Chronic Cerebrospinal Venous Insufficiency in Multiple Sclerosis: The Hydrostatic-Immune Paradigm and the Flow Cytometry as a Diagnostic Tool

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Introduction

Multiple sclerosis (MS) is an inflammatory demyelinating disease of the central nervous system (CNS). Subjects 20-40 years old are affected in a relapsing-remitting, secondary progressive or primary progressive pattern. Although the clinical course of the disease is variable, it usually leads to progressive deterioration and severe disability [1]. Both genetic and environmental factors have been implicated in the pathogenesis [2]. The precise cause remains largely unknown, but there is evidence of a myelin-antigen-specific, Th1-mediated immune process resulting in macrophage-mediated demyelination, axonal loss and gliosis [3].

The vascular controversy

In the last years, a “vascular hypothesis” initially proposed by Charcot [4] has reemerged, causing heated debates among physicians and huge expectations among patients.

Zamboni et al. [5] combined transcranial and extracranial color Doppler ultrasonography to examine internal jugular, vertebral, deep cerebral, and azygous veins and evaluate 5 parameters that assess venous blood flow and vessel anatomy. A significant correlation between MS and the presence of 2 or more criteria of venous outflow abnormalities was reported. This led the authors to a highly criticized hypothesis that chronically impaired venous drainage of the CNS (termed “chronic cerebrospinal venous insufficiency”, CCSVI) may trigger the initiation of an inflammatory process which eventually leads to MS [6,7].

Attempts to treat CCSVI by venous angioplasty and/or stenting, a technique termed “liberation procedure”, have generated conflicting results [8], adding to the general confusion. Published studies so far display remarkable heterogeneity regarding study protocol and patient recruitment, making comparisons and evidence-based recommendations impossible. This is probably due to poor consensus on terminology and sonographic criteria as well as lack of a standardized methodology regarding patient selection and procedural technique [9].

Newer insights into MS pathogenesis

In a study of autopsy-derived MS tissue, Barnett et al. [10] reported that, within newly forming lesions, oligodendrocyte apoptosis and intramyelinic edema was accompanied by active phagocytic demyelination. The latter occurred in the absence of contact-dependent, cell-mediated immunity, suggesting that macrophage activity may be of a scavenger-like nature, in response to phagocytic ligands on apoptotic oligodendrocytes [11]. Whether oligodendrocyte death and demyelination unmask an autoantigen which causes autoimmune, or another factor induces both apoptosis and a systemic inflammatory response, is not known [11]. Either way, according to the paradigm proposed by Barnett et al, a population of microglia differentiates into myeloid-like dendritic cells. These migrate to the cervical lymph nodes (CLNs) and prime the differentiation of naïve T cells towards a central memory and effector phenotype. Such CCR7+CD68+MHCII+ dendritic cells have been observed in CLN tissue [12,13], although their precise ancestry is controversial [14]. Th1 lymphocytes then enter the CNS and are reactivated by antigen-presenting cells (APCs), amplifying the inflammatory response and creating a vicious, ultimately self-sustaining cycle. This temporal diversification of the immune process would explain the heterogeneity of MS lesions observed by Lucchinetti.
et al. [15] and reconcile observed histopathologic differences between active and chronic lesions.

The hydrostatic approach

On a more macroscopic scale, MS has been associated with diffuse brain hypoperfusion [16] and decreased cerebral metabolism [17] even in the initial stages of the disease [18]. Using dynamic susceptibility contrast-enhanced magnetic resonance imaging, Law et al. [19] found significantly decreased cerebral blood flow (CBF) and significantly prolonged mean transit time throughout the normal appearing white matter (NAWM) in patients with relapsing–remitting MS. Adhya et al. [20] also found significantly decreased CBF and cerebral blood volume (CBV) in NAWM regions of patients with either relapsing–remitting or primary progressive MS compared with healthy subjects.

