

Iron in multiple sclerosis – from pathophysiology to disease progression – a narrative literature review

Karolina Kłodnicka¹, Jacek Januszewski^{2*}, Alicja Forma¹, Weronika Pająk¹,
Barbara Teresińska¹, Jacek Baj²

¹ Department of Forensic Medicine, Medical University of Lublin, Lublin, Poland

² Department of Normal, Clinical and Imaging Anatomy, Medical University of Lublin, Lublin, Poland

* Email: jacek.januszewski000@gmail.com

Multiple sclerosis (MS) is a chronic autoimmune illness characterized by demyelination and neurodegeneration, which causes physical disability and severe alterations in the neurological system, including gliosis and neuron loss. The disease primarily affects myelinated parts of the central nervous system (CNS), such as the optic nerves, cerebellum, brain stem, and spinal cord. T cells play an important role in MS pathogenesis by inducing demyelination, and risk factors include genetic predisposition, environmental effects, and lifestyle decisions. The prevalence of MS is rising, especially among women and the elderly population. Iron dysregulation is a critical element in MS pathogenesis, with excess iron causing neurodegeneration *via* ferroptosis and immune response modulation. Excess iron amplifies inflammation by triggering the activation of macrophages with inflammatory properties, and promoting microglial polarization toward the pro-inflammatory phenotype. This causes increased oxidative stress, mitochondrial malfunction, and the release of reactive oxygen species, which harm neurons. Furthermore, proinflammatory cytokines like IL-6 regulate iron metabolism and encourage the formation of Th17 cells, which exacerbates CNS inflammation. Macrophages and microglia, which are implicated in inflammatory responses, collect iron during MS, exacerbating neuroinflammation and demyelination. Disrupted iron homeostasis is a major contributor to MS pathology, with iron deficiency affecting immunological function and changing T-cell responses, both of which are necessary for disease progression. Lumbar puncture, oligoclonal bands analysis, and magnetic resonance imaging are all used to diagnose MS and confirm disease activity and progression. The blood-brain barrier is frequently disrupted in MS, allowing the influx of inflammatory cells. The aim of this paper is to demonstrate the cause-effect relationship between the amount of iron and the health status of patients with MS.

Key words: demyelination, iron, macrophages, microglia, multiple sclerosis, neurodegeneration, neuroinflammation, oxidative stress, reactive oxygen species

INTRODUCTION

The leading cause of nontraumatic neurological dysfunction in young adults is multiple sclerosis (MS). MS is an autoimmune disease of demyelination and neurodegeneration. In addition to its serious impact on physical disability, this disease also causes numerous changes affecting the nervous system, such as gliosis and neuronal loss in the central nervous system (CNS) (Hauser & Cree, 2020; Travers et al., 2022; Haki et al., 2024). Structures of the nervous system, such as the optic nerves, cerebellum, brain stem, and spinal

cord, are among the heavily myelinated areas of the nervous system that are particularly affected by MS, due to the progressive process of destruction of the myelinated regions (Dhanapalaratnam et al., 2022). Of particular importance in this regard are immunogenic T cells responsible for demyelination, resulting in the destruction of myelinated axons of the CNS (Haki et al., 2024). As a result, some patients experience sensorimotor symptoms such as tremors and vision loss, which are the most noticeable clinical symptoms (Makhoul et al., 2020; Stoiloudis et al., 2022). The first symptoms of MS usually appear between the ages

of 20 and 40. However, a growing number of studies show that the epidemiology is changing, with older people experiencing higher incidence and prevalence. Numerous statistics show that MS is becoming more common and widespread worldwide, especially among women. Multiple studies have shown that girls are more likely to develop MS in childhood than boys (Gonzalez-Lorenzo et al., 2024; Zhang et al., 2024; García López et al., 2024; Fernández et al., 2024).

It is estimated that 2.9 million people worldwide suffer from this disease. The annual incidence is about 5–6 per 100,000 people, and the prevalence in the northern regions of North America and Europe is about 0.1%–0.2% of the population. This indicates that the population of people suffering from MS is constantly growing (Valadkeviciene et al., 2019; Sandesjö et al., 2024). Important risk factors for the development of MS include genetic predisposition, environmental variables, lifestyle factors and their interactions (Olsson et al., 2017). The onset of MS is also believed to be influenced by factors like smoking, sunshine, and viral infections (Holz et al., 2024). Up to 30% of monozygotic twins have MS, and 12.6% of all MS patients have familial instances of the disease (Sadovnick et al., 1993).

Genetic predisposition plays a crucial role in the development of MS, with various genes influencing an individual's susceptibility to the disease. The *HLA-DRB1* gene, located on chromosome 6, is the most strongly associated with MS, as certain variants of this gene are believed to increase the likelihood of the immune system attacking the body's own tissues, a key characteristic of MS (Brynedal et al., 2007). Additionally, other genes involved in immune regulation, inflammation and cellular repair mechanisms, such as the *IL7R* gene, contribute to a polygenic risk model, where multiple genetic variations collectively influence the overall risk of developing MS (Zuvich et al., 2010). Among the inflammasomes, the nucleotide-binding oligomerization domain-, leucine-rich repeat- and pyrin domain-containing 3 (NLRP3) inflammasome, which contains the nucleotide-binding oligomerization domain, leucine-rich repeat, and pyrin domain, is well-characterized and contributes to several neurological diseases, including MS (Cui et al., 2022). It is suggested that the NLRP3 inflammasome, a protein complex responsible for activating the inflammatory response, and the cytokine IL-1 β , one of the main mediators of the inflammatory process, may be involved in how patients respond to interferon beta treatment (Cui et al., 2022). In practice, this means that these molecules may be key in regulating the inflammatory response in MS patients, and their activity may affect the effectiveness of the

treatment (Malhotra et al., 2015). Ethnic differences have been noted to influence the study results, with a stronger association observed in the Latin American population. The frequency of NLRP3 polymorphisms may vary by ethnic group (Wu et al., 2021).

Pathophysiology of multiple sclerosis

Hemoglobin

The WHO's 1968 definition of anemia's cut-off hemoglobin values applies to all adults (hemoglobin <12g/dL for women and hemoglobin <13g/dL for men), but it does not specifically address the elderly population (Otálora-Alcaraz et al., 2025). Hemoglobin is a protein composed of four subunits: two alpha and two beta subunits, each containing a heme group attached to an iron atom (Pires et al., 2023). In the serum, both heme and non-heme forms of iron are present. Heme is released during hemolysis, utilized in metabolic processes, and found in high concentrations within erythrocytes. To prevent the harmful pro-oxidant effects of free heme, the blood protein haptoglobin binds to the released hemoglobin and its iron (Schaer et al., 2014). Hemoglobin can cross the compromised blood-brain barrier (BBB) to reach the CNS. In progressive diseases, the breakdown products of hemoglobin, such as iron and heme, contribute to neurodegeneration and brain atrophy (Lewin et al., 2016).

