentation, which is to inform immune cells of the presence of a pathogen, thereby triggering an immune response. It would be inconsistent for microglia to present antigens to T cells but cause immune response suppression. If this were true, microglia would not be accurately described as APCs. However, subsequent research has revealed that microglia do not suppress the immune effects of all T cells; rather, they induce T cells to differentiate into two distinct subpopulations: pro-inflammatory and anti-inflammatory (Wang et al., 2016).

In post-stroke neuroinflammation, T cell subsets can have beneficial or harmful effects. In particular, it has been demonstrated that the pro-inflammatory Th1 and Th17 subsets of Th cells, as well as IL-17-producing T cells, cause secondary neurotoxicity, leading to infarct expansion and functional decline (Gelderblom et al., 2012). Conversely, Treg cells limit excessive inflammatory responses to the brain infarct by acting as anti-inflammatory and neuroprotective cells (Benakis et al., 2022). MHC class II (+) CD40 (dim) CD86 (dim) IL-10 (+) microglia have also been shown to be powerful inducers of Ag-specific CD4 (+) forkhead box protein P3 (+) *in vitro*. Microglia with low MHC II expression, co-stimulatory molecules, and high IL-10 expression promote Treg induction in a manner depending on the amount of IFN- γ challenge and the dose of Ag (Ebner et al., 2013). In contrast, Treg cells are key immunomodulators that protect the brain during acute ischemic stroke. They inhibit the overproduction of pro-inflammatory cytokines, regulate the invasion and/or activation of lymphocytes and microglia, and prevent the spread and growth of secondary infarct areas in the ischemic brain (Liesz et al., 2009). In the pathological process of ischemic stroke, the effect of microglia on the induction of T cells is not unidirectional. Single-cell sequencing has shown that, after an ischemic stroke, microglia may express their chemotactic properties in a T cell-dependent manner.

Distinct T cell subsets differently regulate the early activation of microglia after stroke. Th1 cells increase type I interferon signaling in microglia, driving them toward an antigen-immune functional phenotype, while regulatory T cells boost genes related to chemotaxis in microglia (Benakis et al., 2022). Tregs have been demonstrated to mediate the inhibition of M1-like microglial function by suppressing NF- κ B activation (Reynolds et al., 2009), which reduces migration, phagocytosis, and the generation of neurotoxic factors. For M2 microglia, the osteopontin derived from Treg cells acts through integrin β 1 (ITG β 1) receptors to enhance the repair activity of microglia (Shi et al., 2021). This promotes oligodendrocyte production and white matter regeneration, ultimately rescuing the neurological impairment caused by stroke. Gene set enrichment analysis showed that, after coculture with activated Treg cells, microglia had activated pathways associated with the type II immune response and tissue remodeling. Under Treg cell stimulation, microglia enhance the expression of genes and proteins that are key drivers of oligodendrogenesis, including vascular endothelial growth factor, transglutaminase 2, osteopontin, and arginase-1, proteins that boost the repair of ischemic brain tissue (Zera and Buckwalter, 2021).

Th1 cells release IFN- γ , IL-2, and TNF- β and express the transcription factor T-bet, and thereby promote cell-mediated

immune responses and phagocyte-dependent inflammation. In contrast, Th2 cells secrete IL-4, IL-10, and IL-13 and express the transcription factor GATA3. According to experimental results, the adoptive transfer of M2 cells reduced the number of CD4 T-bet Th1 cells and increased the number of CD4 GATA3 Th2 cells. These findings suggest that M2 microglia can shift the local microenvironment from a Th1-dominated state to a Th2-dominated state, thereby facilitating the inhibition of neuroinflammation.

Increasing evidence suggests that modulating microglia/macrophages through the CD47–SIRP α pathway alters phagocytosis. Signal regulatory protein α (SIRP α) is a typical inhibitory immunological receptor that is expressed on myeloid cells such as macrophages, granulocytes, and microglia (Kharitonov et al., 1997; van den Berg et al., 2005; Matozaki et al., 2009). At the same time, CD47 is expressed on Treg cells (Zhang et al., 2021a). Binding of CD47 to SIRP α immunoreceptor tyrosine-based inhibitory motifs within the cytoplasmic region of SIRP α , leading to the recruitment of Src homologous phosphatases, e.g., SH2-containing protein tyrosine phosphatase-1 (SHP-1), and SHP-2, in macrophages. In turn, these phosphatases inhibit the accumulation of myosin II at phagocytic synapses and inhibit phagocytosis by microglia as a means of attenuating the inflammatory response (Russ et al., 2018). An experimental study demonstrated that the Treg cell-derived exosome miR-709 can be delivered to microglia and plays a key role in inhibiting microglia focal death, the mechanism of which might be related to the negative regulation of NF- κ B-activating protein by miR-709 (Xiong et al., 2022).

