



SLC29A3 (NM_018344.6) - cDNA - 2024-05-05

AGTGGTCCTG GCCGTGCC GGAGGCAGCG GCGGCGTGGC GCAGCGGCGA **-2**
CATGCCGTT GTCTCAGAGG ACGACTTTCA GCACAGTTCA AACTCCACCT **49** **M1?** **V4D S13*** **S15Pfs*86**
ACAGAACCAC AAGCAGCAGT CTCCGAGCTG ACCAGGAGGC ACTGCTTGAG **99** **R18G**
AAGCTGCTGG ACCGCCGCC **CC**CTGGCCTG CAGAGGCC**G** AGGACCGCTT **149** **P41Lfs*60 E47Rfs*54**
CTGTGGCACA TACATCATCT TCTTCAGCCT **GGG**CATTGGC AGTCTACTGC **199** **G61V**
CATGGAACCTT CTTTATCACT GCCAAGGAGT ACTGGATGTT **CAA**A**T**CCGC **249** **K81Nfs**
AACTCCTCCA GCCCAGGCCAC CGGGGAGGAC CCTGAGGGCT CAGACATCCT **299**
GAACTACT**TT** GAGAGCTACC TTGCC**G**TTGC CTCCACCGTG CCCTCCA**T**GC **349** **c.302_303insCTACTTTGAGAGCTACC F103X V109I M116R**
TGTGCCTGGT GGCCA**ACT**TC CTGCTTG**TC** ACAGGGTTGC AGTCCACATC **399**
CGTGTCTGG CCTCA**T**GAC GGT**CAT**CTTG GCCATCTCA TG**GT**GATAAC **449** **R133C R134H L139P**
TGC**ACT**GGTG AAGGTGGACA CTTCCCTCTG GACCCGTGGC TTTTTGCGG **499**
TCACCATTGT CTGC**AT**GGTG ATC**CT**CAGCG G**T**GC**CT**CC**A**C TGT**CT**TCAGC **549** **T180A**
AGCAG**CAT**CT AC**GG**CATGAC CGG**CT**CC**TT** CCTATGAGGA ACTCCCAGGC **599** **S184R**
ACTGATA**T**CA GGAGGAGCCA TGG**GC**GG**AC** G**GT**CAGCG**CC** GTGG**CC**T**CAT** **649** **S203P c.625G>A**
TGGTGGACTT GG**CT**GC**AT**CC AGT**G**A**T**GTGA GGA**AC**A**G****CC** CCTGG**CC**TT**C** **699** **D225G A230T**
TT**CCTG****A****CG** CC**ACT**GT**CT**TT CCTCG**TG**CTC TGC**AT**GG**AC** T**CT**AC**CT**G**CT** **749** **T236M T236T**
G**CT**GT**CC****AG** CTGGAGTATG CC**AG**GT**ACT**TA CATGAG**GC**CT GTT**CT**TG**CG****G** **799**
CC**CAT**GT**GT**TT TT**CT**GG**GT**AA GAGGAG**CT**TC CCCAG**GG**ACTC CCTCAG**GT**GC **849**
C**CT**TC**GG**TGG CCTCCAGATT CATTGATT**CC** CACACAC**CC** CCT**CC**GC**CC** **899**
CAT**CCT**GAAG AAGACGG**CC** GA**C**CTGG**GG**CTT CTG**T**GT**C**ACC **TAC**GT**C**T**T**CT **949** **C310X Y314Tfs F316Y**
TCATCAC**CC**AG CCT**CAT**CT**T**AC **CC**CG**CC****A**T**CT** GC**AC**CAAC**AT** CGAGT**CC**CT**C** **999** **P324S P324L I326V**
AAC**A**GG**GT**TT CGGG**CT**CA**CT** GTGG**AC**CC**AC** AAC**TT**TT**CA** T**CCCC****C**T**CA**C **1049** **c.1045delC**

TACCTTCCTC CTGTACAAC T TGCTGACCT ATGTGGCGG CAGCTCACCG 1099 c.1087C>T R363Q A367T
CCTGGATCCA GGTGCCAGGG CCCAATAGCA AGGCCTCCC AGGGTTCTG 1149
CTCCTCCGA CCTGCCTCAT CCCCCTCTTC GTGCTCTGTA ACTACCAGCC 1199 R386W R386Q c.1172C>A
CCGCGTCCAC CTGAAGACTG TGGTCTTCCA GTCCGATGTG TACCCCGCAC 1249 D412N
TCCTCAGCTC CCTGCTGGGG CTCAGCAACG GCTACCTCAG CACCCTGGCC 1299 c.1269_1270delinsA G427S T431I
CTCCTCTACG GGCCTAACAGAT TGTGCCAGG GAGCTGGCTG AGGCCACGGG 1349 G437R E444X c.1339G>A T449R
AGTGGTGATG TCCTTTATG TGTGCTTGGG CTTAACACTG GGCTCAGCCT 1399
GCTCTACCCCT CCTGGTGCAC CTCATCTAGA AGGGAGGACA CAAGGACATT *21
GGTGCTTCAG AGCCTTGAA GATGAGAAGA GAGTGCAGGA GGGCTGGGGG *71
CCATGGAGGA AAGGCCTAAA GTTTCACTTG GGGACAGAGA GCAGAGCACA *121
CTCGGGCCTC ATCCCTCCCAGATGCCAGT GAGCCACGTC CATGCCATT *171
CCGTGCAAGG CAGATATTCC AGTCATATTA ACAGAACACT CCTGAGACAG *221
TTGAAGAAGA AATAGCACAA ATCAGGGTA CTCCCTTCAC AGCTGATGGT *271
TAACATTCCA CCTTCTTTCT AGCCCTTCAA AGATGCTGCC AGTGTGCC *321
CTAGAGTTAT TACAAAGCCA GTGCCAAAC CCAGCCATGG GCTCTTGCA *371
ACCTCCCAGC TGCGCTCATT CCAGCTGACA GCGAGATGCA AGCAAATGCT *421 c.1893G>A
CAGCTCTCCT TACCCCTGAAG GGGTCTCCCT GGAATGGAAG TCCCCTGGCA *471
TGGTCAGTCC TCAGGCCAA GACTCAAGTG TGCACAGACC CCTGTGTTCT *521
GTGGGTGAAC AACTGCCAC TAACCAGACT GGAAAACCCA GAAAGATGGG *571
CCTTCCATGA ATGCTTCATT CCAGAGGGAC CAGAGGGCT CCCTGTGCAA *621
GGGATCAAGC ATGCTGGCC TGGGTTTCA AAAAAGAGG GATCCTCATG *671
ACCTGGTGGT CTATGGCCTG GGTCAAGATG AGGGTCTTC AGTGTCCCTG *721
TTTACAACAT GTCAAAGCCA TTGGTTCAAG GGCGTAATAA ATACTTGCCTG *771
ATTCAA

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