These findings indicate that globally decreased white matter perfusion may be common in all subtypes of the disease [16]. Blood supply to the grey matter appears to be also compromised [21] either by the same mechanism affecting white matter or as a result of a dissociation between the cerebral cortex and subcortical structures due to the white matter damage. It is not clear whether hypoperfusion in MS is of a primary, vascular aetiology or secondary to reduced metabolic demand from axonal degeneration. Diffusion tensor imaging studies by Saindane et al. [22] point against a secondary insult, while, on a microscopic level, the absence of structural perivascular abnormalities indicates that hypoperfusion of the white matter in MS has a functional vascular origin [23].

Astrocytes occupy a significant proportion of the surface of intracerebral blood vessels and are involved in regulation of vessel diameter and CBF [24,25]. In MS white matter lesions and NAWM, astrocytes were found to be deficient in β2-adrenergic receptors [26]. Decreased CAMP formation due to impaired β2 signaling may lead to vasoconstriction trough several mechanisms [16], implying a key role for astrocyte dysfunction in MS hypoperfusion. The cause of astrocyte dysfunction has not been elucidated yet, but spatial correlation of lesions and blood vessels may provide a clue.

Lightman et al., studying the frequency of retinal vascular abnormalities in patients presenting with acute optic neuritis who went on to develop MS later, noted that “… the presence of perivenular abnormalities in a region free of myelin and oligodendrocytes provides evidence that the vascular changes in MS can occur independently of contiguous demyelination, and may be the primary event in the formation of new lesions” [27].

In initial stages of MS, plaques appear as thin, linear periventricular white matter lesions (Dawson’s fingers), oriented around the long axis of central veins [28]. Ultra-high-field 7T MRI studies showed that the majority of these lesions are associated with centrally coursing veins [29,30]. The close proximity of evolving plaques to veins has been demonstrated histologically in human brain tissue [31,32] showing perivascular cuffs, intramural fibrinoid accumulation, haemosiderin deposition and collagenised thickened vessel walls that were not seen in control subjects [33]. These histological findings are similar to those described in the case of chronic periphlebitis retinae, suggesting a common pathogenetic mechanism and implying that periphlebitis may be an initial event in plaque formation [34].

Another important finding in MS patients early in the course of the disease is evidence of diminished venous vasculature [35]. According to Zivadinov et al. [36], decreased venous vasculature in brain parenchyma of MS patients is strongly related to concomitant presence and severity of extracranial venous stenoses.

So, altered cerebral vasculature and the inflammatory process underlying MS may share common pathophysiological mechanisms [37]. This relationship is still poorly understood but may involve production and release of soluble substances with vasactive properties (NO, substance P, Interleukin-1) [38,39] and cytokines (TNF-a) [40]. Changes in neuronal homeostasis due to blood brain barrier (BBB) disruption have also been implicated [41]. In fact, defective endothelial tight junctions (TJ) have been found to coincide with microglial activation in active lesions and NAWM [42]. According to the authors: "The restriction of “putatively open” junctions to venules in active lesions and microscopically abnormal NAWM and the absence of similar alterations in chronic plaques may point to a specific mechanism of reversible junctional disruption restricted to such vessels, and related to the latter’s involvement in the diapedesis of inflammatory cells". In other words, "… altered endothelial cell function and might be the common final pathway leading to changes in perfusion and BBB leakage” [37]. The cause of endothelial malfunction is not clear but one theory suggests that venous stenoses may contribute [43].

Abnormal venous outflow, as described in patients with CCSVI, is not only due to downstream anatomical restrictions in blood flow (abnormal jugular valve, stenosis etc). The effect of an anatomical restriction may be augmented by a resonant standing wave created through reflection and convergence of transmitted arterial pressure waves along the vein [43]. The result is a localized increase in intravascular pressure at the venous end of the capillary bed (local hypertension), which, coupled with pulsatile shear stress and cyclic strain of the vessel wall can result in rupture of endothelial tight junctions and BBB breakdown [44]. Depending on magnitude of the latter, leakage of intravascular fluid, proteins and blood cells could occur. This would lead to significant changes in neuronal microenvironment, increased ISF and lymph production, local activation of the inflammatory cascade, iron deposition, oligodendrocyte loss, attraction of phagocytes, demyelination and eventually plaque formation [44].