In summary, M2 microglia and Treg cells demonstrate positive anti-inflammatory effects during the acute phase of ischemic stroke. Therefore, future scientific research and clinical trials should prioritize investigations into pro-polarization and the agonistic action of interactions between these two cell types. The mechanism of interaction between T cells and microglia is shown in **Figure 3**. The interactions between microglia and T cells are summarized in **Table 2**.

Conclusion

The mechanisms of ischemic stroke have been the subject of intense investigations in recent years. However, the complex pathogenesis of ischemic stroke makes this task particularly challenging. In this review, we have delved into the intricate role of the immune system in the pathophysiological changes that occur following the onset of ischemic stroke. The CNS and the immune system are intimately linked through a complex network of communication, with the immune system monitoring brain function and responding when homeostasis is disrupted by injury or disease. During the acute phase of ischemic stroke, there are interactions among the various immune cell types involving intricate mechanisms regulated by numerous factors. The triggering of inflammatory processes leads to the rupture of the BBB, allowing peripheral immune cells, particularly T cells, to enter the brain parenchyma and accumulate in the affected ischemic tissue. As these peripheral immune cells accumulate, microglia, pivotal immune cells of the brain, begin to activate in response to the ischemic event.

Upon activation, microglia release a range of pro-inflammatory cytokines that promote the expression of adhesion molecules on endothelial cells. This triggers the recruitment of additional leukocytes from the peripheral blood. T cells that infiltrate the brain release both pro- and anti-inflammatory factors,

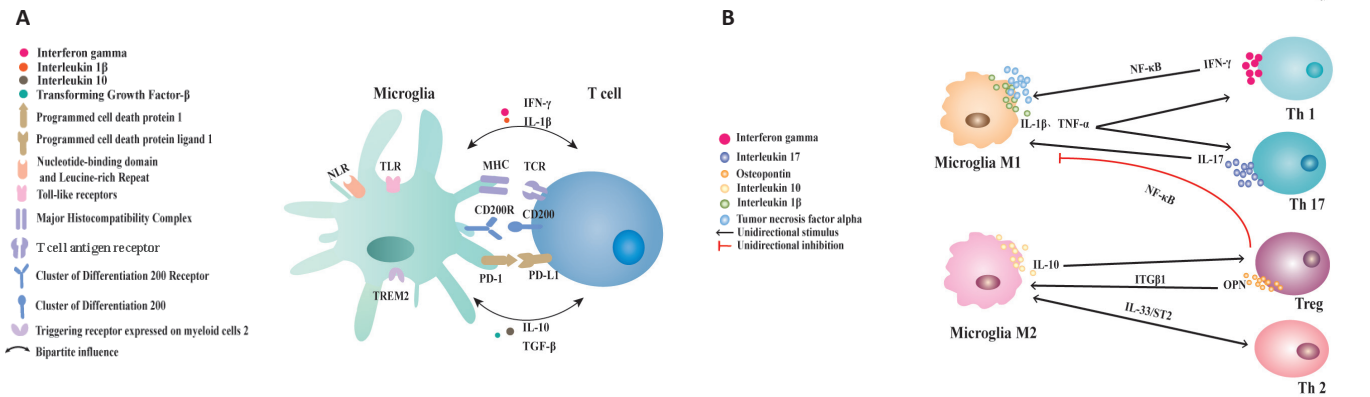


Figure 3 | Mechanisms of interaction between T cells and microglia.