Haptoglobin

Due to the inflammatory nature of certain diseases, various inflammatory markers, including the proinflammatory cytokine IL-6, have been linked to disease symptoms and pathophysiology (Yan et al., 2012). IL-6, produced by hepatocytes and white adipose tissue, induces the expression of haptoglobin (Li et al., 2020). This acute-phase protein binds tightly to hemoglobin released during red blood cell turnover, reducing iron loss, preventing renal injury, and mitigating heme-induced inflammation (Sarpong-Kumankomah et al., 2022). In multiple sclerosis patients, haptoglobin also plays a protective role by acting as a first line of defense against oxidative damage to myelin. This is especially important given the high fragility of erythrocytes in MS, which leads to increased hemoglobin release, alters the BBB, and damages myelin proteins (Bamm & Harauz, 2014; Altinoz et al., 2016). The BBB, along with the blood-cerebrospinal fluid (CSF) barrier, restricts iron transport from the

blood to the brain (Engelhardt & Sorokin, 2009). Iron is believed to enter the brain through transferrin receptor-mediated endocytosis in brain capillaries and is released back into circulation via the CSF (Moos et al., 2006). Elevated iron levels are implicated in protein aggregation in neurodegenerative diseases, further complicating the pathophysiology of MS (Harri-son & Arosio, 1996).

Ferritin

Ferritin serves as a major iron storage protein with the capacity to store up to 4,500 iron atoms within its hollow spherical structure, composed of 24 heavy and light chains (Wilkinson & Pantopoulos, 2014). Ferritin mRNA expression is regulated by the iron-responsive element (IRE)-iron regulatory protein (IRP) system, which responds to cellular iron levels (Arosio et al., 2017). In pathological conditions, ferritin levels increase significantly as iron accumulates within cells (Jhelum & David, 2022). Iron histochemistry can detect CNS iron deposition, and ferritin is a sensitive marker of iron presence (Jhelum et al., 2023). While ferritin itself is inert, it can become harmful when redox-active agents mobilize iron through ferritino-phagy (Santana-Codina & Mancias, 2018). The protein nuclear receptor coactivator 4 (NCOA4) facilitates the transport of ferritin to autophagosomes for degradation. Overexpression of NCOA4 in pathological contexts can result in the release of redox-active iron, which is cytotoxic (David et al., 2023).

Ferroptotic mechanisms

Ferroptosis is an iron-dependent, controlled form of cell death that occurs in MS (Santana-Codina & Mancias, 2018). This process is characterized by mitochondrial shrinkage, iron accumulation, severe lipid peroxidation, and increased membrane density without apparent rupture. Notably, the nucleus remains normal, with no chromatin condensation (Li et al., 2020a; Tang & Kroemer, 2020). Ferroptosis is triggered by oxidative changes in the intracellular environment (Yu et al., 2017). Excessive lipid peroxidation is considered the key mechanism of ferroptotic cell death, distinguishing it from other forms of regulated cell death (Wiernicki et al., 2020). However, both iron overload and a deficiency in the glutathione (GSH) pathway contribute equally to its initiation (Jhelum et al., 2023). Iron plays a critical role in maintaining healthy oligodendrocytes and myelin and may be essential for remyelination (Cheli et al., 2020). In MS,

iron levels are dysregulated, with an accumulation of iron in gray matter (GM) and a reduction in white matter (WM) (Hametner et al., 2018). After oligodendrocyte death, iron is released and can accumulate in neurons, potentially leading to neurotoxicity (Salva-dor et al., 2010; Levi et al., 2024). Excess iron, typically stored in ferritin, can be mobilized to generate free radicals, exacerbating oxidative damage. This dysregulation of cellular iron metabolism results in elevated bioactive iron, which drives the process of ferroptosis. Ferroptosis occurs when iron-mediated free radicals promote lipid peroxidation, especially in the presence of insufficient GSH-mediated antioxidant defenses (Fan et al., 2022; Wang et al., 2024). Polyunsaturated fatty acids in cell membranes are particularly susceptible to oxidation under these conditions (Ayala et al., 2014). In MS patients, brain tissue shows greater damage to myelin lipids than myelin proteins (Barnes-Vélez et al., 2022; Kister & Kister, 2023). A reduction in GSH, an important antioxidant, leads to the accumulation of lipid radicals and a loss of cell viability. This depletion of GSH in MS patients may also result in a decrease in glutathione peroxidase 4 (GPX4), an enzyme crucial for neutralizing lipid peroxides (Hu et al., 2019). Ferroptosis is exacerbated by chronic oxidative stress. In MS lesions, there is significant immune cell infiltration, which increases immune cell presence in the brain tissue compared to normal brain tissue. Higher ferroptosis scores have been correlated with phagocytic activation in WM lesions (Wu et al., 2024). Iron and lipid peroxides are central to the initiation of ferroptosis, with iron acting as a catalytic regulator. Ferroptosis can be inhibited by iron chelators like deferiprone or lipophilic antioxidants such as α -tocopherol. Additionally, reactive oxygen species (ROS) generated by the iron-catalyzed Fenton reaction contribute to ferroptosis onset (Feng et al., 2023).

MS diagnosis

Lumbar puncture and oligoclonal bands

A lumbar puncture is a key diagnostic test for MS (Ford, 2020). It involves drawing a sample of CSF from the subarachnoid space in the lumbar region to assess abnormalities linked to the disease (Gomes, 2022). One of the primary markers examined during this procedure is the presence of oligoclonal IgG bands (OCB), which suggest local antibody production in the CNS, a hallmark of MS-associated inflammation (Hümmert et al., 2019). Detection of OCB is strongly indicative of MS, as it is present in approximately 90%

of MS patients, although it is not exclusive to MS and can appear in other inflammatory disorders (Katsarogiannis et al., 2023). The revised McDonald criteria include OCB detection in CSF as a standard step in diagnosing MS (Kim, 2022). While OCB is not required for a definitive diagnosis, its presence, especially in combination with other diagnostic tools such as magnetic resonance imaging (MRI), provides crucial supporting evidence for MS (Joseph et al., 2009). Several types of OCB exist, with the most common being type 2 and occasionally type 3 (Sánchez-Vera et al., 2023). However, it is important to note that the presence of OCB in the CSF does not rule out non-inflammatory neurological diseases. Some neurological conditions, even those that are non-inflammatory, may also show OCBs, emphasizing the need for a comprehensive diagnostic approach (Pannewitz-Makaj et al., 2020). Despite its high diagnostic value, a lumbar puncture is an invasive procedure that carries potential risks, which must be considered before proceeding with the test (Graner et al., 2020).

Free kappa chains

Free kappa chains (κ -FLC), proteins produced by plasma cells and components of immunoglobulins (Rao et al., 2012), are another sign used to diagnose MS. Determining κ -FLC levels in CSF can reveal inflammatory activity in the CNS (Di Filippo et al., 2024). Elevated amounts of these chains in MS patients may indicate the existence of locally generated antibodies and disease activity. Recently, κ -FLC has emerged as a biomarker for disease monitoring and chronic inflammation assessment (Gudowska-Sawczuk & Mroczko, 2023). κ -FLCs, like other proteins, can be taken from blood or generated in the subarachnoid space under pathological conditions, which is especially important in MS (Hegen et al., 2022). Testing for κ -FLC using nephelometry or turbidimetry is simple, accurate, and cost-effective. Elevated κ -FLC levels in CSF can predict disease progression to clinical MS (Arrambide et al., 2022).

