



A Rare Cause of Ataxia: SPG7 Mutation

Nadir Bir Ataksi Nedeni: SPG7 Mutasyonu

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Abstract

Hereditary spastic paraplegias (HSPs) are neurodegenerative disorders characterized by progressive weakness and spasticity in the lower limbs and are clinically and genetically heterogeneous. They are clinically classified as pure and complicated forms. HSP may be inherited as autosomal dominant, autosomal recessive, X-linked, or mitochondrial disorders. A 61-year-old man presented with progressive stiffness, weakness, ataxia in the lower limbs, dysarthria and asymmetric ptosis. None of the family members had such neurological symptoms. Brain magnetic resonance imaging showed cerebellar atrophy. Genetic analysis with whole exome sequencing revealed a homozygous p.Ala572Val (c.1715C>T) exchange mutation in *SPG7* gene. SPG 7 mutation is an important cause of adult-onset undiagnosed ataxia. Although the availability of exome sequencing with targeted analysis helps molecular diagnosis of SPG7 easier, it is important to consider SPG7 in the differential diagnosis of adult onset ataxias. In this paper we report a patient with complicated form of HSP with SPG7 mutation.

Keywords: Hereditary spastic paraplegia, ataxia, SPG7

Öz

Hereditör spastik paraplajiler (HSP) alt ekstremitelerde ilerleyici güçsüzlük ve spastisite ile karakterize nörodejeneratif natürde klinik ve genetik olarak heterojen bir grup hastalıktır. Klinik olarak pür ve komplike formları bulunmaktadır. Genetik olarak da otozomal dominant, otozomal resesif, X'e bağılı ve mitokondriyal olarak kalıtılabilir. Olgumuz 61 yaşında erkek hasta olup bacaklarda ilerleyici güçsüzlük, sertlik ve dengesizlik, konuşmada bozukluk ve göz kapağında düşme ile başvurdu. Ailesinde benzer bir öykü bulunmayan hastanın beyin manyetik rezonans görüntülemesinde serebellar atrofi izlendi. Tüm ekzom dizileme ile *SPG7* geninde homozigot p.Ala572Val (c.1715C>T) mutasyonu saptandı. SPG7 mutasyonu erişkin başlangıçlı tanı koyulamayan ataksinin önemli sebeplerindendir. Her ne kadar tüm ekzom dizileme gibi yeni nesil dizileme yöntemleri ile moleküler olarak daha kolay tanı koyulabilse de erişkin başlangıçlı ataksilerin ayırıcı tanısında SPG7'yi düşünmek gerekir. Bu yazıda SPG7 mutasyonlu komplike HSP formu tanısı alan bir hastayı sunuyoruz.

Anahtar Kelimeler: Hereditör spastik paraplaj, ataksi, SPG7

Introduction

Hereditary spastic paraplegias (HSPs), also known as spastic paraplegias (SPGs) are a group of clinically and genetically heterogeneous neurodegenerative disorders characterized by progressive weakness and spasticity in the lower extremities (1). These key symptoms are due to retrograde degeneration of the longest nerve fibers in the corticospinal tracts and posterior columns (2).

The HSPs are classified clinically as pure and complicated forms. Patients with pure HSP show progressive SPG with hyperreflexia, extensor plantar response, ankle clonus and increased tone in the lower limbs, which can be associated with sphincter disturbances and deep sensory loss. Complicated HSPs consist of SPG and additional neurologic or systemic abnormalities such as cerebellar ataxia, dysarthria, mental retardation, peripheral neuropathy, optic

atrophy, deafness, dementia, severe amyotrophy, epileptic seizures, ichthyosis or thin corpus callosum (1,2,3).

To date, over 70 distinct loci and 50 *SPG* genes have been identified. The mode of inheritance can be autosomal dominant (AD), autosomal recessive (AR), X-linked, or mitochondrial. Also, sporadic HSP is not uncommon (4). The prevalence of HSP is estimated to be 1.8 cases per 100,000 individuals for both ADHSP and ARHSP (5).