(A) The MHC molecules expressed by microglia can be specifically recognized by T cell antigen receptors (TCRs) present on the surface of T lymphocytes. When PD1 binds to its receptor PD-L1, it initiates a suppressive signaling cascade in T cells, ultimately leading to lymphocyte apoptosis. Microglia maintain a quiescent state through local interactions with the CD200 receptor and its corresponding ligand, thus mitigating inflammation. (B) Microglia interact with T cells to regulate the homeostasis of inflammation by secreting various pro- and anti-inflammatory factors. Th1 cells promote the activation of M1 microglia by secreting pro-inflammatory factors, and NF-κB is an essential pathway for microglia polarization. IL-17 can be secreted by Th17 cells and activates microglia, while microglia also secrete IL-17 to re-stimulate Th17 cells. Upon activation, microglia release IL-1, TNF-α, and IL-6, which have the potential to promote the differentiation of Th1 and Th17 cells. Through the suppression of NF-κB activation, Tregs have been demonstrated to mediate the inhibition of M1-like microglia function. Treg cell-derived OPN acts through ITGB1 receptors to enhance the repair activity of microglia. T cells and microglia can achieve a suppressive effect on neuroinflammation after ischemic stroke through the IL-33/ST2 axis. Microglia overexpress IL-10, which promotes the activation of Treg. CD: Cluster of differentiation; IL: interleukin; INF-γ: interferon gamma; ITGB1: integrin β1; MHC: major histocompatibility complex; NF-κB: nuclear factor kappa-B; NLR: nucleotide-binding domain and leucine-rich repeat; OPN: osteopontin; PD1: programmed cell death protein 1; PDL1: programmed cell death protein ligand 1; ST2: growth stimulation expressed gene 2; TGF-β: transforming growth factor-β; Th: helper T cell; TLR: Toll-like receptors; TNF-α: tumor necrosis factor alpha; Treg: regulatory T cells; TREM2: triggering receptor expressed on myeloid cells 2.

Table 2 | Interactions between microglia and T cells

T cells	Interactions	References
Th1–M1	Th1-derived factors trigger pro-inflammatory (M1) gene expression in microglia, causing microglia to differentiate toward the M1 phenotype and inducing a pro-inflammatory response.	Prajeeth et al., 2014
	The Th1-derived effector molecule IFN-γ upregulates the expression of the gene encoding MerTK, a functional regulator of myelin phagocytosis, in microglia. Thus Th1 may be able to regulate microglia phagocytosis.	Prajeeth et al., 2018
	Th1 cell-derived IFNγ activates microglia to release IL-1β.	Watanabe et al., 2016
	M1 microglia secrete TNF-α, IL-6, IL-12, and IL23 to recruit and induce the differentiation of Th1 cells.	Murray and Wynn, 2011
Th17–M1	IL-17A secreted by Th17 cells can bind to IL-17AR on the surface of microglia, thereby promoting microglia activation.	Huo et al., 2019
Th2–M2	Excessive transfer of M2 microglia can change the local microenvironment from being dominated primarily by Th1 cells to being dominated by Th2 cells.	Ma et al., 2015
	IL-33 receptors are expressed on microglia, and IL-33 induces protection by promoting IL-4 secretion from T cells, driving Th2-type metastasis in the periphery.	Korhonen et al., 2015
Treg–M2	Treg cell-derived osteopontin acts through integrin receptors on microglia to enhance microglia repair activity, promoting oligodendrocyte production and white matter repair.	Shi et al., 2021
	Microglia with high expression of IL-10 can promote the induction of Treg differentiation.	Ebner et al., 2013
	The Treg cell-derived exosome miR-709 inhibits microglia death.	Xiong et al., 2022

IFN-γ: Interferon gamma; IL: interleukin; IL-17AR: interleukin 17A receptor; MerTK: tyrosine-protein kinase Mer; miR-709: mmu-miR-709; Th: helper T cell; TNF-α: tumor necrosis factor-α; Treg: regulatory T cells.

influencing microglia to differentiate into either the pro-inflammatory phenotype M1 or the anti-inflammatory phenotype M2. M1 microglia interact with Th1 and Th17 cells, secreting pro-inflammatory factors that amplify inflammatory responses. Conversely, M2 microglia interact with Th2 and Treg cells, secreting anti-inflammatory factors that mitigate the inflammatory response. A recent study has highlighted the crucial role of neuroinflammation in determining the prognosis of stroke (Tao et al., 2020). The research showed that suppressing the inflammatory response after acute stroke can prevent brain damage and improve long-term neurological outcomes. The interaction between microglia and T cells affects the direction of neuroinflammation progression, which in turn determines the prognosis of ischemic stroke. This suggests that exploring the mechanisms of microglia-T cell interactions may offer novel therapeutic targets for the treatment of ischemic stroke in the future.

