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Mutation of SELENOI and LAMA1 Causes Hereditary Spastic Paraplegia

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Introduction

Hereditary Spastic Paraplegia (HSP) is a rare neurodegenerative disorder characterized by weakness and spasticity in the lower limbs. HSP primarily affects the muscle function in the lower limbs, making everyday movements challenging. It is caused by genetic factors, with various gene mutations contributing to its development. HSP has various forms and often follows an autosomal dominant inheritance pattern. Treatment focuses on symptom management and relief, including physical therapy and the use of assistive devices. While there is no cure, appropriate management strategies can improve patients' quality of life. Additionally, genetic counseling is important for assessing familial risk and preparing necessary precautions. The SELENOI and LAMA1 genes play essential roles in regulating cellular functions, both internally and externally. The SELENOI gene is associated with neuronal function and primarily operates within the cell, regulating intracellular signaling pathways and influencing cell cycle and motor impairment. The LAMA1 gene, on the other hand, controls the formation and structure of the extracellular matrix, impacting neuronal growth and stability. While mutations in these genes are associated with various neurological disorders, there is no direct evidence linking to HSP.

Case report

We report a 19-year-old male diagnosed with HSP. The patient was born at 40 weeks of gestation and showed lower limb weakness and developmental delay one year after birth, undergoing outpatient rehabilitation therapy regularly since then. Despite interventions such as botulinum toxin injection and tenotomy surgery for lower limb spasticity, his condition persisted. Manual muscle testing revealed overall poor grade in the lower limbs, and spasticity was graded as 2 on the Modified Ashworth Scale (Table 2). The modified Barthel Score (MBI) and Functional Independence Measure (FIM) to measure the level of independence in activities of daily living. Berg Balance Scale (BBS), a clinical test of a person's static and dynamic balance abilities, was also administered. The results of these assessments, as presented in Table 3, indicated notable deficiencies in ambulation and a lack of dynamic balance. Next-generation sequencing (NGS) revealed a homozygous variant in the SELENOI gene (NM_033505.4:c.797C>T:p.(Pro266Leu)) and a heterozygous variant in the LAMA1 gene (NM_005559.4:c.4579C>T:p.(Gln1527Ter)) (Table 1). The patient has a 24-year-old female sibling diagnosed with HSP. To confirm genetic association, NGS was performed on her as well, revealing a heterozygous variant in the SELENOI gene and a homozygous variant in the LAMA1 gene. Brain image of the patient and the sibling show diffuse cerebral and cerebellar atrophy (Figure 1).

ne Patient	NGS						MBI		
Gene	Accession	Nucleotide	Amino acid	Zygosity	dbSNP	Inheritance	Personal hygiene		5
ELENOI	NM_033505.4	c.797C>T	p.Pro266Leu	Homo		AR	Bathing self		3
							Feeding Toilet		10 10
AMA1	NM_005559.4	c.4579C>T	p.Gln1527Ter	Hetero	rs746183130	AR	Stair climbing		5
The Sibling NGS							Dressing		8
							Bowel control		10
Gene	Accession	Nucleotide	Amino acid	Zygosity	dbSNP	Inheritance	Bladder control Ambulation		10 8
ELENOI	NM_033505.4	c.797C>T	p.Pro266Leu	Hetero		AR	Chair/bed transfer		15
AMA1	NM 005559.4	c.4579C>T	p.Gln1527Ter	Homo	rs746183130	AR		Total	84
						A100100	FIM		
Table 2.	Modified Ashw	orth Scale						Eating Grooming	77
		Righ	t Left				Self-care	Bathing	4
Sh	ouldor floxor	G0	G0	_			Sell-care	Dressing-Upper	7
Shoulder flexor								Dressing-Lower	6
Shoulder extensor		G0	G0					Toileting	6
Elbow flexor		G0	G0				Sphincter Control	Bladder	7
Elbow extensor		G0	G0				22 • CONTRACTOR AND	Bowel	7
Hip flexor		G1+						Bed, Chair, W/C	6
							Transfers	Toilet	6
Hip extensor		G1+	G1+					Tub, Shower	6
Knee flexor		G1+	G1+				Locomotion	Walk, Wheelchair	1
Knee extensor		G1+	G1+				Locomoton	Stairs	1
Ankle dorsi flexor		G0	G0				Comprehension	Comprehension	7
							Comprehension	Expression	7
Ankle	e plantar flexor	G1	G1					Social Interaction	7
4)			(B)				Social Cognition	Problem Solving Memory	4 7
			6	100	CLP X		BBS Sitting to standing Standing unsupported	Total	101 2 2
P.	SZ (4)	3.1		and y	· m		Sitting unsupported		4
P	S. 18 8	1 1	LF AL	ver 1	1 2 1	St -	Standing to sitting		1
18	2.4	a st			A Star	1	Transfers		3
	ECT.		ia.	Ter .	3 8		Standing with eyes closed Standing with feet together		2
	1 6 3	1	tion (1)	Control 1	1218	10cm	Reaching forward with outstretched	larm	1
	A B	11		0.00	1913		Retrieving object from floor		0
					A CONTRACTOR OF THE	8 %	Turning to look behind Turning 360 degrees		1
:)	*		(D)				Placing alternate foot on stool		0
	1			100	and and		Standing with one foot in front		0
	AUGH	and a la		11		1	Standing on one foot Total		0 16
	5.5	PO.		-	1354		10tai		16
			u			er Nor			
gure 1. Bra id cerebella I-weighted :	in MRI of the patient r atrophy, with thinni sagittal view (D)] sho	[T2-weighted axia ng of the corpus ci ws diffuse cerebra	I view (A), T1-weig allosum. Brain MRI and cerebellar at	hted sagittal vie of the sibling [T ophy, with ventr	w (B)] shows diffusi 2-weighted axial vie iculomegaly.	e cerebral aw (C),			
						Con	clusion		
	ed mutatio		se genes,			he SELNC	DI and LAMA1 genes, exhil	biting typical HSP symptom ns behind HSP onset relate	