MRI in the diagnosis of multiple sclerosis

In the brain, iron primarily builds up in certain areas, including the striatum (Ward et al., 2014), thalamus (Rodrigue et al., 2020), and caudate nucleus (Schipper, 2012). This metal is crucial for numerous biological functions in certain brain regions, such as metabolism and neurotransmitter activity. The protein ferritin is the primary form of iron storage in the brain, and as people age, its concentration rises

(Beard, 2001). The globus pallidus, caudate nucleus, and substantia nigra are the basal nuclei with the largest concentrations of iron, which are implicated in the production of neurotransmitters, including glutamate and dopamine (Jamwal & Kumar, 2019). Other brain regions, particularly those impacted by myelin loss, may also experience iron buildup. Iron builds up in damaged places, particularly in the WM and demyelination-affected areas, in conditions like MS (LeVine et al., 2013). When myelin and oligodendrocytes are destroyed, activated microglia, involved in the brain's immune system accumulate iron, which can cause long-term inflammation and neurotoxicity (Yong, 2022). The placement of these alterations in deep brain nuclei is a sign of inflammatory activation, and they are frequently discernible on MRI scans, including T2-weighted sequences, where hyperintensity is apparent (Cacciaguerra et al., 2022). The main diagnostic criteria for MS based on MRI imaging are as follows:

- **Temporal Dissemination:** Lesions must be present at different points in time. This means that MRI changes should be observed at various stages during the disease, reflecting the relapsing-remitting nature of MS (Brownlee et al., 2025).
- **Spatial Dissemination:** Lesions should appear in at least two different locations in the CNS. Commonly affected areas include the periventricular regions, brainstem, cerebellum, and spinal cord (Filippi et al., 2019).
- **Characteristic Demyelinating Lesions:** The primary lesions in MS are demyelinating plaques, which typically show as bright spots on T2 or FLAIR MRI sequences. These lesions are most commonly seen on T2-weighted or Fluid-Attenuated Inversion Recovery (FLAIR) images. Active lesions may also be visible on contrast-enhanced T1 images, indicating inflammation or edema (Tillema & Pirko, 2013).
- **Presence of New Active Lesions:** New demyelinating lesions should be visible on MRI, ideally enhanced with contrast, indicating ongoing activity and inflammation (Kaunzner & Gauthier, 2017).
- **Exclusion of Other Causes:** The MRI should reveal changes consistent with MS and not be attributed to other neurological conditions (e.g., infections, tumors, or other demyelinating disorders) (Pirko & Noseworthy, 2007).

To diagnose MS, the McDonald criteria must be met. These criteria consider both spatial and temporal dissemination of lesions in the CNS and may involve the presence of new lesions on MRI, suggesting disease progression (Marcus & Waubant, 2013).

Table 1. Summary of possible methods of MS diagnosis.

Topic	Description
Lumbar Puncture	A diagnostic test for MS involving the collection of CSF from the subarachnoid space (Gomes, 2022). The presence of oligoclonal IgG bands, which indicate inflammation in the CNS, is expected (Katsarogiannis et al., 2023). Over 90% of MS patients present these bands (Hümmert et al., 2019).
Oligoclonal Bands (OCB)	OCB indicate the local production of antibodies in the CNS, which is characteristic of MS (Kim, 2022). The presence of OCB is a key indicator for MS diagnosis, especially when other test results are inconclusive (Kim, 2022). OCB is currently regarded as the gold standard for diagnosing MS (Pannewitz-Makaj et al., 2020).
Free Kappa Chains (κ-FLC)	Free kappa chains are proteins produced by plasma cells, with elevated levels in CSF indicating inflammatory activity in the CNS (Di Filippo et al., 2024). An increase in κ-FLC levels can predict disease progression and is used to monitor MS (Arrambide et al., 2022).
MRI in MS Diagnosis	Fe accumulates in the brain, especially in the basal nuclei, which may indicate damage caused by myelin loss in MS (Jamwal & Kumar, 2019). These changes are visible on MRI scans, particularly on T2-weighted sequences, where hyperintensity is observed (Cacciaguerra et al., 2022). Iron accumulation is associated with inflammatory activation and neurotoxicity (Yong, 2022).

Iron deficiency and weakened immune response

Iron deficiency can be caused by a low-iron diet, or a diet high in substances that inhibit iron absorption (Zimmermann & Hurrell, 2007). Healthy persons in high-income environments typically have serum iron levels ranging from 10–30 $\mu\text{mol/l}$ and show minimal variation with age (Garcia-Casal et al., 2021). However, systemic iron availability may be “functionally” low due to chronically high hepcidin levels in the context of inflammation, which limit iron absorption by the duodenum and serum iron availability, regardless of ferritin iron storage (Cianetti et al., 2010). Iron is thought to be essential not only for DNA synthesis and RBC creation but also for immune system health (Miller, 2013; Roemhild et al., 2021; AlRajeh et al., 2022). The findings revealed that iron-deficient patients with low hemoglobin and/or ferritin levels had a significantly higher risk of recurrent infections. Clinical studies have revealed that iron availability plays a critical role in the development of T and B cell responses to infection and vaccination (Muchowska et al., 2019). They discovered that iron deficit patients had an increase in T-cell proliferation. Increased iron intake is required to fuel T-cell responses, which is consistent with iron’s many roles in cellular metabolism (Andreini et al., 2018; Li et al., 2021). In an *in vitro* model of Th17 polarization, iron shortage affects the elimination of the main inhibitory histone methylation mark as well as differentiation (Teh et al., 2021). We found iron-interacting proteins in CD4⁺ and CD8⁺ T-cell proteomes that were differentially expressed during activation, implying that iron deficit may hamper pathways enriched for such proteins, such as histone demethylation. Iron requirements in T cells are immediately and significantly increased upon ac-

tivation for usage in a variety of cellular processes, including demethylation and DNA synthesis (Jabara et al., 2016). *In vitro* studies of T-cells exposed to iron-depleted media or iron chelators show that iron is crucial for cell proliferation, activation, and production of effector molecules like granulocyte-macrophage colony-stimulating factor (GM-CSF) and interferon- γ (IFN- γ) (Leung et al., 2005; Yarosz et al., 2020). Hepcidin, an acute-phase protein, binds to ferroprotein in enterocytes, liver, and splenic macrophages and then degrades. Overall, its activity lowers blood iron levels by retaining iron in macrophages and inhibiting its absorption from the gastrointestinal tract. It can be deduced that hepcidin overexpression causes iron deficiency anemia, whereas low expression causes iron overload (Collins et al., 2008; Kowdley et al., 2021). Hepcidin has been discovered in mice to decrease antigen-specific CD8 T-cell responses, as well as T follicular helper cell, germinal center B cell, and plasma cell responses in the ovalbumin vaccination paradigm. Mice treated with minihepcidin have impaired T-cell ability to produce interferon- γ and TNF. In mice with a mutation in the gene producing the iron transporter transferrin receptor, the investigators discovered a deficiency in lymphocyte proliferation (Valentine et al., 2021). Iron supplementation has also been shown to reduce iron deficiency and improve vaccination responses in humans (Stoffel et al., 2020; Drakesmith et al., 2021). Because iron is necessary for T and B cells to function properly and because its absence interferes with T cell proliferation and the synthesis of effector molecules like interferon- γ , iron deficiency can impair immunity. Both *in vitro* and clinical studies have demonstrated that increased hepcidin activity, which restricts iron absorption, can worsen iron deficiency and compromise immunological responses.

