

Title: MEGDEL Syndrome *GeneReview* Table 4

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Note: The following information is provided by the authors listed above and has not been reviewed by *GeneReviews* staff.

Table 4. Selected *SERAC1* Pathogenic Variants

DNA Nucleotide Change	Protein Amino Acid Change	Reference	Reference Sequences
c.442C>T	p.Arg148Ter	Wortmann et al [2012b]	NM_032861.3 NP_116250.3
c.466_467insGCGGGAAATGT	p.Ser156fsCysTer16		
c.576delT	p.Leu193SerfsTer9		
c.1167_1170delTCAG	Altered splicing		
c.1202G>A	p.Gly401Asp		
c.1211G>A	p.Gly404Glu		
c.1309_1313dupACATG	p.Trp438Ter		
c.1403+1G>C	Altered splicing		
c.1435_1437delCTT	p.Leu479del		
c.1493G>C	p.Ser498Thr		
c.1598_1599dupATAGTGTCCCTCATCAT	p.Gly536IlefsTer56		
c.1627_1628dupTC	p.Val544LeufsTer43		
c.1822_1828+10delTCAGCAGGTATTCACTCinsACCAACAGG	Altered splicing		
c.1924C>T	p.Gln642Ter	Sarig et al [2013]	
c.698-9TG>AGTGATA	p.Leu233Ter		
c.128+4A>G	Altered splicing		
c.202C>T	p.Arg68Ter	Tort et al [2013]	

Note on variant classification: Variants listed in the table have been provided by the authors. *GeneReviews* staff have not independently verified the classification of variants.

References

Sarig O, Goldsher D, Nousbeck J, Fuchs-Telem D, Cohen-Katsenelson K, Iancu TC, Manov I, Saada A, Sprecher E, Mandel H. Infantile mitochondrial hepatopathy is a cardinal feature of MEGDEL syndrome (3-methylglutaconic aciduria type IV with sensorineural deafness, encephalopathy and Leigh-like syndrome) caused by novel mutations in SERAC1. *Am J Med Genet A*. 2013;161:2204-15.

Tort F, García-Silva MT, Ferrer-Cortès X, Navarro-Sastre A, Garcia-Villoria J, Coll MJ, Vidal E, Jiménez-Almazán J, Dopazo J, Briones P, Elpeleg O, Ribes A. Exome sequencing identifies a new mutation in SERAC1 in a patient with 3-methylglutaconic aciduria. *Mol Genet Metab*. 2013;110:73-7.

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