

Case Report



CADASIL or MS? Consider "Red Flags" but Avoid a Misdiagnosis: Case **Series of a Concomitant Diagnosis**

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Abstract

Cerebral autosomal dominant arteriopathy with subcortical infarcts and leukoencephalopathy (CADASIL) is a monogenic autosome-dominant disease with chronic clinical course. Rarely, CADASIL may present with atypical relapsing-remitting manifestations, cerebral and spinal white matter lesions, mimicking inflammatory CNS disease as multiple sclerosis (MS). The rarely co-occurrence of MS and CADASIL may represent a hard challenging diagnosis even for an expert neurologist. Here, we present a case series of two patients with CADASIL showing MRI pattern overlapping MS. They were the only case of co-occurrence of CADASIL and MS in their own family. Both patients were treated with anti-inflammatory and anti-platelet drugs, mostly with good response. Pathogenic hypothesis highlights that genetic events, related to monogenic disease, may expose CNS antigens with a consequent self-immune attack. In CADASIL, the function of Notch3 receptor showed a consistent interplay with immune system activity. Indeed, certain mutations of Notch3 receptor show abnormal upregulation of specific pro-inflammatory patterns. However, even if it is not possible to determinate if the proinflammatory activity may be promoted by pathogenic mutations in Notch3, the "apparent" difference between MS and "inflammatory CADASIL" could be considered more semantic than etiologic.

Keywords: CADASIL; CNS Inflammation; Inflammatory CADASIL; Multiple Sclerosis; Monogenetic Disease; Neurodegeneration

Background

Cerebral autosomal dominant arteriopathy with subcortical infarcts and leukoencephalopathy (CADASIL) is an autosomal-dominant, small-vessel disease, clinically characterized by transient ischemic attacks and/or strokes, migraine and cognitive impairment possibly leading to pseudobulbar paralysis and progressive dementia. CADASIL is caused by pathogenic mutations in NOTCH3 gene, located on chromosome 19p13, which encodes a large single-pass transmembrane receptor expressed in vascular smooth muscle cells. More than 200 different pathogenic mutations, localized throughout exons 2-24 of the NOTCH3 gene have been already described in CADASIL patients [1]. Most of them lead to a variation in the number of cysteine aminoacidic residues in one of the 34 epidermal growth factor-like repeats (EGFrs) domains of the Notch3 protein. In patients the Notch3 receptor extracellular domain has been demonstrated to accumulate in the vessel wall with a direct or indirect toxic effect on vascular smooth muscle cells, leading to degeneration [2]. A clinical diagnosis of CADASIL is supposed when the patient shows the typical clinical manifestations associated with a

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specific magnetic resonance imagining (MRI) pattern of the brain: 1) diffuse white matter lesions involving in particular the temporal pole, 2) microbleeds in the semioval center, thalamus, basal ganglia and pons and 3) lacunar infarcts [3]. The clinical suspect, complicated by the extreme phenotypic variability, may be supported by the identification of GOM (Granular Osmiophilic Material) on skin biopsy (specificity of 100%, and variable sensitivity from 45% to 100%) [4]. However, only the genetic test can confirm the diagnosis. CADASIL may present with a relapsing-remitting course and with cerebral and brainstem white-matter lesions, being among the several white matter diseases mimicking multiple sclerosis (MS) [5]. The co-occurrence of MS and CADASIL, that are known rarely to co-exist [6], represents a hard challenge for the diagnostic process even for an expert neurologist.

Case 1

A 38 years-old woman came to our attention in April 2011 for rigidity and numbness in right hand and leg. She had no familial history of neurological diseases. Her medical history was unremarkable except for celiac disease, diagnosed in 2010 by esophagogastroduodenoscopy (EGDS) and digiunal biopsy (Marsh IIIC *) [7]. She performed a lumbar puncture that showed an increased cellularity in the absence of oligoclonal bands, and a brain and spinal MRI that showed multiple T2/FLAIR hyperintense lesions in periventricular white matter, semioval centrum, bilateral putamen, frontal white matter and spinal cord with no enhancement after gadolinium contrast. Autoantibody panel for vasculitis resulted negative. Somatosensory and Visual Evoked Potentials were normal, while Motor Evoked Potentials evidenced a delay of central nervous system motor conduction in right leg. When she came at our board, the neurological examination evidenced Babinski sign, brisk tendon reflexes and light spastic hypertonus on her right side. In May 2011, after a gastrointestinal infection, the patient presented acute stiffness in right leg; a new MRI revealed three additional lesions (two cerebral and one new spinal lesions), without contrast enhancement. A diagnosis of MS was posed based on McDonald Criteria 2010 [8] and a therapy with Glatiramer Acetate three times a week was started. In February 2012, the patient presented a new clinical relapse with hypoesthesia on her left side and was given intravenous high dose corticosteroids treatment with subsequent complete regression of symptoms. After three months, she manifested a new relapse characterized by paresthesia in both hands and Lhermitte sign. Spinal MRI showed a new cervical (C4-C5) lesion with contrast-enhancement. Symptoms partially regressed after a new treatment with i.v. corticosteroids. Considering the progression of the disease course, she switched to a second-line therapy with Natalizumab with a good clinical response for three years, without changes in neurologic examination (EDSS 1) and MRI. Following JCV