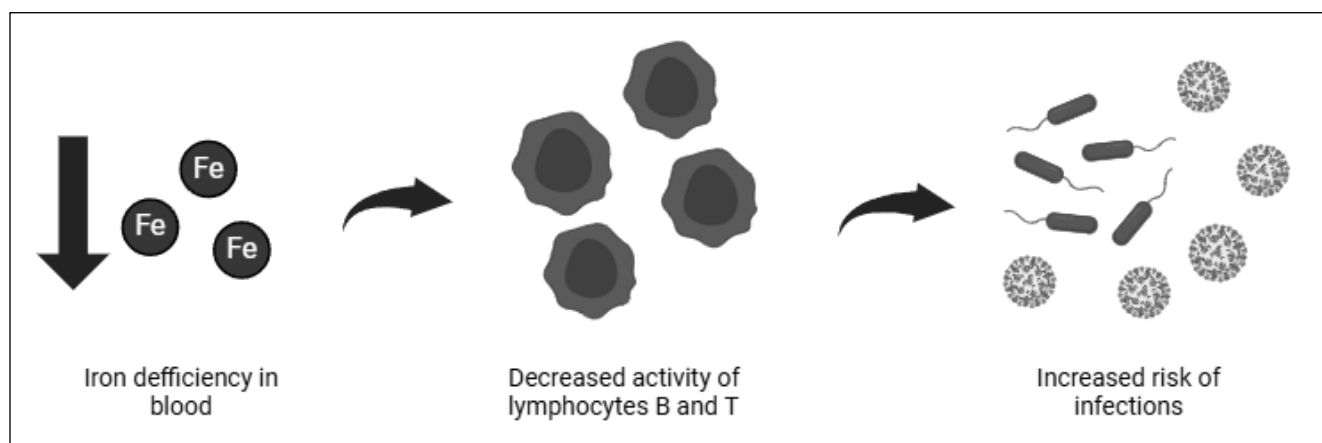


Fig. 1. The effect of iron deficiency on the immune response. A decrease in blood iron levels leads to reduced availability of this element to immune cells. In response to iron deficiency, T and B lymphocytes show increased proliferation, which symbolizes their activation. Additionally, hepcidin, by limiting iron absorption in the intestines, reduces the availability of this element, which weakens the immune response and increases the risk of infection.

Weakening of macrophage function

MS lesions are distinguished by the presence of various immune cells, such as lymphocytes, macrophages, and dendritic cells (Cheng et al., 2017; Cavallo, 2020). All of these cells are thought to play an important role in the development of MS lesions, and they do so through a variety of coordinated inflammatory pathways (Haase & Linker, 2021). Macrophages have a vital role in pathogen phagocytosis, the synthesis of cytokines that govern inflammatory responses, and the maintenance of the innate immune system (Hirayama et al., 2017). Macrophages are categorized into two types based on their activation state: those that are conventionally activated, with inflammatory functions, and those that are alternatively activated, playing a role in immune regulation (Strizova et al., 2023). In the context of MS, inflammatory alterations are related to macrophage activation, which can be distinguished depending on iron levels and other environmental factors (Mehta et al., 2013). For example, high levels of intracellular iron activate M1 macrophages, thereby reducing the number of anti-inflammatory M2 macrophages. Consequently, lower iron levels may limit M1 macrophage activation, leading to an enhanced inflammatory response and an imbalance between M1 and M2 macrophages (Feng et al., 2024). Chronic demyelinating lesions in MS show infiltration of myelin-laden macrophages, which causes deterioration of the myelin sheaths that surround neurons (Podbielska et al., 2013). These lesions are frequently linked with lymphocytic infiltrates, which work alongside macrophages in the demyelination process (Love, 2006). This mechanism is selective, as axons remain largely intact

while macrophages destroy myelin (Park et al., 2020). Monocyte infiltration across the BBB is an important stage in the formation of these lesions (Zhang et al., 2025). In active demyelinating plaques, macrophages swallow myelin debris, causing additional damage to the myelin sheaths (Höftberger & Lassmann, 2017). This phenomenon is caused by the interaction between macrophages and myelin, which also involves T lymphocytes (Legroux & Arbour, 2015). The presence of monocytes in the perivascular space in progressive MS lesions stimulates the establishment of lymphoid tissue, which can produce pathogenic autoantibodies (Lassmann, 2018). In the context of experimental MS investigations, macrophages interact with myelin in response to BBB disruption, triggering myelin phagocytosis (Kopper & Gensel, 2018). Furthermore, interleukin-10 (IL-10) secreting B cells can stimulate microglia and macrophages, encouraging phagocytosis and remyelination. In this process, macrophages and microglia exhibit various activation patterns, which can be both pro-inflammatory and helpful in terms of controlling the inflammatory response (Touil et al., 2023). Myelin breakdown begins when IgG-positive macrophages come into touch with mostly intact myelin sheaths (Prineas & Parratt, 2021). There is additional evidence that microglia and macrophages collaborate to remove myelin and promote remyelination. It should be highlighted, however, that the specific involvement of these cells in the pathophysiology of MS, as well as the mechanisms that regulate their activation, are still being studied intensively (Mado et al., 2023; Baaklini et al., 2023). Iron deficiency can limit their ability to secrete cytokines and remove pathogens, making them more susceptible to infections.

The role of iron in microglia

Neuroinflammation is an inflammatory reaction in the CNS that is induced by both external and endogenous causes (Cronin et al., 2019). An early inflammatory response following CNS injury may be advantageous because it allows for the elimination of toxic chemicals and inhibits the progression of injury, hence aiding tissue repair (Köllicker-Frers et al., 2021). However, chronic inflammation in neurological illnesses causes damage to nervous system structures, and its long-term presence may contribute to neurodegeneration (Sochocka et al., 2017). Furthermore, as one gets older, the risk of acquiring neurodegenerative illnesses increases, and neuroinflammation plays an important role (Zhang et al., 2023). The CNS's original immune cells are microglia, which are formed from primary macrophages in the embryonic yolk sac (Kempuraj et al., 2016). Microglia are the innate immune system's initial line of defense, responsible for phagocytosis and the elimination of abnormal protein aggregates, dead cells, and other foreign particles (Colonna & Butovsky, 2017). Microglia activate M1 type microglia in response to inflammatory stimuli like lipopolysaccharide (LPS) and IFN- γ , producing proinflammatory cytokines like interleukin-1 β (IL-1 β), interleukin-6 (IL-6), and tumor necrosis factor (TNF- α), resulting in CNS tissue damage. IL-4 and IL-13 can transform microglia into the M2 type, producing anti-inflammatory factors like transforming growth factor- β , interleukin-4 (IL-4), IL-10, and in-

