Abstract

Neuromyelitis optica spectrum disorders (NMOSD) are central nervous system inflammatory disorders. Recently, autoimmunity against aquaporin-4 (AQP4) water channel is identified to be the underlying immunopathogenetic mechanism of the majority of NMOSD patients. This is evidenced by the detection of IgG autoantibodies against aquaporin-4 (AQP4-IgG) in the serum of ~75%-80% of typical neuromyelitis optica patients. Detection of AQP4-IgG is highly specific for NMOSD, facilitates diagnosis of NMOSD and their distinction from classical multiple sclerosis. Besides its diagnostic value, AQP4-IgG is likely directly pathogenic in NMOSD. This review focuses on the immunopathological effects of AQP4-IgG in NMOSD.

Introduction

Neuromyelitis optica (NMO) and NMO spectrum disorders (NMOSD) are central nervous system inflammatory demyelinating disorders (CNS IDD) characterized by monophasic or relapsing optic neuritis (ON), acute myelitis (AM) and less commonly encephalitis [1]. NMO-IgG, an autoantibody detected in serum of 73% of NMO but not in classical multiple sclerosis (CMS) patients [2]. The autoantigen targeted by NMO-IgG is aquaporin-4 (AQP4), the major CNS water channel protein which is abundantly expressed in foot processes of astrocytes [3]. The discovery of these IgG autoantibodies against aquaporin-4 (AQP4-IgG) detected in the serum of ~75% of NMO patients has clarified that NMOSD seropositive for AQP4-IgG are not CMS, but an autoimmune disorder affecting CNS astrocytes (astrocytopathy). NMOSD patients seropositive for AQP4-IgG have underlying autoimmunity against CNS astrocytic AQP4. A small proportion of patients with typical NMO (relapsing ON and extensive AM without brain involvement suggestive of CMS) and some with restricted forms such as relapsing extensive myelitis without ON are seronegative for AQP4-IgG. The immunopathogenesis of these patients are uncertain. A proportion of these patients (~20-25%) are seropositive for IgG autoantibodies against myelin oligodendrocyte glycoprotein (MOG-IgG). The pathogenesis of MOG-IgG positive patients await clarification from further studies, especially histopathological studies of affected CNS tissues.

In NMOSD, patients typically have monophasic or relapsing severe AM and ON, with relatively less frequent brain involvement especially in the early phase, and CSF OCB are infrequent [1]. NMO patients typically have severe neurological disability after attacks of extensive myelitis, severe ON and brainstem encephalitis [4], no secondary progression and worse clinical outcome than CMS patients [1,5]. Pathologically, spinal cord tissues of NMO patients exhibit necrosis in both gray and white matter, infiltrating leucocytes (macrophages, polymorphonuclear cells [neutrophils, eosinophils], and lymphocytes), activated microglia, demyelination, axonal loss, thickened hyalinized vessel walls with deposits of IgM, IgG and complement activation products in a vasculocentric rim and rosette pattern. These hyalinized vessel walls with deposits of immunoglobulins and complement activation products in the characteristic vasculocentric rim and rosette pattern is not observed in lesions of CMS patients.

Clinical usefulness of AQP4-IgG

Distinction of NMOSD from CMS can be difficult especially in the early stage [5]. In addition, brain involvement in NMO is increasingly recognized, and NMO patients with MRI brain lesions fulfilling criteria for CMS are reported [6]. Early diagnosis of NMOSD is important as disability in NMOSD is relapse-related [1], hence relapse prevention is the key in treatment. Importantly, commonly used disease modifying drugs for relapsing MS including β-interferon, fingolimod and natalizumab may be ineffective or even harmful in NMOSD by precipitating development of extensive hemispheric lesions [7]. Immunosuppression by azathioprine/mycophenolate mofetil and corticosteroids or B-cell depletion therapy (e.g. rituximab) should be initiated early in NMOSD to prevent relapses and disabilities [8]. Detection of AQP4-IgG greatly facilitates early diagnosis and prompt treatment of NMOSD.

Immunopathologic effects of AQP4-IgG

AQP4-IgG is predominantly of IgG1 subclass, and binds to AQP4 abundantly expressed as transmembrane protein in astrocytic end feet processes at glial limitans abutting capillaries in pia, subpia and subependyma, and less abundantly in ependymal cells [3]. AQP4 has six transmembrane regions and three extracellular
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 loops with intracytoplasmic amino and carbonyl terminals. AQP4 exists as heterotetramers formed by the M1 and M23 isoforms (the 2 major isoforms). CNS AQP4 play important roles in functional integrity of blood-brain barrier (BBB) and CNS water homeostasis, regulating water transport across the interfaces between blood and brain, and between brain and CSF [9]. Using sera from 32 NMOSD patients with high serum concentrations of AQP4-IgG, Iorio et al. [10] showed that AQP4-IgG from NMOSD patients bind to extracellular loops, monomers, tetramers and high order orthogonal arrays of particles (OAP) of AQP4, and disease-specific epitopes reside in extracellular loop C more than in loop A or E. Importantly, AQP4-IgG bind most avidly to membrane AQP4 epitopes formed by loop interactions between tetramers and intermolecular interactions within OAP [10]. AQP4-IgG are likely directly pathogenic in NMOSD. IgG from serum of AQP4-IgG positive NMO patients bound to extracellular region of membrane AQP4 of transfected HEK293 cells and triggered