seroconversion, the patient stopped natalizumab on June 2015 and started dimethylfumarate after 5 weeks. Clinical examinations and MRI were stable in the next years, but an important decrease of lymphocyte count appeared in February 2017, leading to a new switch to interferon beta (IFN) i.m. once a week, which was started in August 2017. In December 2017, her mother had a stroke at the age of 62. The absence of conventional cerebrovascular risk factors and the brain MRI pattern (a recent capsular infarct, associated to multiple lacunar infarcts, severe diffused white matter hyperintensities involving temporal lobes and several microbleeds on gradient echo images – T2 weighted imaging) were strongly suggestive for CADASIL. The suspect was subsequently confirmed by the identification of the heterozygous c.421C>T, p. (Arg141Cys) substitution in NOTCH3 gene (rs1174625611). The same pathogenetic variant was detected in our patient by Sanger sequencing in February 2018. Given this result and the stability of the clinical-MRI conditions, we decided to stop the Disease-modifying therapy (DMT). However, three months later the patient developed hyposthenia in her lower limbs, and her spinal cord MRI showed a new enhancing lesion (Figure 1). This new clinical and radiological activity prompted to perform a new lumbar puncture: we found unmatched oligoclonal bands (OCBs) in the CSF and an elevated Kappa Free Light Chain Index (KFLC) index [9]. Considering the coexistence of MS and CADASIL and the report of side effects to INF beta and acetylsalicylic acid, we started treatment with subcutaneous glatiramer acetate 40 mg three times a week and antiplatelet therapy with clopidogrel 75 mg a day. The patient has been clinically and radiologically stable during the last two years of follow-up, and she remains currently worsening-free.

Case 2

In 2016 a 49-year old healthy man experienced progressive bilateral legs weakness, with gait limitation. He performed a brain MRI showing confluent hyperintensities in T2 sequences in periventricular, deep white matter and basal ganglia areas. After contrast injection, some small lesions got enhancement. Multiple spinal hyperintensities in T2 sequences were found at C2, C2-C3, C4-C5, between D3-D6 and D8-D9, without any enhancement. Clinical and MRI investigations allowed to pose diagnosis of MS, referring to revised McDonald's criteria of 2010 [8]. At that time, the patient presented a EDSS of 4.0 and he started oral therapy with Teriflunomide. Due to radiological disease activity without clinical worsening (stable EDSS 4.0), he shifted to higher efficacy DMT in May 2019, starting fingolimod. Further MRI activity, along with clinical disease progression in July 2020 (EDSS 5.0) and in September 2020 (EDSS 6.0), required further vertical therapy shift to ocrelizumab in 2020. When his dizygotic twin showed a symptomatology characterized by recurrent migraine and memory loss episodes, a brain MRI was performed and a diagnosis of CADASIL was considered. The



molecular analysis of NOTCH3 identified the heterozygous c.3226C>T (p.Arg1076Cys) pathogenic variant in NOTCH3 (rs1438626607), confirming the clinical diagnosis of CADASIL. The case 2 thus came at our neurogenetics unit for a genetic counseling about CADASIL in 2020. He had never reported CADASIL-related clinical features such as migraine, stroke, cognitive impairment, epilepsy, gate disturbances or other symptoms. However, considering his family history, he underwent molecular analysis to investigate the presence of the familial variant on his DNA. As in his brother, the NOTCH3 c.3226C>T (p.Arg1076Cys) substitution was confirmed in the heterozygous state also in our patient. At that point, we decided to deepen his previous MS diagnosis performing a lumbar puncture and a new brain and spine MRI (Figure 2), 5 months far from the last ocrelizumab infusion. The CSF analysis showed the presence of 14 OCB, with a unique intracranial synthesis pattern, and a high G-globulin antibody concentration (308.8 mg/dL, with a normal value < 34.0). Considering the co-occurrence of MS and CADASIL, we decided to continue the DMT with Ocrelizumab and we started an antiplatelet therapy with acetylsalicylic acid 100 mg/die. To date, patient repeated a new brain and spinal MRI in 2021 showing absence of inflammatory activity and is clinically stable (EDSS 6.0).

