



WDR1 (NM_017491.5) - cDNA - 2025-06-03

ACTCCACTCC CGGCACGCC 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 CCTAAGGGC AACAAATTTTC TGTACACCAA TGGAAAGTGC 117 D26N
GTCATCCTAA GGAACATCGA CAACCCAGCC CTTGCTGACA TCTACACAGA 167
GCACGCCCAT CAGGTGGTGG TGGCCAAGTA TGCGCCCAGC GGATTCTACA 217
TTGCCCTCCGG AGATGTGTCT GGAAGCTGA GGATCTGGGA TACCACGCAG 267
AAGGAGCACC TGTTGAAGTA TGAGTACCAG CCTTTCGCTG GGAAGATCAA 317
AGACATTGCT TGGACTGAAG ACAGTAAGAG GATCGCCGTG GTCGGGAAG 367 G121R
GAAGGGAGAA GTTTGGAGCA GTCTTCTCTT 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 GATTTACGCA 717
ATTAGTTGGA GTCCCGACAG CACCCATTTG CTTTCTGCTT CTGGGGACAA 767
AACTTCCAAG ATTTGGGACG TCAGCGTGAA CTCCGTGGTC AGCACATTTT 817
CCATGGGCTC CACGGTTCTG GACCAGCAGC TGGGCTGCC TATGGCAGAAG 867 L286V
GACCACCTGC TCAGTGTCTC CCTGTCCGGG TACATCAACT ATCTGGACAG 917 L293F
AAACAACCCC AGCAAGCCCC TGCACGTCAT CAAGGGTCAC AGTAAATCGA 967

TCCAGTGTCT GACGGTGCAT AAAAACGGCG GCAAGTCCTA CATTACTCT 1017
GGGAGCCACG ACGGACACAT TAATTACTGG GATTCAGAGA CGGGGGAGAA 1067
CGACTCCTTC GCTGGGAAAG GCCACACGAA CCAGGTGTCC AGGATGACCG 1117
TGGATGAGTC GGGGCAGCTC ATCAGCTGCA GCATGGACGA CACCGTGCGG 1167
TACACCAGCC TCATGCTGCG GGACTACAGC GGACAAGGAG TTGTGAAACT 1217
GGACGTTTTCAG CCAAAGTGCG TAGCCGTCGG CCCCAGGGGA TACGCCGTGG 1267 [Y422M](#)
TCGTGTGCAT TGGACAGATT GTCCGTCTGA AGGATCAGAG GAAGTGCTTC 1317 [Y424M](#)
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 TACGCGTTGCT GACGGCTACT 1567
CGGAGAACAA TGTTTTTTAT GGACACCATG CAAAAATCGT CTGCCTGGCC 1617
TGGTCCCCAG ACAATGAACA CTTTGCCCTC GGTGGCATGG ACATGATGGT 1667
GTATGTTTGG ACCCTGAGTG ACCCGGAAAC CAGAGTCAAG ATCCAAGATG 1717
CACACCGGCT GCACCATGTC AGCAGCCTGG CCTGGCTGGA CGAGCACACG 1767
CTGGTCACGA CCTCCCATGA TGCTCTGTGTC AAGGAGTGG CAATCACCTA 1817 [A597V](#)
CTGAGGAGCC CCACCCCGC CTCTGGATGG ACCGAATCAG GGACTAGAGT *46
TTAACTGCAG CGGAACATGT CATTCTCTA 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 GCCCCTGGCC CCGCCGCGGC TCTCACTATC *346
TGTCAGGGGA GGTGTGACAG GTGAATGAGC CGGGGGGCTC ATGTTCTGTC *396
CTGCAGAACA TTTCTGTACT AGTGAGAAGA GGAATATGC ATTGCAGTTC *446
AGCAAAGCCG GAATTCTGTG TTGAACAGAT GTCTGTCTCC CTAGTGTGTG *496
ACTCACACCT TGTGGCTGCC TTCAGAGCGC CACCTCCAGA TCAGATGGGG *546

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

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