



SLC29A3 (NM\_018344.6) - cDNA - 2026-07-11

AGTGGTCCTG GCCGTGCGCC GGAGGCAGCG GCGGCGTGGC GCAGCGGCCA -2  
CATGGCCGTT GTCTCAGAGG ACGACTTTCA GCACAGTTCA AACTCCACCT 49 M1? V4D S13\* S15Pfs\*86  
ACAGAACCAC AAGCAGCAGT CTCCGAGCTG ACCAGGAGGC ACTGCTTGAG 99 R18G R25\*  
AAGCTGCTGG ACCGCCC GCCCTGGCCTG CAGAGGCCCG AGGACCGCTT 149 P41Lfs\*60 E47Rfs\*54  
CTGTGGCACA TACATCATCT TCTTCAGCCT GGGGCATTGGC AGTCTACTGC 199 G61V  
CATGGAACCTT CTTTATCACT GCCAAGGAGT ACTGGATGTT CAAACTCCGC 249 K81Nfs  
AACTCCTCCA GCCCAGCCAC CGGGGAGGAC CCTGAGGGCT CAGACATCCT 299  
GAACTACTTT GAGAGCTACC TTGCCGTTGC CTCCACCGTG CCCTCCATGC 349 c.302\_303insCTACTTTGAGAGCTACC F103X V109I M116R  
TGTGCCTGGT GGCCAACCTC CTGCTTGTCA ACAGGGTTGC AGTCCACATC 399  
CGTGTCTTGG CCTCACTGAC GGTCATCCTG GCCATCTTCA TGGTGATAAC 449 R133C R134H L139P  
TGCACCTGGTG AAGGTGGACA CTTCTCCTG GACCCGTGGC TTTTTTGCGG 499  
TCACCATTGT CTGCATGGTG ATCCTCAGCG GTGCCCTCCAC TGTCTTCAGC 549 T180A  
AGCAGCATCT ACGGCATGAC CGGCTCCTTT CCTATGAGGA ACTCCCAGGC 599 S184R  
ACTGATATCA GGAGGAGCCA TGGGCGGGAC GGTCAGCGCC GTGGCCTCAT 649 S203P c.625G>A  
TGGTGGACTT GGCTGCATCC AGTGAATGTGA GGAACAGCGC CCTGGCCTTC 699 D225G A230T  
TTCTGACGG CCACTGTCTT CCTCGTGCTC TGCATGGGAC TCTACCTGCT 749 T236M T236T  
GCTGTCCAGG CTGGAGTATG CCAGGTACTA CATGAGGCCT GTTCTTGCGG 799 R258T  
CCCATGTGTT TTCTGGTGAA GAGGAGCTTC CCCAGGACTC CCTCAGTGCC 849  
CCTTCGGTGG CCTCCAGATT CATTGATTCC CACACACCCC CTCTCCGCCC 899  
CATCTTGAAG AAGACGGCCA GCCTGGGCTT CTGTGTCCACC TACGTCTTCT 949 C310X Y314Tfs F316V  
TCATCACCAG CCTCATCTAC CCCGCCATCT GCACCAACAT CGAGTCCCTC 999 P324S P324L I326V  
AACAAAGGGTT CGGGCTCACT GTGGACCACC AAGTTTTTCA TCCCCCTCAC 1049 c.1045delC

TACCTTCCTC CTGTACAAC T TGCTGACCT ATGTGGCCGG CAGCTCACCG 1099 c.1087C>T R363Q A367T  
CCTGGATCCA GGTGCCAGGG CCCAATAGCA AGGCGCTCCC AGGGTTCGTG 1149  
CTCCTCCGGA 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 GGCCTAAGAT TGTGCCCAGG GAGCTGGCTG AGGCCACGGG 1349 G437R E444X c.1339G>A T449R  
AGTGGTGATG TCCTTTTATG TGTGCTTGGG CTTAACACTG GGCTCAGCCT 1399  
GCTCTACCCT CCTGGTGCAC CTCATCTAGA AGGGAGGACA CAAGGACATT \*21  
GGTGTTCAG AGCCTTTGAA GATGAGAAGA GAGTGCAGGA GGGCTGGGGG \*71  
CCATGGAGGA AAGGCCATAA GTTTCACCTG GGGACAGAGA GCAGAGCACA \*121  
CTCGGGCCTC ATCCCTCCCA AGATGCCAGT GAGCCACGTC CATGCCCATT \*171  
CCGTGCAAGG CAGATATTCC AGTCATATTA ACAGAACACT CCTGAGACAG \*221  
TTGAAGAAGA AATAGCACAA ATCAGGGGTA CTCCCTTCAC AGCTGATGGT \*271  
TAACATTCCA CCTTCTTTCT AGCCCTTCAA AGATGCTGCC AGTGTTCGCC \*321  
CTAGAGTTAT TACAAAGCCA GTGCCAAAAC CCAGCCATGG GCTCTTTGCA \*371  
ACCTCCCAGC TGCCTCAT T CCAGCTGACA GCGAGATGCA AGCAAATGCT \*421 c.1893G>A  
CAGCTCTCCT TACCCTGAAG GGGTCTCCCT GGAATGGAAG TCCCCTGGCA \*471  
TGGTCAGTCC TCAGGCCCAA GACTCAAGTG TGCACAGACC CCTGTGTTCT \*521  
GTGGGTGAAC AACTGCCAC TAACCAGACT GGAAAACCCA GAAAGATGGG \*571  
CCTTCCATGA ATGCTTCATT CCAGAGGGAC CAGAGGGCCT CCCTGTGCAA \*621  
GGGATCAAGC ATGCTTGCC TGGGTTTTCA AAAAAAGAGG GATCCTCATG \*671  
ACCTGGTGGT CTATGGCCTG GGTCAAGATG AGGGTCTTTC AGTGTTCCCTG \*721  
TTTACAACAT GTCAAAGCCA TTGGTTCAAG GGCCTAATAA ATACTTGCCT \*771  
ATTCAA

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