



TBXAS1 (NM_001061.7) - cDNA - 2026-06-20

ACTCCATGTG ATGTTTGCTT GGTTCCTGT TCCCTTTTCT ACCTGCAGAG -78
CACGGTTCCC ATAAGGGCGG CGAGATCAGC CTCCTGTCTC ATCTGGAAGA -28
CCACCACCTCT GGGGTCTCAG AGGAATGATG GAAGCCTTGG GGTTCCTAAA 23
ATTGGAAGTG AATGGCCCCA TGGTGACGGT GGCCCTGTCA GTGGCTCTCT 73
TGGCCCTCCT GAAATGGTAC TCCACATCAG CATTCTCAAG ACTGGAGAAG 123
TTAGGCCTCA GACATCCCAA GCCTTCTCCT TTCATTGGAA ACTTGACATT 173
TTTCCGCCAG GGTTTTTTGGG AAAGCCAAAT GGAGCTCAGA AAGCTGTATG 223 R60Pfs*33 R71Efs*25
GACCTCTGTG TGGGTACTAT CTTGGTCGTC GGATGTTTAT TGTATTCTCT 273 p.L77Cfs*19
GAGCCAGACA TGATCAAGCA GGTGTTGGTT GAGAACTTCA GTAACCTTAC 323
CAACAGAATG GCGTCGGGTT TGGAGTTCAA GTCGGTAGCC GACAGCGTTC 373
TGTTTTTACG TGACAAAAGA TGGGAAGAGG TCAGAGGTGC CCTGATGTCT 423
GCTTTCAGTC CTGAAAAGCT GAACGAGATG GTTCCCCCTCA TCAGCCAAGC 473
CTGCGACCTT CTCCTGGCTC ATTTAAAACG CTATGCGGAA TCTGGGGACG 523
CATTTGACAT CCAGAGGTGC TACTGCAATT ACACCACAGA TGTGGTTGCC 573
AGCGTCGCCT TTGGCACCCC GGTGGACTCC TGGCAGGCC CTGAGGATCC 623
CTTTGTGAAA CACTGCAAGC GTTCTTTCGA ATTCTGCATC CCCAGACCTA 673
TCCTGGTTTT ACTCTTATCA TTTCCATCCA TAATGGTCCC ACTGGCCCCG 723
ATTTTGCCCA ATAAGAACCG AGACGAACTG AATGGCTTTT TTAACAAACT 773
CATTAGGAAT GTGATTGCCT TGCGGGACCA GCAAGCTGCC GAAGAGAGGC 823
GGAGAGACTT CCTCCAAATG GTCCTGGATG CCCGACATC TGCAAGTCCC 873
ATGGGCGTGC AAGACTTTGA CATCGTCAGA GACGTTTTCT CCTCTACTGG 923
GTGCAAGCCG AACCTTCCC GGCAACACCA GCCCAGCCCT ATGGCCAGGC 973

CTTTGACTGT GGATGAGATT GTGGGCCAGG CCTTCATCTT CCTCATCGCT 1023
GGCTATGAAA TCATCACCAA CACACTTCTT TTTGCCACCT ACCTACTGGC 1073
CACCAACCCT GACTGCCAAG AGAAGCTTCT GAGAGAGGTA GACGTTTTTA 1123
AGGAGAAACA CATGGCCCCCT GAGTTCGTCA GCCTCGAGGA AGGCCTGCCC 1173
TATCTGGACA TGGTGATTGC AGAGACGCTG AGGATGTACC CGCCAGCTTT 1223
CAGATTCACA CGGGAGGCAG CTCAGGACTG CGAGGTGCTG GGGCAGCGCA 1273
TCCCCGCAGG CGCTGTGCTA GAGATGGCCG TGGGTGCCCT GCACCATGAC 1323
CCTGAGCACT GGCCAAGCCC GGAGACCTTC AACCCTGAAA GGTTCACGGC 1373 P448L
TGAGGCCCGG CAGCAGCACC GGCCCTTCAC GTACCTGCCC TTCGGGGCCG 1423 G473W
GCCCACGGAG CTGCCCTCGGG GTGCGTCTAG GGCTGCTTGA GGTCAAGTTG 1473
ACACTGCTCC ACGTGCTGCA CAAGTTCGGG TTCCAAGCCT GCCCTGAGAC 1523
CCAGGTACCG CTGCAGCTAG AATCCAAATC TGCCCTAGGT CCAAAAAATG 1573
GTGTCTATAT CAAGATCGTA TCCCGCTGAC ACAGAAGGCT GCCGGGTGGG *21
GGGAGGGCAC CCCCAAATC AAAGAAAACC CTAAGTGTGG ATGTTCAGAA *71
TTTTTGAAAA ATGTCACTGA AGTGATTGAA AGAGTGCCCTG GCATGCAAGG *121
ATAAGAGGTT CTTTACATAA CATTTCTAA ATGCTTAATA AACGTTTGTT *171
GCACTTGGTT TTGACATTGC CAA

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