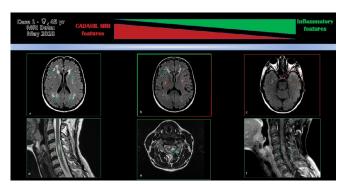


Figure 1: MRI images. T2/FLAIR (a,b,c): a. Centrum Semiovale (CSO); b. Deep basal ganglia; c. Anterior temporal lobe. T2-weighted images (d,e): d. Sagittal section; e. Axial section. T1 post gadolinium-enhancement: f. Sagittal section. Green arrow: inflamation lesions; Red circle: CADASIL footures.

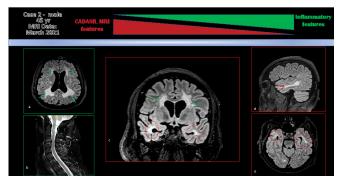


Figure 2: MRI images. T2/FLAIR: a. Centrum Semiovale (CSO); b. Tervical spinal cord; c. CSO and polar lobe. d. Temporal lobe, sagittal section; f. Temporal lobe, axial section. Green arrow: inflammatory lesions; Red circle: CASASIL features.

Conclusion

We observed two cases of CNS inflammatory disease with a concomitant diagnosis of CADASIL. These cases may be difficult to diagnose considering the rarity of CADASIL condition, the poor genotype-phenotype correlation, the heterogeneity of symptoms and age of onset of this inherited disease. The first case of co-occurrence of CADASIL and MS was reported by Carone et al. as an atypical presentation of CADASIL, misdiagnosed with MS [10]. A diagnosis of MS, according to the McDonald's criteria, requires evidence of spatial and temporal dissemination of the disease, but also "no better explanation" for other etiologies [8]. Instead, migraine and familial history are two feature guides for CADASIL diagnostic suspect. In both our cases, the clinical presentation, the MRI pattern and the excellent clinical response to i.v. steroid therapy favored an inflammatory diagnostic hypothesis. No clinical features instead leaned for a CADASIL diagnosis except for the positive familial history. Concerning MRI findings, they were of difficult interpretation for the differential diagnosis. In both cases, some atypical features for MS, like hyperintensities in deep basal ganglia or in anterior temporal lobe, that seemed more typical for CADASIL, co-existed with brain and spinal cord lesions (that are rarely seen in CADASIL [11]), supporting the diagnosis of MS (Figure 1, 2). The description of CADASIL and MS co-existence cases increased in the recent years, opening a debate about a possible relationship between monogenic diseases and inflammatory disease of CNS [12]. Pathogenic hypotheses highlight that genetic defects, like mutation of Notch3 protein [13], or cytopathies such as Leber's disease [14], or POLG mutations [15], could promote demyelinatinglike diseases, suggesting a key role of the exposure of CNS antigens with a consequent self-immune attack [16]. Notably, the function of Notch3 receptor in CADASIL disease showed a consistent interplay with immune system activity. Notch3 appears involved in T cell response regulation, specifically in myelin reactive T-cells, contributing to the encephalitogenic potential of autoreactive T cells [13]. Moreover, its abnormal upregulation showed a pro-inflammatory activity [17]. It is known that the Notch3 receptor activity favors the activation of the Nuclear Factor kappa B (NF-κB), promoting proinflammatory gene expression in activated macrophages, and its inhibition can reduce myelin-activated T-cell response [18]. Taking these data into account, it might be expected that mutations which increase, or do not change, Notch3 activity can predispose to inflammation. Intriguingly, functional studies in CADASIL patient with NOTCH3 p.R1076C mutation showed an abnormal upregulation of pro-inflammatory NF-kB target genes in VSMCs, partly attributed to constitutive activation of Notch3 signalling [17]. This mutation, present in patient of case 2, is localized in the EGFr 27 domain and it has been already described in further 7 unrelated CADASIL families worldwide [17], [19], [20]. Conversely, the more common pathogenic c.421C>T



(p.Arg141Cys) variant in the EGFr 3 domain, identified in our case 1 and in many other families with CADASIL worldwide, causes a reduced expression of the Notch3 receptor on the cell surface, higher intracellular aggregates but no alteration of Notch3 signalling, suggesting that the pathology may be due to abnormal Notch3 trafficking and localization rather than signalling dysfunction [21]. Overall, to our knowledge, this is the first time that inflammatory phenotypic features have been described in patients carrying these two variants. To date it is not possible to determinate if the proinflammatory activity may be promoted only by the pathogenic mutations in NOTCH3 or need the contemporary intervention of other genetic and non-genetic modifier factors. Nevertheless, the relevant role for NOTCH3 in inflammation has been already proven. Maybe the presence of modifier factors could explain the intrafamilial phenotypic variability described in the two reported families. Considering these hypothesis, the "apparent" difference between MS and "inflammatory CADASIL" could be semantic rather than etiologic. In this perspective DMTs used in MS might be beneficial for these patients [7] and, reciprocally, the selective modulation of Notch3-dependent pathways may represent a possible therapeutic option in MS [13].

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