

Erratum to: The neurological and ophthalmological manifestations of *SPG4*-related hereditary spastic paraplegia

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There is unfortunately a mistake in the protein notation for one of the *SPG4* mutations in Table 1. The c.1253_1255 delAAG mutation was identified in Patient 5. The amino acid change resulting from this deletion is p.Glu418del, not p.E418fsX198.

The correct table is given in the following page.

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Table 1 Molecular genetic and ophthalmological features of the *SPG4* patient cohort

Patient	Sex	Age (years)	<i>SPG4</i> mutation		BCVA RE-LE	Optic discs/OCT measurements	Eye movements	Visual electrophysiology
			Exon	cDNA change/consequence				
1	F	31	5	c.743C>G/p.S245X	20/20-20/20	Normal/no RNFL thinning	Normal	Normal
2	M	53	5	c.743C>G/p.S245X	20/20-20/20	Normal/no RNFL thinning	Horizontal SWJ/saccadic pursuit	Normal
3	F	50	6	c.937delG/p.D313fsX1	20/20-20/20	Normal/no RNFL thinning	Normal	Normal
4	F	55	4–17	del exon 4-17/large-scale deletion	20/20-20/20	Normal/no RNFL thinning	Normal	Normal
5	F	29	10	c.1253_1255delAAG/p.Glu418del	20/20-20/20	Normal/no RNFL thinning	Horizontal SWJ/saccadic pursuit	Normal
6	F	25	11	c.1442_1443insA/p.V482fsX5	20/20-20/20	Normal/no RNFL thinning	Normal	Normal
7	F	55	11	c.1442_1443insA/p.V482fsX5	20/20-20/20	Normal/no RNFL thinning	Horizontal SWJ/saccadic pursuit	Normal
8	F	49	11	c.1414G>A/p.V472I	20/20-20/20	Normal/no RNFL thinning	Normal	Normal
9	F	72	11	c.1384A>G/p.K462E	20/60-20/30	Normal/no RNFL thinning	Normal	Normal
10	M	65	11	c.1081C>A; c.1082T>A/p.L361N	20/20-20/20	Normal/no RNFL thinning	Normal	Normal

BCVA best-corrected visual acuities, *cDNA* complementary DNA, *LE* left eye, *OCT* optical coherence tomography, *RE* right eye, *RNFL* retinal nerve fibre layer, *SWJ* square wave jerks