The mechanisms of crosstalk between T cells and microglia are complex, and we have consulted both experimental and clinically validated articles to gain a deeper understanding. Currently, the most well-defined mechanism of interaction between T cells and microglia is the release of inflammatory cytokines. To some extent, there is a synergistic effect between T cells and microglia. Th1 cells secrete IFN-γ, which promotes the polarization of M1 microglia, while the TNF-α secreted by M1 microglia can induce the differentiation of Th1 cells. Together, Th1 and M1 microglia produce high levels of pro-inflammatory cytokines, iNOS, and neurotoxic substances that promote inflammation and brain injury following ischemic stroke. During cerebral ischemia, Th2 cells secrete IL-4 and IL-10, while Tregs secrete IL-10 and TGF-β. These cytokines promote the activation of M2 microglia and protect the brain from damage. M2 microglia can also secrete IL-10 and TGF-β, which induce the differentiation of Th2 and Treg cells. Th2, Treg, and M2 microglia work together to produce high levels of anti-inflammatory cytokines and growth

factors, limiting secondary inflammation-mediated damage and promoting CNS repair. Therefore, in the treatment of ischemic stroke, the targeted regulation of inflammatory factors may play a moderating role in the interaction between the central and peripheral immune systems. This modulation can regulate the developmental direction of neuroinflammation and promote the repair of brain injury. In addition, T cells and microglia can also interact through certain signaling molecules. The surface receptor TLR4 on T cells triggers the downstream NF- κ B signaling pathway, which is essential for the activation of M1 microglia. After cerebral ischemia, microglia activate the TLR17/IL-2/IL-23 signaling pathway, leading to the secretion of IL-17 and subsequent neuronal damage. IL-1 β and TNF- α secreted by microglia activate the JAK2/STAT3 pathway, promoting Th17 cell differentiation and intensifying brain injury. Th2-derived IL-4 triggers the polarization of M2 microglia through the IL-4 Ra/JAK/STAT signaling pathway. In the future, exploring targeted drugs that specifically block these key signaling pathways could offer novel therapeutic approaches. In both clinical and experimental settings, various scholars have investigated drugs that can influence the immune roles played by microglia and T cells in ischemic stroke.

Fu et al. (2014) treated 11 patients with acute ischemic stroke using standard therapy plus fingolimod. By employing CET1-WI, they observed a significant reduction in infarct zones and microvascular permeability in patients who received fingolimod compared to those who received standard therapy alone. This clinical finding suggests that fingolimod, an immunosuppressant, effectively blocks T cell infiltration into the brain parenchyma. This attenuation of the inflammatory response in the nervous system may slow down the progression of secondary injury related to post-ischemic stroke. Consequently, fingolimod could serve as a potential therapeutic agent in the acute phase of ischemic stroke. In an animal model of middle cerebral artery occlusion, dimethyl fumarate administration was associated with the death of fewer neurons in the brain, reduced microglia activation, and the migration of fewer immune cells to the affected region. This led to lower systemic levels of immune cytokines, indicating that dimethyl fumarate may reduce brain inflammation through both direct and indirect mechanisms (Kunze et al., 2015). These findings suggest that administering dimethyl fumarate can lead to the reduced expression of various cytokines or chemokines. For instance, IL-17A, secreted by Th17 cells, promotes the production of inflammatory cytokines and chemokines, leading to microglia recruitment and activation (Lin et al., 2016). Additionally, dimethyl fumarate mitigates plasma levels of IL-12 and brain levels of IL-12p40. IL-12 is secreted by activated microglia and promotes inflammatory activity in ischemic stroke. Li et al. (2013) infused purified Treg cells intravenously into a mouse middle cerebral artery occlusion model of cerebral ischemia, and the number of Treg cells in the peripheral blood and peripheral immune organs increased. These cells, which survived for at least 12 days, significantly reduced the levels of inflammatory cytokines IL-6 and TNF- β induced by cerebral ischemia and improved cellular immune function after ischemic stroke.

Microglia, representative cells of central immune responses, and T cells, which typify peripheral immune responses, play pivotal roles in the prognosis of ischemic stroke. From a broad perspective, interactions between the central and peripheral immune systems hold the key to the effective treatment of ischemic stroke. These interactions, however, are complex, and while we have partially unraveled the mechanisms and verified their functions, a full explanation of the interplay between these two cell types remain elusive. At the clinical trial level, there

are also gaps in our knowledge, as there are a lack of targeted treatment options. It is our hope that, in the future, we can identify practical and effective drug targets that can regulate the interactions between T cells and microglia. By promoting their anti-inflammatory functions, reducing neuroinflammation, and fostering nerve regeneration and repair following ischemic stroke, we can pave the way for more effective treatment options.

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