Increased pressure at the venous end of a capillary bed has also important consequences on organ perfusion pressure and lymph production. According to Darcy’s law, the rate of blood flow in a vessel of specific length is proportional to the pressure difference between its ends. Increased venous capillary pressure in the face of normal arterial capillary pressure would lead to hypoperfusion [43] and “virtual hypoxia” [45], both of which have been described in the setting of MS.

Increased filtration pressure due to venous hypertension and intravascular fluid and protein leakage due to BBB breakdown can lead to increases in interstitial fluid. Any factor that increases interstitial fluid also increases lymph flow [46]. Lymphatic drainage of the CNS regulates the balance of ISF and solutes within the CNS microenvironment and represents an accessory route through which excess fluid and proteins can flow from the interstitial spaces back into the blood. It also plays an important role in neuroimmunological reactions, through physiological drainage of antigens from the brain to regional lymph nodes, mostly cervical and lumbar [47]. In other words, altered lymph flow due to increased production or decreased outflow may affect both neuronal milieu and immune response.

The immune approach: Unlike most other organs, the CNS lacks an anatomically well-defined lymphatic system. Cerebrospinal fluid (CSF)
and ISF drainage pathways are different in humans but both drain partly or wholly to regional lymph nodes.

CSF production and outflow through the arachnoid villi and granulations to the venous sinuses has been extensively studied in the past. Even though it is the primary drainage route in humans [48,49], a small fraction of CSF drains through the cribriform plate into the nasal submucosa and ultimately to cervical lymph nodes [50].

ISF and solutes can initially diffuse through brain parenchyma, drain out of the brain along basement membranes of capillaries and cerebral arteries [51] and finally leave the artery walls and drain to adjacent cervical lymph nodes at the base of the skull [52].

The motive force for drainage along artery walls in a direction opposite to blood flow may be related to the pulsations of arteries and the creation of a contrary reflection wave [53]. A decrease in the amplitude of the contrary wave due to changes in arterial wall elasticity or other reasons could result in impediment of perivascular lymphatic drainage [53].

This unique anatomic arrangement, coupled with the presence of the blood brain barrier and the low expression of Major Histocompatibility Complex class II (MHC II) molecules creates an optimally sheltered environment and contributes to what is termed "immunological privilege" of the CNS [54].

Immune surveillance in the CNS differs somewhat from that of other organs, contributing to the "immune privilege". It consists of an afferent and an efferent arm.

The afferent arm involves antigen transfer and presentation to naïve lymphocytes in regional lymph nodes or the spleen, resulting in their priming and activation. Antigen can be transferred by a fluid or a cellular route [55]. Soluble antigens (in ISF or CSF) can follow the fluid route through perivascular spaces [50-52] or along cranial (olfactory, optic, trigeminal, acoustic) and spinal nerves to regional lymph nodes [56]. CSF antigens can also gain direct access to the systemic circulation and to the spleen by absorption in arachnoid villi and granulations [47].

The cellular route involves antigen transfer by specialized antigen presenting cells (APCs) such as macrophages or dendritic cells (DCs). Macrophages containing myelin antigens have been described within CLNs of patients with MS [12]. This suggests that either APCs containing antigen migrate from the CNS to regional lymph nodes or that antigens draining from the CNS are taken up by APCs within the lymph nodes themselves [57]. To this date, there is no convincing evidence that inflammatory cells containing antigens can exit the healthy CNS parenchyma [55,58], even though this cannot be ruled out in the setting of inflammation.

Peripheral stimulation in lymphoid organs is another unresolved issue. Dendritic cells and macrophages containing myelin antigens have been found in CLNs, close to T-cells, suggesting interaction [12]. T cell responses to CNS antigens may depend on where this interaction takes place. Within CNS-draining CLNs, constitutive drainage of antigen from the healthy CNS can result in suppression of CNS-specific naïve T cell priming and induction of tolerance [59]. On the other hand, T cell responses initiated within peripheral, non-CNS draining lymph nodes, in response to systemic CNS antigen exposure, leads to T cell activation which eclipses the basal tolerogenic effect of soluble antigen drainage to CLNs [57].

