Title: ALS2-Related Disorders GeneReview, Table 3

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Updated: January 2016

Note: The following information was provided by the original authors (Bertini ES, Eymard-Pierre E, Boespflug-Tanguy O, Cleveland DW, Yamanaka K) and was not reviewed by *GeneReviews*

staff.

Table 3. Selected ALS2 Pathogenic Allelic Variants

| DNA Nucleotide Change (Alias ¹) | Protein Amino Acid Change | Reference Sequences ² | Pheno- type | References |
|------------------------------------|---|-------------------------------------|----------------|--|
| c.138delA (A261del) | p.Ala47ProfsTer4 | NM 020919.2 NP_065970.2 | JALS | Hadano et al [2001], Yang et al [2001] |
| c.470G>A | p.Cys157Tyr | | IAHSP | Eymard-Pierre et al [2006] |
| c.553delA | p.Thr185fsTer5 | | JALS | Kress et al [2005] |
| c.1007_1008delTA | p.lle336ThrfsTer5 | | IAHSP | Eymard-Pierre et al [2002] |
| c.1425_1426delAG (1548delAG) | p.Glu476GlyfsTer71 | | JPLS | Hadano et al [2001], Yang et al [2001] |
| c.1472_1481delTTTCCCC CAG | p.Val491GlyfsTer3 | | IAHSP | Eymard-Pierre et al [2002] |
| c.1619G>A | p.Gly540Glu | | JPLS | Panzeri et al [2006] |
| C;.1825_1826insCAGTG/ c.3529G>T | p.Glu609fsTer9/ p.Gly1177Ter | | IAHSP | Sztriha et al [2008] |
| c.1867_1868delCT | p.Leu623ValfsTer24 | | JPLS | Yang et al [2001] |
| c.2143C>T | p.Gln715Ter | | IAHSP | Verschuuren-Bemelmans et al [2008] |
| IVS9-2A>T | p.Glu724fsTer32 | | IAHSP | Herzfeld et al [2008] |
| c.2537_2538delAT (2660delAT) | p.Asn845llefsTer13 | | IAHSP | Eymard-Pierre et al [2002] |
| c.2980-2A>G | p.993fsTer7 | | JPLS | Mintchev et al [2009] |
| c.2992C>T (C3115T) | p.Arg998Ter | | IAHSP | Devon et al [2003] |
| c.3565delG / IVS22+5G>C | p.Val1189TrpfsTer19/ p.Gly1172GlufsTer29 | | JALS | Shirakawa et al [2009] |
| c.3619delA (3742delA) | p.Met1206Ter | | IAHSP | Eymard-Pierre et al [2002] |
| c.4721delT (4844delT) | p.Val1574AlafsTer44 | | IAHSP | Gros-Louis et al [2003] |

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

Note on nomenclature: *GeneReviews* follows the standard naming conventions of the Human Genome Variation Society (www.hgvs.org). See Quick Reference for an explanation of nomenclature.

- 1. Variant designations that do not conform to current naming conventions
- 2. Reference sequence (www.ncbi.nlm.nih.gov/Genbank/index.html)

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