I. Internalization and endolysosomal degradation of bound AQP4, and

II. Complement activation with deposition of terminal lytic membrane attack complex, C9neo, causing cell lysis [11].

AQP4-IgG induced necrosis of cultured astrocytes in a complement-dependent manner. AQP4-IgG binding to extracellular domains of AQP4 have isoform-specific outcomes, M1 isoform is completely internalized while M23 isoform aggregates into larger OAP via cross-linking and resist internalization. The FC portion of the AQP4-IgG bound to these OAP triggers complement activation followed by astrocyte cytotoxicity, proliferation of foot processes. These suggest:

- AQP4-IgG in peripheral blood become accessible to CNS AQP4 AQP4 are the initial step in neuroinflammation of NMOSD. IgG from NMOSD patients showed evidences of astrocytic activation with marked proliferation of foot processes. These suggest:

I. CFA and PTx alone lead to BBB breakdown,

II. AQP4-IgG from NMOSD patient’s induced asymptomatic AQP4 loss and astrocytic activation without inflammatory cell infiltration, demyelination or astrocytic cytotoxicity in the absence of complement activation, and

III. AQP4-IgG can activate astrocytes in the absence of complement activation [14].

Colleagues and I studied the pathogenic role of AQP4-IgG in the absence of complement activation by passive transfer of IgG isolated from sera of NMO patients into mice (human IgG cannot activate mouse complements). Breakdown of BBB was induced before transfer of human IgG by subcutaneous CFA and intraperitoneal pertussis toxin (PTx). We observed that mice treated with IgG from AQP4-IgG positive NMOSD patients had areas of AQP4 loss in spinal cord while mice treated with IgG from AQP4-IgG negative NMOSD patients and IgG from healthy subjects did not. All mice had no clinical features of encephalomyelitis and no inflammatory cell infiltration, demyelination or loss of glial fibrillary acidic protein (GFAP) immunoreactivity. In addition, mice treated with IgG from AQP4-IgG positive NMOSD patients showed evidences of astrocytic activation with marked proliferation of foot processes. These suggest:

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A recent animal study showed that chronic infusion of IgG from AQP4-IgG positive NMOSD patients to CSF of rats results in motor impairment of the animal associated with astrocyte alteration characterized by loss of AQP4, myelin basic protein, axons and reduced glutamate uptake but no immune cell infiltration, microglial activation or complement activation.

AQP4 expression is markedly reduced in cord lesions of NMOSD patients but preserved or increased in cord lesions of CMS patients [15]. Recent histopathological studies further confirm that NMOSD have distinct pathologies from CMS. Although different pathologies are observed in different sites of lesions, AQP4 loss and astrocytic injury or cytotoxicity (losses) are the key pathologies in cord and brain lesions of NMOSD whereas normal AQP4 level and gliosis are observed in brain and cord lesions of CMS patients [15,16]. Loss of AQP4 is always observed in CNS lesions of NMOSD patients, whereas loss of astrocyte, complement activation and necrosis are variable. Histopathological studies of brainstem lesions at the dorsal medulla (area postrema without intact BBB) from a small number of NMOSD patients revealed loss of AQP4, prominent astrocytic activation, lymphocytic infiltration, microglial reactivation, some complement activation products but no obvious axonal or neuronal injury. It is proposed that AQP4-IgG binding to astrocytic AQP4 at this site is followed by rapid internalization of antigen-antibody complex, possibly explained by predominantly AQP4-M1 expressed in this site. The astrocytes react with inflammatory response leading to inflammatory cell infiltration without obvious demyelination, axonal or neuronal injury [17].

Conclusion

Current evidence suggests AQP4-IgG binding to astrocytic AQP4 are the initial step in neuroinflammation of NMOSD. AQP4-IgG in peripheral blood become accessible to CNS AQP4
at sites without intact BBB (such as the area postrema) or during inflammatory condition such as infection during which proinflammatory cytokines in circulation lead to breakdown of BBB. The astrocytic activation triggered upon binding of AQP4-IgG to AQP4 M1 and internalization of antigen-antibody complex release proinflammatory cytokines and chemokines leading to inflammatory cell infiltration. Other immune cells are activated with secretion of complements which will be activated especially at sites with high level of AQP4 M23 and hence OAP.

References