The efferent arm involves inflammatory cell (monocytes, B- and T-cells) migration to the CNS and initiation of the inflammatory response. The process of cell activation and CNS homing is poorly understood. Cervical lymph node drainage may enhance CNS specificity of T cells by providing these cells with a CNS "homing beacon" in the form of signature molecules, such as CCR7 and CD68 [60]. Crossing of the BBB occurs by endothelial transmigration, similarly to what happens in other tissues: a sequential process of cellular rolling, adhesion and diapedesis, mediated and guided by adhesion molecules and chemokines [61]. Astrocytes are involved in this process, modulating BBB permeability and affecting lymphocyte recruitment [62]. At least three different routes of cell entry into the CNS have been described [63] but the fate of lymphocytes and macrophages once within the brain parenchyma is still controversial. They may remain in perivascular spaces, migrate through the basement membrane of the perivascular glia limitans into the CNS parenchyma and interact with target antigens [64] or simply undergo apoptosis [65].

Once inside the brain parenchyma, it is not clear how antigen presentation occurs in the absence of professional APCs and in the face of low expression of MHC II molecules [66]. T lymphocytes that have been pre-activated peripherically may release pro-inflammatory cytokines (IFN-γ, TNF-α) and induce MHCII molecules in CNS cells [67]. Alternatively, sentinel monocytes/macrophages stationed in perivascular spaces may act as APCs [66], thus initiating autoimmunity and closing the circle of immune surveillance.

Auto-reactive T and B cells are fundamental for development of MS and require antigenic presentation of myelin antigens by APCs in order differentiate into effector cells [68]. In the initial stages of the disease, before lymphoid neogenesis occurs and tertiary lymphoid structures within the meningeal compartment are able to sustain a continuous inflammatory process [69], these interactions probably take place in CLNs [3] and are dependent on CNS antigen transport to those sites [57]. So, CLNs may act as loci of antigen presentation and T- or B-lymphocyte priming [54] but also as sites of tolerance induction towards CNS antigens [70,71]. In this case, alterations in lymph flow towards CLNs could theoretically lead to alterations in antigen presentation and processing with unknown consequences for CNS immune surveillance.

The combined hydrostatic-immune paradigm of CCSVI: Current data does not support the view that CCSVI is the sole cause of MS. However a wider role in MS pathogenesis cannot, at the present time, be confirmed or refuted. If CCSVI does correlate with MS, then alterations in CNS venous outflow must somehow be associated with induction or attenuation of inflammatory and immune responses. The only direct link between the vascular and immune systems is the lymphatic system.

We have described previously how the presence of venous stenoses (CCSVI) may lead to changes in venous outflow, which may in turn affect central nervous system (CNS) interstitial fluid (ISF) production and composition. The latter would also affect lymph production and outflow, altering CNS antigen presentation to regional lymph nodes. Altered antigen presentation could lead to altered immune responses in the form of increased autoimmunity or decreased tolerance to self-antigens (Figure 1).
More specifically, according to this theory, the presence of anatomic or functional venous stenoses leads to absolute or relative (functional) capillary venous hypertension due to reflected pressure waves. This results in dysfunction of tight junctions, perivenous edema and compromised BBB function, affecting parenchymal homeostasis and causing astrocyte and oligodendrocyte dysfunction. Local phagocytic mechanisms activated by the death of these cells initiate demyelination. At the same time, increased ISF and lymph production coupled with an increased antigen load from apoptotic cells result in augmented antigen presentation to T-lymphocytes in CLNs, overwhelming their inherent tolerizing capacity and kick starting autoimmunity. This fuels the inflammatory process and perpetuates myelin damage.

