

Title: Childhood Ataxia with Central Nervous System Hypomyelination/Vanishing White Matter *GeneReview* Table 3  
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Note: The following information is provided by the authors and has not been reviewed by *GeneReviews* staff.

Table 3. Pathologic Allelic Variants

Gene	Exon/Intron	Mutation	Allelic Percentage	Protein Mutation
EIF2B1	Intron 2	IVS2+1G>A	<1%	S84ins22aa, stop
	Exon 6	A622T	<1%	N208Y
	Exon 7	A622T		N208Y
EIF2B2	Exon 4	C512T	<1%	S171F
	Exon 4	C547T	<1%	R183fs
	Exon 4	548Gdel	<1%	R183fs
	Exon 4	C586T		P196S
	Exon 5	G599T	<1%	G200V
	Exon 5	G599C	<1%	G200A
	Exon 5	607-612delins TG	<1%	M203fs
	Exon 5	A638G	8.2%	E213G
	Exon 6	A818G	<1%	K273R
	Exon 7	C871T		P291S
	Exon 8	G910T	<1%	E304X
	Exon 8	T947A	<1%	V316D
	Exon 8	G986T	<1%	G329V
EIF2B3	Exon 1	G136A	<1%	V46I
	Exon 2	C260T	<1%	A87V
	Exon 3	A407C	<1%	Q136P
	Exon 6	G674A	<1%	R225W
	Exon 8	T4023G	<1%	H341Q
	Exon 9	11931195delTG	<1%	V398fs
EIF2B4	Exon 7	C683T	<1%	A228V
	Exon 7	G626A	1.5%	R209Q
	Exon 7	C625T	<1%	R209X
	Exon 8	C728T	3%	P243L
	Exon 9	T806G		L269R
	Exon 11	C1069T		R357W
	Exon 11	G1070A		R357Q

Gene	Exon/Intron	Mutation	Allelic Percentage	Protein Mutation
	Exon 11	G1091T		R364Q
	Exon 11	C1120T	1.1%	R374C
	Exon 11	C1172A	<1%	A391D
	Intron 11	IVS11+1G>A	<1%	E397ins11aa
	Exon 13	T1393C	<1%	C465R
	Exon 13	C1447T	<1%	R483W
	Exon 13	T1465C	1.1%	Y489H
EIF2B5	Exon 1	C47A	<1%	A16D
	Exon 1	G161C	<1%	R54P
	Exon 1	T166G	<1%	F56V
	Exon 2	T203C		L68S
	Exon 2	T218G	1.5%	V73G
	Exon 2	G220A		A74T
	Exon 2	G241A	<1%	E81K
	Exon 2	A271G	7.1%	T91A
	Exon 2	A318T	<1%	L106F
	Exon 3	G338A	27.4%	R113H
	Exon 3	C337T	<1%	R113C
	Exon 3	C406T	<1%	R136C
	Exon 3	453-454del		Y152fsX12
	Exon 4	C545T	<1%	T182M
	Exon 4	G584A	2.2%	R195H
	Exon 4	C583T	<1%	R195C
	Exon 4	G592A	<1%	E198K
	Exon 6	792delTinsACA	<1%	F264fs
	Exon 6	G806T	<1%	R269L
	Exon 6	C805G		R269G
	Exon 6	G806A		R269Q
	Exon 7	G896A	<1%	R299H
	Exon 7	G925C	1.5%	V309L
	Exon 7	G929T		C310F
	Exon 7	C943G	<1%	R315G
	Exon 7	G944A	<1%	R315H
	Exon 7	C943T	<1%	R315C
	Exon 7	C967T	<1%	P323S

Gene	Exon/Intron	Mutation	Allelic Percentage	Protein Mutation
	Exon 7	T1003C		C335R
	Exon 7	C1015T	3.4%	R339W
	Exon 7	G1016C	<1%	R339P
	Exon 7	G1016A	<1%	R339Q
	Exon 7	G1016T		R339P
	Exon 7	A1028G	<1%	Y343C
	Exon 7	A1153G	<1%	I385V
	Exon 8	G1157T	<1%	G386V
	Exon 8	A1160G	<1%	D387G
	Exon 8	C1264T	<1%	R422X
	Exon 8	T1274G	<1%	L425R
	Exon 8	C1280T	<1%	P427L
	Exon 8	T1289C	<1%	V430A
	Exon 9	C1340T	<1%	S447L
	Exon 9	G1444ins17	<1%	G481fs493X
	Exon 9	G1459A		E487K
	Exon 10	A1484G	<1%	Y495C
	Exon 13	1813delC		L605fs
	Exon 14	T1882C	<1%	W628R
	Exon 14	G1884A	<1%	W628X
	Exon 14	G1948A	1.2%	E650K
	Exon 14	1996del21	<1%	664del7

[www.vumc.nl/whitematter](http://www.vumc.nl/whitematter)

Mutations can affect all regions of the five eIF2B subunits. From Leegwater et al [2001], van der Knaap et al [2002], van der Knaap et al [2003], Fogli et al [2004b], Ohtake et al [2004], Ohlenbusch et al [2005], Fogli & Boespflug-Tanguy [2006], Pronk et al [2006], Scali et al [2006], Matsui et al [2007].

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