terleukin-13 (IL-13). This supports nerve repair and regeneration, contributing to CNS homeostasis (Kwon & Koh, 2020; Shao et al., 2022; Guo et al., 2022; Qin et al., 2023). Iron buildup in microglia, caused by inflammatory cytokines, is a key part of the neuroinflammatory process (Li et al., 2024). Stimulating microglia with LPS, TNF- α , or IL-6 activates the production of proteins such as divalent metal transporter 1 (DMT1) and hepcidin, which transport and accumulate iron in these cells (Vela, 2018; Ward et al., 2022). Increased iron concentration in microglia causes TNF- α production, leading to inflammation (Rosenblum & Kosman, 2022). In multiple sclerosis, iron accumulation in microglia at the edges of demyelinating lesions leads to a shift toward a pro-inflammatory state, contributing to the worsening of pathogenic changes (McIntosh et al., 2019). Histological investigations have demonstrated that within these lesions, microglia collect iron, which produces more severe inflammation and intensifies the damage in CNS structures (Mehta et al., 2013). To summarize, iron buildup in microglia in chronic neuroinflammatory illnesses such as MS is an essential mechanism that contributes to the progression of CNS damage (Rock et al., 2004). Microglia, with their bidirectional function - both defensive and damaging - play an important part in the etiology of neuroinflammatory diseases. Future research into the mechanisms of iron buildup and the involvement of microglia in neuroinflammation may pave the way for new therapeutic approaches to slow the progression of neurodegenerative disorders.

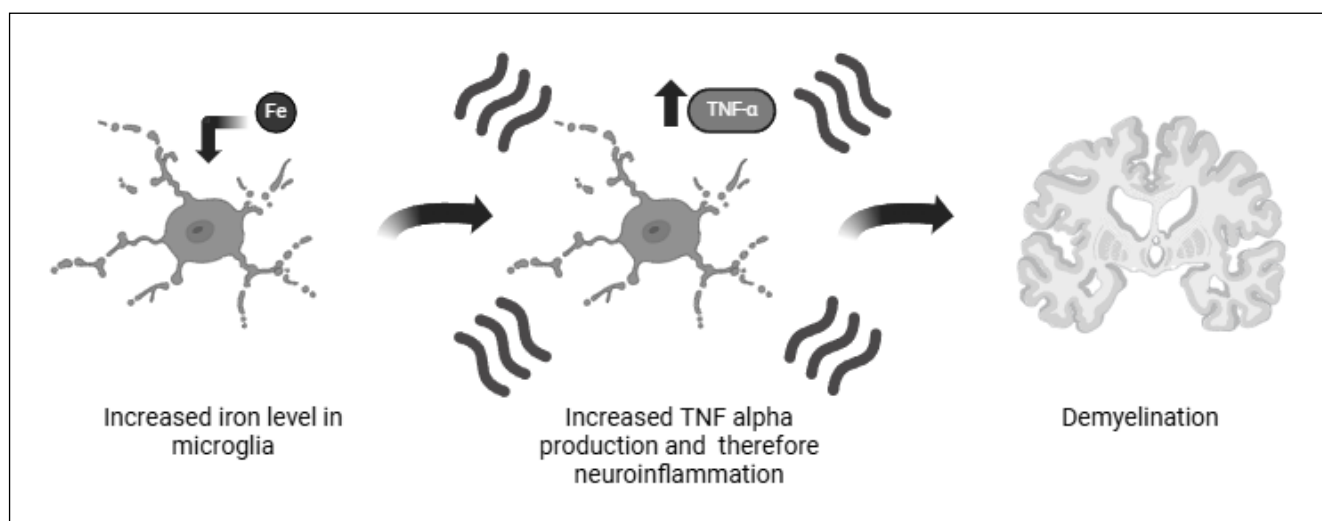


Fig. 2. The mechanism by which increased iron levels in microglia lead to the production of tumor necrosis factor alpha (TNF- α). Increased iron concentration in microglia stimulates an inflammatory response, which results in the activation of inflammation within the nervous system. This process leads to neuroinflammation and, consequently, to demyelination, which is particularly evident in diseases such as multiple sclerosis. Iron accumulation in microglia increases damage to brain and spinal cord structures, contributing to disease progression.

Mitochondrial dysfunction – reactive oxygen species origin

Oxidative stress results from excess iron in cells and tissues, which upsets redox equilibrium and promotes the growth of ROS (Williams et al., 2012). Age-related iron buildup in the human brain exacerbates the effects of oxidative damage and mitochondrial dysfunction, which also contribute to brain damage in aging and age-related ischemic vascular disorders (Galaris et al., 2019). Glutamate Ca^{2+} excitotoxicity, oxidative stress, and metabolic dysfunction are all brought on by ROS, which are produced when the CNS's divalent iron (Fe^{2+}) content rises, for instance, as a result of microhemorrhages or the deterioration of oligodendrocytes and myelin (Chakrabarti et al., 2011). Iron is present in microglia/macrophages and astrocytes in active and chronic lesions, but it is also stored in oligodendrocytes and myelin in normal-appearing WM and GM in MS patients (Spaas et al., 2021). While iron is comparatively elevated in the perivascular tissue, it appears to decline with age in the subcortical WM of MS patients compared to healthy controls (Popescu et al., 2017). On the other hand, too much iron can cause neurotoxicity, inflammation, and oxidative stress, which can harm and damage the function of neurons. Cognitive impairment and disability in MS are exacerbated by iron deposition in the basal ganglia, which can happen concurrently with or independently of demyelination (Schweser et al., 2021).

ROS are created when electrons are moved from one material (the reactant) to another in an oxidation reaction. ROS constitute the most significant kind of radicals generated in biological systems. The three primary forms are hydroxyl radical (OH^\cdot), hydrogen peroxide (H_2O_2), and superoxide anion (O_2^\cdot). While much larger concentrations of ROS can harm cellular components such as proteins, lipids, and nucleic acids, immune cells produce high local concentrations of ROS to destroy invaders (Williams et al., 2012). Oxidative species are produced in part by NADPH oxidase and the mitochondrial electron transport chain (Afzal et al., 2023). The trimeric Nrf2-Keap1-Cul3 complex regulates ROS levels in mammalian cells and the production of antioxidant enzymes. According to reports, ROS may play a role in the development and course of autoimmune illnesses such as MS (Dikalov, 2011).

The absence of the nuclear factor erythroid 2-related factor 2 (Nrf2) signaling pathway, a crucial antioxidant transcription factor, is another feature shared by neurodegenerative and/or neurological diseases, such as MS. Nrf2 inhibits oxidative stress, neuroinflammation, and mitochondrial failure (Tavassolifar et al., 2020). Axonal degeneration, neuro-

nal death, and a variety of neurological impairments that present with wildly disparate symptoms are all caused by demyelination. Therefore, in MS patients, primary demyelination results in neurodegeneration, which causes severe neurological damage and impairment (Villavicencio et al., 2021). By regulating the synthesis of adenosine triphosphate (ATP), redox balance, and intracellular free Ca^{2+} concentration in neuronal cells, mitochondria are essential organelles in charge of energy supply. In order to sustain and carry out membrane excitability, neurotransmission, and plasticity, as well as to regulate variations in (Ca^{2+}), which aid in the release of neurotransmitters, neurons rely heavily on mitochondrial function (Correale et al., 2019).

