



*IKBK*G (NM_003639.4) - cDNA - 2026-02-28

AGCCC GTTCC TGCTCCGCGC TTCTGGAGCA CTGGCCAAGG CGGGCCGATT -209
CAGGACCCAG GTTACTTGGG CGGCGAGCTG GACTGTTTCT ACTCCTCCCT -159
CCTCCTCCAC TGC GGGGTCT GACCCTACTC CTTGTGTGAG GACTCCTCTA -109
GTT CAGAGAC ATATTCTGTT CACCAAAC TT GACTGCGCTC TATCGAGGTC -59
GTTAAAT TCT TCGGAAATGC CTCACATATA GTTTGGCAGC TAGCCCTTGC -9
CCTGT TGGAT GAATAGGCAC CTC TGGGAAGA GCCAACTGTG TGAGATGGTG 42
CAGCCCAGTG GTGGCCCGGC AGCAGATCAG GACGTACTGG GCGAAGAGTC 92
TCCTCTGGGG AAGCCAGCCA TGCTGCACCT GCCTTCAGAA CAGGGCGCTC 142
CTGAGACCCT CCAGCGCTGC CTGGAGGAGA ATCAAGAGCT CCGAGATGCC 192
ATCCGGCAGA GCAACCAGAT TCTGCGGGAG CGCTGCGAGG AGCTTCTGCA 242
TTTCCAAGCC AGCCAGAGGG AGGAGAAGGA GTTCC TCATG TGCAAGTTCC 292
AGGAGGCCAG GAAACTGGTG GAGAGACTCG GCCTGGAGAA GCTCGATCTG 342
AAGAGGCAGA AGGAGCAGGC TCTGCGGGAG GTGGAGCACC TGAAGAGATG 392
CCAGCAGCAG ATGGCTGAGG ACAAGGCCTC TGTGAAAGCC CAGGTGACGT 442
CCTTGCTCGG GGAGCTGCAG GAGAGCCAGA GTCGCTTGGG GGCTGCCACT 492
AAGGAATGCC AGGCTCTGGA GGGTCGGGCC CGGGCGGCCA GCGAGCAGGC 542
GCGGCAGCTG GAGAGTGAGC GCGAGGCGCT GCAGCAGCAG CACAGCGTGC 592
AGGTGGACCA GCTGCGCATG CAGGGCCAGA GCGTGGAGGC CGCGCTCCGC 642 [V199V](#) [Q205*](#)
ATGGAGCGCC AGGCCGCC TC GGAGGAGAAG AGGAAGCTGG CCCAGTTGCA 692
GGTGGCCTAT CACCAGCTCT TCCAAGAATA CGACAACCAC ATCAAGAGCA 742
GCGTGGTGGG CAGTGAGCGG AAGCGAGGAA TGCAGCTGGA AGATCTCAA 792
CAGCAGCTCC AGCAGGCCGA GGAGGCCCTG GTGGCCAAAC AGGAGGTGAT 842

CGATAAGCTG AAGGAGGAGG CCGAGCAGCA CAAGATTGTG ATGGAGACCG 892
TTCCGGTGCCT GAAGGCCAG GCGGATATCT ACAAGGCCGA CTCCAGGCT 942
GAGAGGCAGG CCCGGGAGAA GCTGGCCGAG AAGAAGGAGC TCCTGCAGGA 992
GCAGCTGGAG CAGCTGCAGA GGGAGTACAG CAACTGAAG GCCAGCTGTC 1042
AGGAGTCGGC CAGGATCGAG GACATGAGGA AGCGGCATGT CGAGGTCTCC 1092
CAGGCCCCCT TGCCCCCGC CCCTGCCTAC CTCTCCTCTC CCCTGGCCCT 1142
GCCCAGCCAG AGGAGGAGCC CCCCCGAGGA GCCACCTGAC TTCTGCTGTC 1192
CCAAGTGCCA GTATCAGGCC CCTGATATGG ACACCCTGCA GATACATGTC 1242
ATGGAGTGCA TTGAGTAGGG CCGGCCAGTG CAAGGCCACT GCCTGCCGAG *32
GACGTGCCCC GGACCGTGCA GTCTGCGCTT TCCTCTCCCG CCTGCCTAGC *82
CCAGGATGAA GGGCTGGGTG GCCACAACCT GGATGCCACC TGGAGCCCCA *132
CCCAGGAGCT GGCCGCGGCA CCTTACGCTT CAGCTGTTGA TCCGCTGGTC *182
CCCTCTTTTG GGGTAGATGC GGCCCCGATC AGGCCTGACT CGCTGCTCTT *232
TTTGTTCCTT TCTGTCTGCT CGAACCACCT GCCTCGGGCT AATCCCTCCC *282
TCTTCCTCCA CCCGGCACTG GGAAGTCAA GAATGGGGCC TGGGGCTCTC *332
AGGGAGAACT GCTTCCCTG GCAGAGCTGG GTGGCAGCTC TTCCTCCAC *382
CGGACACCGA CCCGCCCGCT GCTGTGCCCT GGGAGTGCTG CCCTCTTACC *432
ATGCACACGG GTGCTCTCCT TTTGGGCTGC ATGCTATTCC ATTTTGCAGC *482
CAGACCGATG TGTATTTAAC CAGTCACTAT TGATGGACAT TTGGGTTGTT *532
TCCCATCTTT TTGTTACCAT AAATAATGCG ATAGTAAAAA TCCTTGTGCA *582
TTA

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