Here, we report a case of a Turkish patient with SPG7 mutation, complicated form of HSP and a recessive pattern of inheritance.

Case Reports

Our patient was a 61-year-old right-handed Caucasian male. At age of forty one, he had noticed gait disturbance and slow and slurry speech. Ten years later, he started using a walking stick and

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He already had been screened for vitamin E levels, anti-GAD antibodies, Friedreich ataxia, SCA1, SCA2, SCA3, SCA6, SCA7, SCA17, during his admissions to different hospitals. The results were negative. We carried out whole exome sequencing analysis of genomic DNA from the patient. A homozygous p.Ala572Val (c.1715C>T) exchange mutation was detected in *SPG7* gene.

We diagnosed his disorder as an autosomal-recessive form of hereditary SPG7 based on the characteristic clinical symptoms, MRI findings, and the mutation in *SPG7* gene.

Discussion

SPG7, OMIM#602783 is one of the most common form of ARHSP caused by mutations in the *SPG7* gene. Rare AD transmission has also been described (6). *SPG7* gene is located in chromosome 16q24 and encodes paraplegin protein, a subunit of a m-AAA-protease complex located within the inner mitochondrial membrane (7,8). Paraplegin plays a significant role in different cellular activities including membrane trafficking, protein maturation and degradation (8,9). *SPG7* often results in complicated forms of HSP with cerebellar involvement and/or cerebellar atrophy, optic atrophy, ptosis, supranuclear palsy, ophthalmoplegia, cognitive impairment and dysexecutive syndrome (8,10,11,12,13). Our patient had a complicated form of HSP without cognitive impairment. Although ptosis seen in HSP has been reported as bilateral symmetrical ptosis in the literature, it is rarely asymmetrical as in the case we presented (14). Also, *SPG7* mutations are indicated as a rare cause of AD optic neuropathy and an important cause of sporadic progressive external ophthalmoplegia with multiple mitochondrial DNA mutations (6,15). In 2019, a review including 241 patients with *SPG7* mutation was published. Results of the study states that spasticity-predominant phenotype is related with loss of function variants and that more frequent cerebellar ataxia and later onset is observed in patients carrying at least 1 Ala510Val variant. In this study, neurologic follow up was done only in 98 of these patients. Electromyography was performed only in 23 of the 241 patients and among these 23 patients 20 had sensorimotor axonal polyneuropathy, but in our case nerve conduction studies were normal (16). A recent study showed that *SPG7* gene mutations can be responsible for 18.6% of patients with undiagnosed ataxia presenting in mid-adult life. Ataxia was the predominant feature in these patients and all of them did not have pyramidal signs initially, but during long-term follow-up pyramidal signs were found on repeat examinations (17,18).

Exome sequencing analysis provides a high yield approach for the molecular diagnosis of undiagnosed ataxia. Sun et al. (19) identified pathogenic and suspected diagnostic variants in 88 patients of 170 patients with undiagnosed ataxia with a positive diagnostic rate of 52%. Among mutation positive patients, *SPG7* gene mutation rate was 9% (19). Recent studies which were carried out among different populations showed that *SPG7* mutation was a frequent cause in undiagnosed cerebellar ataxias (17). Thus, *SPG7* should be considered in undiagnosed cerebellar ataxias after ruling out the most common etiologies.

We reported a rare cause of ataxia in a Turkish patient with HSP *SPG7* with clinical features, MRI findings and genetic diagnosis which were achieved by whole exome sequencing in compatible with recent literature.

Ethics

Informed Consent: Written consent was obtained.

Peer-review: Externally peer-reviewed.

Authorship Contributions

Surgical and Medical Practices: A.Ç., B.Ö.B., R.A.İ., Concept: A.Ç., B.Ö.B., Design: B.Ö.B., Data Collection or Processing: A.Ç., Analysis or Interpretation: A.Ç., B.Ö.B., Literature Search: A.Ç., B.Ö.B., Writing: A.Ç.

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