When compared to age-matched healthy brain samples, post-mortem examination of MS brain tissues revealed fractured neurofilaments, decreased organelle content, and defective axoplasm (Duarte et al., 2023). It has also been observed that neurons with a higher mitochondrial content have significantly greater levels of organelle content compared to demyelinated axons, which results in impaired mitochondrial transport. In both CSF samples and animal models, MS-based studies have commonly proposed oxidative damage caused by elevated ROS and reactive nitrogen species (RNS) (Maldonado et al., 2022). The Nrf2 pathway, a transcription factor encoded by the nuclear factor erythroid 2-related factor 2 (*NFE2L2*) gene and a member of the Cap'n'Collar family of transcription factors, regulates the balance of ROS. These genes, which encode glutathione S-transferase (GST), glutamate-cysteine ligase (GCL), heme oxidase-1 (HO-1), NAD(P)H quinone oxidoreductase-1 (NQO1), and glutathione peroxidase, are a defense response to ROS overproduction (Campbell & Mahad, 2011). The Nrf2 pathway is unable to sustain ROS at physiological levels during redox abnormalities, such as those seen in MS (Tebay et al., 2015). However, there is mounting evidence that Nrf2 may play a role in persistent neuroinflammation. In fact, Nrf2 regulates nuclear factor kappa-light-chain-enhancer of activated B cells (NF κ B) -dependent responses, another important route triggered by oxidative stress. Its activation constitutes a critical antioxidant checkpoint for astrocytes, conferring neuroprotective properties during neuroinflammation (Maldonado et al., 2022).

Lymphocyte CD3+ in MS pathogenesis

MS is caused by the loss of immunological tolerance to self-proteins, which results from the interplay of hereditary vulnerability and environmental

variables. This results in the production of autoreactive T and B cells, which are critical to the disease's progression (Maldonado et al., 2022). Th17 cells that produce glutamate, in particular, can cause damage to oligodendrocytes, resulting in an attack on myelin antigens. However, in MS, BBB function is compromised (Saha et al., 2020), which promotes T lymphocyte migration to the CNS. Special attention should be devoted to CD8⁺ T-cells, which have been found in both the CNS and the CSF (Lazibat et al., 2018). These cells accumulate in the CNS and are found in the blood of MS patients, confirming their function in the demyelination process (Larochelle et al., 2021; Nishihara et al., 2022). A decrease in the number of T-regulatory cells (Tregs) during the course of MS is associated with an increase in symptoms (Liu et al., 2022). Their number fluctuates with the stage of the disease, decreasing during relapses and increasing during remission (Pirko & Noseworthy, 2007). In turn, histological studies show the presence of leukocyte infiltrates in demyelinating plaques in the CNS, providing additional evidence for T-cell involvement in MS pathophysiology (Calahorra, et al., 2022). An increase in the number of CD20⁺ T-cells is particularly prominent in individuals with the primary progressive type of MS (PPMS), and it also correlates with the intensity of demyelination and disease severity in relapsing-remitting multiple sclerosis (RRMS) (Benallegue et al., 2022). Furthermore, extra iron in the body may alter the activation of T lymphocytes, particularly Th1 cells, which secrete proinflammatory cytokines like interferon-gamma. High iron levels stimulate uncontrolled activation of these cells, resulting in greater inflammation and additional nervous system damage (Findling et al., 2018). Iron overload can exacerbate MS by altering T lymphocyte activation, particularly Th1 cells, which contribute to inflammation and neurodegeneration. Elevated iron levels in the body may enhance the production of proinflammatory cytokines, further promoting myelin damage and disease progression.

Lymphocyte CD8⁺ in MS pathogenesis

Although T-cells are commonly acknowledged as the primary contributors to inflammatory demyelination in MS, emerging data suggests that B lymphocytes play an important role in the disease's pathogenesis (Maldonado et al., 2022). MS may be caused by B cells through various mechanisms, including antigen presentation to T-cells, which drives autoproliiferation of brain-homing T-cells, production of proinflammatory cytokines and chemokines such as

lymphotoxin- α , TNF- α , IL-6, and GM-CSF, which propagate inflammation throughout the CNS and generation of soluble toxic factors that damage oligodendrocytes and neurons. These pathways contribute to the persistent inflammation and neurodegeneration associated with MS (Pirko & Noseworthy, 2007; McLaughlin & Wucherpfennig, 2008; Lazibat et al., 2018; Findling et al., 2018; Milo, 2019; Saha et al., 2020; Larochelle et al., 2021; Ni et al., 2022; Nishihara et al., 2022; Liu et al., 2022; Calahorra, et al., 2022; Benallegue et al., 2022; von Essen et al., 2023). Excessive release of lymphotoxin- α , TNF- α , IL-6, and GM-CSF by B cells in patients' peripheral immune systems can worsen inflammatory processes (van Langelaar et al., 2020; Comi et al., 2021). There is also some evidence that B lymphocytes can produce autoantibodies against myelin, which contributes to disease development (de Gruijter et al., 2022). Clinical studies have proven the role of B cells in the development of MS, with anti-CD20 monoclonal antibody therapy demonstrating substantial efficacy in decreasing disease activity (Pala et al., 2018). MS patients' peripheral B cells produce anti-inflammatory cytokines such as IL-10, interleukin-35 (IL-35), and transforming growth factor β (TGF- β), perhaps maintaining immunological homeostasis (von Büdingen et al., 2011; de Sèze et al., 2023). Their presence in the brain and meninges, particularly in areas of demyelination, suggests a critical involvement in the local immune response, which may contribute to disease worsening (Gharibi et al., 2020). Iron overload may exacerbate inflammatory processes in MS by promoting oxidative stress, which damages oligodendrocytes and neurons. This increased oxidative stress can enhance the activation of B lymphocytes, worsening neuroinflammation and contributing to disease progression.

Toxic consequences of Fe

Neuronal metabolism associated with iron

Iron is a cofactor for the enzyme tyrosine hydroxylase (TH), which catalyzes the first and limiting step in dopamine production (Li et al., 2016). This enzyme requires iron (Fe²⁺) to function, hence iron levels are directly tied to dopamine release. Additional TH activity is typical, resulting in the cofactor BH₄, complicating the process of dopamine production (Lubetzki & Stankoff, 2014). The onset of iron deficiency can interfere with neuronal metabolism, resulting in dopaminergic dysfunction and associated behavioral issues, which may exacerbate MS symptoms in individuals with MS (Lozoff, 2011). In turn, dopamine in-

fluences the steering system, including in the context of MS, where it modulates the response, particularly in the regulation of Th17 cells (Xiao et al., 2021).

Dopamine has been found in studies to limit the release of proinflammatory cytokines by Th17 cells, which may have anti-inflammatory effects in MS. In this scenario, stimulation of the D2 dopamine receptor type inhibits the Th17 immune response (Juárez Olguín et al., 2016). Furthermore, dopamine influences the functioning of both the innate and adaptive immune systems, including effects on macrophages, dendritic cells, monocytes, and CD4+ and CD8+ T-cells that can generate dopamine and exhibit dopaminergic autoregulation (Melnikov et al., 2016). Dopamine affects neuropsychological symptoms in MS, including fatigue, cognitive impairment, and depression (Melnikov et al., 2022). Understanding the role of dopamine in the course of this disease, where dopamine can be identified based on the severity of individual neurological symptoms, and above all, because the occurrence of dopamine can be determined, as a result of which the release of proinflammatory cytokines will not be adequately inhibited has a direct impact on inflammation. This is crucial for the understanding of neuronal homeostasis and has consequences in the aftermath of MS prognosis, causing disease progression and harming brain structures and the spinal cord.

