



WDR1 (NM_017491.5) - cDNA - 2024-05-15

ACTCCACTCC CGGCACGCCC GGTGCCGCCT TCCGGCTCCA GTCCCCGGGC -84
TCGGCCTCGG CGAGGTGTAA TTCGCAGCGC GGGCCGGCCC CGGAGGCTCT -34
CGGCGAGCGC GGC GCGGTAA CAAGTGGGCG AGGATGCCGT ACGAGATCAA 17
GAAGGTGTTT GCCAGCCTCC CGCAGGTGGA GAGGGGCGTC TCCAAGATCA 67 K7del
TCGGCGGC GA CCCTAAGGGC AACAATTTTC TGTACACCAA TGAAAAGTGC 117 D26N
GTCATCCTAA GGAACATCGA CAACCCAGCC CTTGCTGACA TCTACACAGA 167
GCACGCCCAT CAGGTGGTGG TGGCCAAGTA TGCGCCCAGC GGATTCTACA 217
TTGCCTCCGG AGATGTGTCT GGAAGCTGA GGATCTGGGA TACCACGCAG 267
AAGGAGCACC TGTTGAAGTA TGAGTACCAG CCTTTCGCTG GGAAGATCAA 317
AGACATTGCT TGGACTGAAG ACAGTAAGAG GATCGCCGTG GTCGGGGAAG 367 G121R
GAAGGGAGAA GTTTGGAGCA GTCTTCTCT GGGATAGTGG CTCTTCTGTG 417
GGCGAGATTA CAGGACA CAAAGTCATC AACAGCGTGG ACATCAAGCA 467 H145Q
GAGCCGGCCA TACCGGCTGG CCACGGGAAG CGATGATAAC TGC GCGGCAT 517
TCTTTGAGGG ACCCCCATTC AAGTTCAAGT TCACAATTGG CGACCACAGC 567
CGCTTTGTCA ACTGTGTGCG ATTCTCTCCT GATGGGAACA GATTTGCCAC 617
AGCCAGTGCT GACGGCCAGA TATACATCTA TGACGGGAAG ACTGGGGAGA 667
AGGTGTGCGC GCTGGGCGGA AGCAAGGCC ACGACGGTGG GATTACGCA 717
ATTAGTTGGA GTCCCGACAG CACCCATTTG CTTTCTGCTT CTGGGGACAA 767
AACTTCCAAG ATTTGGGACG TCAGCGTGAA CTCCGTGGTC AGCACATTTT 817
CCATGGGCTC CACGGTTCTG GACCAGCAGC TGGGCTGCC TATGGCAGAAG 867 L286V
GACCACCTG TCAGTGTCTC CCTGTCCGGG TACATCAACT ATCTGGACAG 917 L293E
AAACAACCCC AGCAAGCCCC TGCACGTCAT CAAGGGTCAC AGTAAATCGA 967

TCCAGTGTCT GACGGTGCAT AAAAACGGCG GCAAGTCCTA CATTTACTCT 1017
GGGAGCCACG ACGGACACAT TAATTACTGG GATTCAGAGA CGGGGGAGAA 1067
CGACTCCTTC GCTGGGAAAG GCCACACGAA CCAGGTGTCC AGGATGACCG 1117
TGGATGAGTC GGGGCAGCTC ATCAGCTGCA GCATGGACGA CACCGTGCGG 1167
TACACCAGCC TCATGCTGCG GGA CTACAGC GGACAAGGAG TTGTGAAACT 1217
GGACGTTTCA CCAAAGTGCG TAGCCGTCGG CCCC GGGGGA TACGCCGTGG 1267
TCGTGTGCAT TGGACAGATT GTCCTGCTGA AGGATCAGAG GAAGTGCTTC 1317 [V424M](#)
AGCATCGACA ACCCCGGCTA CGAGCCCGAA GTTGTGGCAG TGCACCCCGG 1367
CGGGGACACG GTGGCAATTG GGGGTGTGGA CGGCAACGTC CGCCTGTATT 1417
CCATCCTGGG CACCACGCTG AAGGATGAGG GCAAGCTCCT AGAGGCCAAG 1467
GGCCCCGTGA CCGACGTGGC CTACTCCAC GACGGCGCCT TCCTCGCGGT 1517
GTGCGACGCC AGCAAGGTGG TCACAGTGT CAGCGTTGCT GACGGCTACT 1567
CGGAGAACAA TGT TTTTAT GGACACCATG CAAAAATCGT CTGCCTGGCC 1617
TGGTCCCCAG ACAATGAACA CTTTGCCTCC GGTGGCATGG ACATGATGGT 1667
GTATGTTTGG ACCCTGAGTG ACCCGGAAAC CAGAGTCAAG ATCCAAGATG 1717
CACACGGCT GCACCATGTC AGCAGCCTGG CCTGGCTGGA CGAGCACACG 1767
CTGGTCACGA CCTCCCATGA TGCCTCTGTC AAGGAGTGG CAATCACCTA 1817
CTGAGGAGCC CCACCCCGC CTCTGGATGG ACCGAATCAG GGA CTAGAGT *46
TTAACTGCAG CGGAACATGT CATTTCTCTA TTTCTGTGAC GCGCCCCAT *96
GCCCCACCC CACCACAAGA GGCAGGAGGG CCCAGTCATG ACCCTCGTCT *146
CTGCAGGGTG TCTGTACACG TTCTTCTGAA AGCTTTAGAC AGTAACAGTT *196
TGCACATGAA AAATAAAGCG AGCACCTAAA CAATGTGTGG AGCATAACTA *246
AAACCCACAG CCCAACCAA CCTTGAGAAT GCGAAACATT CCAGAGGCAG *296
TAGCCTCCAA AGCACACAGA GCCCTGGCC CCGCCGCGGC TCTCACTATC *346
TGTCAGGGGA GGTGTGACAG GTGAATGAGC CGGGGGGCTC ATGTTCCCTG *396
CTGCAGAACA TTTCTGTACT AGTGAGAAGA GGAATATGC ATTGCAGTTC *446
AGCAAAGCCG GAATTCTGTG TTGAACAGAT GTCTGTCTCC CTAGTGTGTG *496
ACTCACACCT TGTGGCTGCC TTCAGAGCG CACCTCCAGA TCAGATGGGG *546

ACACACAACC CCTGGATATG TTTCATTGTC AGATTTTGTG CTTGATTTTA *596
AGAATGGAAT TGTGGGTATC TTTCCTTTTT TTTAATGTA TCTTAACTGT *646
TGCCTGTCAG TGTTTACAAA CTAGTGCCTT GACGGCACCG TGTCCAAGTT *696
TTTAGAACCC TTGTTAGCCA GACCGAGGTG TCCTGGTCAC CGTTTCACCA *746
TCATGCTTTG ATGTTCCCCT GTCTTTCCCT CTTCTGCTCT CAAGAGCAAA *796
GGTTAATTTA AGGACAAAGA TGAAGTCACT GTAAACTAAT CTGTCATTGT *846
TTTTACCTTC CTTTTCTTTT TCAGTGCAGA AATTAAAAGT AAGTATAAAG *896
CACCGTGATT GGGAGTGTTT TTGCGTGTGT CGGAATCACT GGTAAATGTT *946
GGCTGAGAAC AATCCCTCCC CTTGCACTTG TGAACAACTT TTGAGCGCTT *996
TAAGAGATTA GCCTGAGAAA TAATTAAATA TCTTTTCTCT TCA

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