Treatment of venous abnormalities (liberation procedure) would attenuate this process by alleviating venous hypertension and BBB damage. This would remove at least one of the hypothetical factors contributing towards astrocyte and/or oligodendrocyte apoptosis. The decreased perfusion pressure would also result in decreased ISF and lymph production and ultimately in decreased antigen transport and presentation to regional lymph nodes. In the initial stages of the disease, before the autoimmune process becomes compartmentalized in the Virchow-Robin spaces, altered antigen flow could lead to altered immune responses.

**Implications for research**

Unfortunately, study of the human lymphatic system is hampered by the difficulty to readily image it. Older techniques such as lymphangiography required considerable technical skill and were associated with complications. Newer, water soluble contrast agents minimize these problems but because of the difficulties in delivering sufficient amounts of agents necessary to produce X-ray contrast, classic lymphangiography is not routinely used today [72]. Lymphoscintigraphy is another imaging technique that enables two-dimensional planar visualization of the lymphatic network after injection of Tc-99m-labelled colloid, unfortunately with a low spatial resolution [73]. Magnetic Resonance Imaging (MRI) of the lymphatic system (MR-lymphography) is easier to perform and does not require exposure to ionizing radiation. It involves interstitial or intravenous injection of gadolinium-labeled contrast agents or iron oxide particles and provides a very high spatial resolution [74]. The most promising imaging modality today is through optical imaging of the lymphatics. It is based upon the administration of contrast agents (fluorescein, indocyanine green) that, when excited by light of specific wavelength, exhibit fluorescence. This technique has been used successfully in mapping human lymphatics [75].

To this day, there have been no studies in MS patients regarding CNS lymph outflow or its changes after angioplasty treatment for CCSVI (liberation treatment). Nevertheless, reversal of abnormal lymphoscintigraphy after placement of venous stents for correction of associated venous obstruction in the lower limbs of MS patients has been reported [76], supporting the hydrostatic aspect of our theory.

It would be interesting to further explore whether lymph production and antigenic composition is different in MS patients with or without CCSVI and whether liberation treatment causes a change in these parameters. However, studies like that would have to involve injection of contrast agent directly into the brain parenchyma, something which is currently ethically unacceptable in humans. Extracranial lymph outflow or changes in CSF flow are much easier to image and could be used instead, but no such studies have been conducted and it is not known whether they accurately reflect intracranial lymph flow. Future improvements in lymphatic system imaging technique may provide this opportunity.

**Flow Cytometry as a diagnostic tool**

We could investigate the possible prognostic value of ploidy in humans and the disruptions occurring inside the cell cycle with flow cytometry as a diagnostic tool. Flow cytometry enables rapid quantification of DNA content of individual cells, and the cellular DNA content provides useful information about the ploidy, expressing the modal DNA value, and the proliferative activity in a tissue. The ability of flow cytometry to estimate cellular DNA content is based on the measurement of fluorescence from dyes which bind in a stoichiometric manner to DNA [77].

As the DNA content is duplicated prior to cell division, mathematical models have been derived which can estimate the percentage of cells in different phases of the cell cycle. The use of flow cytometry for DNA analysis between family members with genetically linked diseases provides fast results, permits multiparameter analysis correlating DNA content with antigen expression, and also provides sensitivity for detecting near-diploid aneuploid peaks [78].

In cases like this, we consider that a genetic profile analysis should be performed to patients and their first degree relatives for more efficient therapies and follow ups of the disease. Flow Cytometry as an analysis method is rapid, sensitive, accurate and efficient for the genetic analysis and the progress of the disease.

**Conclusions**

MS is considered a primarily autoimmune disease but this paradigm is slowly shifting towards a combined autoimmune-
neurodegenerative model [79]. The pathogenetic mechanism is almost certainly multi-factorial and even though the initial trigger is still unknown, some of the waypoints on the road leading to demyelination and axonal loss have been discovered. It is not clear yet to what extend venous outflow abnormalities contribute to this process. Altered neuronal microenvironment and changes in antigen presentation to regional lymph nodes may provide a theoretical framework combining hydrostatic and immune effects. Lymph nodes are a site of sensitive immune regulation where both immune suppression and immune activation take place [80,81] CCSVI and its treatment may act via tipping the balance towards one or the other side of the scales.

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