Inflammatory processes associated with IL-6, STAT3

Chronic inflammation in MS disrupts iron metabolism, which can lead to poor iron management. Macrophages and microglia, immune cells found in the CNS, release iron in response to inflammation, and incorrect management of this process can result in increased oxidative stress, which harms cells and tissues (Arreola et al., 2016). Proinflammatory cytokines, such as IL-6, stimulate the IL-6R-JAK2-STAT3 pathway, increasing the expression of hepcidin, an acute-phase protein that blocks iron release from macrophages and promotes iron retention in the body (Margoni et al., 2023). High levels of hepcidin cause inflammatory anemia defined by high serum ferritin and decreased erythropoiesis due to a lack of accessible iron (Williams et al., 2012). Furthermore, IL-6 is required for the development of T-cells into Th17 cells, which

produce proinflammatory cytokines such as interleukin-17 (IL-17), interleukin-21 (IL-21), GM-CSF, granulocyte colony-stimulating factor (G-CSF), and interleukin-22 (IL-22). Th17 cells can pass the blood-brain barrier, causing neuroinflammation and demyelination in MS (Tesmer et al., 2008; Zepp et al., 2011; Paganini et al., 2019; Camaschella et al., 2020). Increased IL-6 levels in the CSF are associated with reduced synaptic plasticity, which leads to clinical signs of brain injury, such as cognitive impairment and exhaustion (Gruol, 2015). Furthermore, elevated levels of IL-6 in the peripheral circulation can worsen inflammation, increasing disease activity in MS (Stampanoni Bassi et al., 2020). As a result, modulating inflammatory cytokine levels, particularly IL-6, may have substantial therapeutic implications for regulating chronic inflammation and improving iron metabolism, both of which may help alleviate MS symptoms (Maggio et al., 2006).

Damage to the blood-brain barrier

Iron is essential for the BBBs proper function, particularly in the case of MS, where the disease's pathophysiology heavily relies on iron damage. By regulating the entry of chemicals and cells into the CNS, the BBB serves as a physical barrier for the brain (Schreiner et al., 2022). Damage to this barrier encourages inflammatory cells and cytotoxic chemicals to infiltrate, which exacerbates inflammation and damages brain structures in MS (Miljković & Spasojević, 2013). The BBB's integrity is jeopardized by iron, a strong catalyst for free-radical processes, which can raise oxidative stress (Rand et al., 2021). Additionally, extra iron in the brain might contribute to additional cellular damage and increase clinical symptoms in MS (Williams et al., 2012). Conversely, iron is essential for axonal myelination, and oligodendrocytes may store and transfer iron throughout the BBB because they employ ferritin-iron to build myelin. Myelination effectiveness may be impacted, and the disease's brain damage may worsen, if this process is disrupted, for example, by rupturing the BBB (Cheli et al., 2020). Therefore, maintaining appropriate brain function and preventing iron overload depend on the brain's iron levels being properly regulated.

Table 2. Summary of different pathways of Fe toxic influence on neuronal tissue.

Neuronal Metabolism and Iron	Inflammatory Processes (IL-6, STAT3)	Blood-Brain Barrier (BBB) Damage
<ul style="list-style-type: none"> - Iron is a cofactor for tyrosine hydroxylase (TH), essential for dopamine production (von Büdingen et al., 2011; de Sèze et al., 2023) - Dopamine production and release are tied to iron levels, affecting various systems, including immune response (Gharibi et al., 2020) - Dopamine modulates Th17 cell responses and proinflammatory cytokine release, potentially having anti-inflammatory effects in MS (Lubetzki & Stankoff, 2014) - Dopamine impacts neuropsychological symptoms in MS, including cognitive impairment and depression (Xiao et al., 2021) - Deficiency or excess of dopamine can exacerbate neuroinflammation, impacting MS progression and neuronal homeostasis (Xiao et al., 2021) 	<ul style="list-style-type: none"> - Chronic inflammation disrupts iron metabolism in MS, leading to oxidative stress and cellular damage (Juárez Olguín et al., 2016) - IL-6 activates the IL-6R-JAK2-STAT3 pathway, which increases hepcidin expression, causing iron retention and limiting available iron for erythropoiesis (Melnikov et al., 2016) - Elevated IL-6 levels contribute to Th17 cell development, which causes neuroinflammation and demyelination in MS (Arreola et al., 2016; Margoni et al., 2023; Williams et al., 2012; Camaschella et al., 2020) - High IL-6 levels in CSF and peripheral circulation are associated with increased disease activity, cognitive impairment, and exhaustion in MS (Pagani et al., 2019) - Modulating IL-6 levels may help alleviate MS symptoms by reducing inflammation and improving iron metabolism (Zepp et al., 2011) 	<ul style="list-style-type: none"> - Iron is crucial for BBB function and plays a role in MS pathophysiology, as BBB damage allows immune cells and toxins to enter the CNS, worsening inflammation and brain damage (Gruol, 2015) - Iron-induced oxidative stress can compromise BBB integrity, further exacerbating MS symptoms (Maggio et al., 2006) - Iron is also vital for myelination and is stored by oligodendrocytes for this process. Disruption of BBB or iron regulation can worsen brain damage in MS (Miljković & Spasojević, 2013)

Animal models focused on iron metabolism and MS

Animal models (primarily rodents like mice or rats) are commonly used to study various aspects of iron metabolism and how it relates to MS (Wang et al., 2019). MS is a neuroinflammatory disease where the immune system attacks the protective covering of nerve fibers (myelin) (Tafti et al., 2024). The role of iron in MS pathology can be studied through these models by manipulating iron levels in the body and assessing the effects on MS symptoms (Stüber et al., 2016).

Experimental Autoimmune Encephalomyelitis (EAE) is one of the most common conditions studied in animal models for MS research. It mimics the demyelination process of MS. By manipulating iron levels (e.g., *via* iron supplementation or depletion), researchers can examine how changes in iron metabolism affect disease severity and progression in MS (Constantinescu et al., 2011). Iron is essential for myelin formation and function, but too much or too little can lead to detrimental effects (Wallace, 2016). Animal studies can explore the effects of iron overload or iron deficiency in the brain and spinal cord and how it affects inflammatory responses, oxidative stress, and neurodegeneration (Dash et al., 2025). There are models where animals have altered iron metabolism due to mutations in genes like ferroportin or hepcidin (which regulate iron absorption and distribution) (Nicolas et al., 2002). These models help in studying the relationship between iron dysregulation and MS-like symptoms or neurodegenerative conditions.

Cell-based models focused on iron metabolism and MS

Cell-based models are used to study the cellular processes involved in iron regulation, inflammation, and myelination (Santacreu-Vilaseca et al., 2025). Microglial and astrocyte cultures: Since microglia and astrocytes play a significant role in neuroinflammation, these cells are often studied in the context of MS (Mado et al., 2023). They are also crucial in the brain's handling of iron. Iron may influence the inflammatory response of microglia in MS models, and cell cultures can help understand this relationship (McCarthy et al., 2018). Oligodendrocyte precursor cells (OPCs): These cells are responsible for forming the myelin sheath (Benarroch, 2023). Iron is crucial for myelin production, and disruptions in iron homeostasis can affect OPC differentiation and myelination (Cheli et al., 2020). Cell models can be used to study the impact of iron on MS. Iron plays a role in redox reactions and can catalyze the production of reactive oxygen species (ROS) (Galaris et al., 2019). Elevated ROS levels contribute to cellular damage and neurodegeneration in MS (Ohl et al., 2016). *In vitro* models with neurons or glial cells can help examine how iron-induced oxidative stress influences MS progression (Czpakowska et al., 2024). Impairment of wound healing and fibrosis results in the buildup of cells and extracellular matrix constituents, with fibroblasts and their activated forms—myofibroblasts—present at injury sites (Klingberg et al., 2013). Disruptions in iron metabolism could influence the functioning of these cells, as iron plays a vital role in various inflam-

matory responses and healing processes (Ni et al., 2022). Fibroblasts in MS lesions contribute to extracellular matrix deposition and fibrosis. They produce collagen and fibronectin, contributing to the function of oligodendrocyte precursors (Lozinski et al., 2024). In MS, where there is damage to myelin and tissue within the CNS, alterations in iron metabolism may lead to disruptions in cell function, including that of fibroblasts, impacting repair processes and scar formation.

Pharmacological and non-pharmacological methods of modifying iron levels

The level of iron in the blood can be modified both through pharmacological and non-pharmacological methods. In the case of iron deficiency, it can be supplemented orally in the form of compounds such as iron sulfate, iron gluconate, or iron fumarate. This is the most commonly used way of modifying iron levels currently applied in medicine. The range between 50 mg and 200 mg of elemental iron per day for three to twelve weeks is usually necessary for effective iron substitution with oral supplements (Stoffel et al., 2017; Pantopoulos, 2024). In exceptional cases, intravenous iron supplementation may be used. This type of supplementation is typically indicated by poor tolerance to oral preparations, severe iron deficiency, impaired iron absorption from food, as well as women in advanced stages of pregnancy or heavy menstrual bleeding in women (Gupta et al., 2016; Pantopoulos, 2024). Furthermore, combined supplementation of iron with vitamin C can help improve iron absorption from the gastrointestinal tract due to its ability to convert ferric ions into ferrous ones. Vitamin C improves the absorption of iron even when the intake is modest compared to what is found in a typical diet (Li et al., 2020b; Doseděl et al., 2021).

Improved iron absorption can also be achieved through non-pharmacological methods. One such method may be modifying the diet. Products that contain notable amounts of iron include meat, fish, cereal products, and green plants such as spinach. By altering the diet to include the aforementioned items in meals, it is possible to support the increase of iron levels in the blood (Skolmowska et al., 2022). Avoiding products such as coffee and tea in the diet may help with iron supplementation. Studies have also shown that certain antinutritional factors found in these beverages, such as tannins and phytates, impair the bioavailability of essential nutrients (Shaw et al., 2022). Cryostimulation may affect iron levels through mechanisms such as improved circulation, reduced

inflammation, or effects on metabolism, although this area still requires further research (Dulian et al., 2015). Intense physical activity itself may also contribute to increased iron loss due to sweat and tissue microtrauma, which is why iron deficiency is particularly common among athletes, especially those involved in endurance sports. Although there is currently no direct evidence that kinesiotherapy affects blood iron levels, this remains a subject of future research (Solberg & Reikvam, 2023).

DISCUSSION

The pathogenesis of multiple sclerosis involves disruptions to critical components of the nervous system, such as iron homeostasis, which are linked to neurodegeneration and impaired motor function. MS is a complex neurological disorder in which iron dysregulation may play a role in the disease's pathogenesis, potentially influencing its progression and increasing the risk of developing neurological damage. It is getting more and more serious globally, particularly in women. When hemoglobin is released into the body's main system, it causes neurodegeneration and brain shrinkage. In contrast, haptoglobin protects the body from hemoglobin release. An important regulatory component, ferritin is a source of iron storage, and when it malfunctions, can lead to neurotoxicity and disease damage. The disease progresses as a result of iron buildup and lipid peroxidation, which cause neurons to develop and myelin to be destroyed. Myelin deterioration and the formation of MS lesions are accelerated by oxidative stress and immunological activation brought on by dysregulation of the supply in MS. A low iron diet, or an application that prevents its absorption, can cause the deficit. The impact and burden on the immune system's operation and the deterioration of T and B lymphocytes have a crucial impact on the immune response.

Iron is necessary for the balance of macrophage types and for the utilization of macrophages, which power the immune response. A compromised response can result from inadequacies, raising the chance of autoimmune illnesses like MS. The CNS's defensive microglia are triggered by inflammation and have the ability to collect iron, which accelerates lymphocyte destruction. The buildup of components in microglia can deepen the structure of systemic systems in neuroinflammatory illnesses like MS. ROS can generate too much energy in cells by sending mitochondria and secreting ROS, which exacerbate neurodegenerative processes. The pathogenesis of MS is closely linked to alterations in iron homeostasis, ROS

production, and subsequent neuronal degeneration. Disruptions in iron regulation within the CNS can exacerbate inflammation and contribute to the progression of MS. The imbalance leads to the activation of autoreactive T and B lymphocytes, which play a crucial role in driving disease progression.

Special attention should be given to the role of CD8⁺ T cells, which accumulate in the CNS and contribute to demyelination. Additionally, B cells play a key role in MS pathophysiology by releasing proinflammatory cytokines and presenting antigens to T lymphocytes, further driving the inflammatory response. They are important when a local immune reaction takes place, which can happen as the disease progresses. They are found in the brain, particularly in cases of demyelination. Dopamine and inflammation cause iron to function as a neural mediator. Increased oxidative stress in MS can result from metabolic instability, which triggers the development of neurons and the disease. Hepcidin may act and cease operating as a result of chronic inflammation from MS that interferes with metabolic activity. Consequently, elevated IL-6 levels, a pro-inflammatory cytokine, can accelerate neurodegenerative and inflammatory processes, including demyelination in MS. Maintaining the BBB's integrity and controlling the amount of iron in the brain are essential for halting the disease's progression. A lumbar puncture to check for the presence of OCB in the cerebrospinal fluid is one of the procedures used to diagnose MS. When there is a high level of κ -FLC in CSF, it can be utilized as a biomarker to track the course of MS. The accumulation of iron in these locations, as seen in the MRI imaging, results from activation and neurotoxicity, indicating the disease's progressive nature.

CONCLUSIONS

MS is a chronic neurological disease characterized by neurodegeneration and autoimmune demyelination, particularly in young adults. Central to the pathophysiology of MS is the dysregulation of iron homeostasis, which contributes to neuronal damage. Key factors include the accumulation of iron in affected brain areas, leading to ferroptosis, a form of cell death associated with oxidative stress. This process accelerates neurodegeneration, exacerbating the loss of myelin and neurons. In addition to iron dysregulation, inflammatory responses and metabolic disruptions play significant roles in disease progression. The alteration of iron metabolism and the associated oxidative damage highlight the importance of targeting iron-related pathways as potential therapeutic strategies in MS.

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