



HCK (NM_002110.5) - cDNA - 2026-03-07

ACCACGTCCC TGGTCCCAGC TCGGGAGCAC ATCAGAGGCT TAGAGGCGAG -134
TGGGAAGGGA CTCAGACAGT GCAGGACGAG AAACGCCCGC GGCACCAAAG -84
CCCCTCAGAG CGTCGCCCCC GCCTCTAGTT CTAGAAAGTC AGTTTCCCGG -34
CACTGGCACC CCGGAACCTC AGGGGCTGCC GAGCTGGGGG GCGCTCAAG 17
CTGCGAGGAT CCGGGCTGCC CGCGAGACGA GGAGCGGGCG CCCAGGATGG 67
GGTGCATGAA GTCCAAGTTC CTCCAGGTCG GAGGCAATAC ATTCTCAAAA 117
ACTGAAACCA GCGCCAGCCC ACACTGTCTT GTGTACGTGC CGGATCCCAC 167
ATCCACCATC AAGCCGGGGC CTAATAGCCA CAACAGCAAC ACACCAGGAA 217
TCAGGGAGGC AGGCTCTGAG GACATCATCG TGGTTGCCCT GTATGATTAC 267
GAGGCCATTC ACCACGAAGA CCTCAGCTTC CAGAAGGGGG ACCAGATGGT 317
GGTCCTAGAG GAATCCGGGG AGTGGTGGAA GGCTCGATCC CTGGCCACCC 367
GGAAGGAGGG CTACATCCA AGCAACTATG TCGCCCGCGT TGACTCTCTG 417
GAGACAGAGG AGTGGTTTTT CAAGGGCATC AGCCGGAAGG ACGCAGAGCG 467
CCAAC TGCTG GCTCCCGGCA ACATGCTGGG CTCCTTCATG ATCCGGGATA 517
GCGAGACCAC TAAAGGAAGC TACTCTTTGT CCGTGCAGAG CTACGACCCT 567
CGGCAGGGAG ATACCGTGAA ACATTACAAG ATCCGGACCC TGGACAACGG 617
GGGCTTCTAC ATATCCCCC GAAGCACCTT CAGCACTCTG CAGGAGCTGG 667
TGGACCACTA CAAGAAGGGG AACGACGGGC TCTGCCAGAA ACTGTCGGTG 717
CCCTGCATGT CTTCCAAGCC CCAGAAGCCT TGGGAGAAAG ATGCCTGGGA 767
GATCCCTCGG GAATCCCTCA AGCTGGAGAA GAAACTTGGG GCTGGGCAGT 817
TTGGGGAAGT CTGGATGGCC ACCTACAACA AGCACACCAA GGTGGCAGTG 867
AAGACGATGA AGCCAGGGAG CATGTCGGTG GAGGCCTTCC TGGCAGAGGC 917

CAACGTGATG AAAACTCTGC AGCATGACAA GCTGGTCAAA CTCATGCGG 967
TGGTCACCAA GGAGCCCATC TACATCATCA CGGAGTTCAT GGCCAAAGGA 1017
AGCTTGCTGG ACTTTCTGAA AAGTGATGAG GGCAGCAAGC AGCCATTGCC 1067
AAAACCTCATT GACTTCTCAG CCCAGATTGC AGAAGGCATG GCCTTCATCG 1117
AGCAGAGGAA CTACATCCAC CGAGACCTCC GAGCTGCCAA CATCTTGGTC 1167
TCTGCATCCC TGGTGTGTAA GATTGCTGAC TTTGGCCTGG CCCGGGTCAT 1217
TGAGGACAAC GAGTACACGG CTCGGGAAGG GGCCAAGTTC CCCATCAAGT 1267
GGACAGCTCC TGAAGCCATC AACTTTGGCT CCTTCACCAT CAAGTCAGAC 1317
GTCTGGTCCCT TTGGTATCCT GCTGATGGAG ATCGTCACCT ACGGCCGGAT 1367
CCCTTACCCA GGGATGTCAA ACCCTGAAGT GATCCGAGCT CTGGAGCGTG 1417
GATACCGGAT GCCTCGCCA GAGAACTGCC CAGAGGAGCT CTACAACATC 1467
ATGATGCGCT GCTGGAAAAA CCGTCCGGAG GAGCGGCCGA CCTTCGAATA 1517
CATCCAGAGT GTGCTGGATG ACTTCTA_CAC GGCCACAGAG AGCCAGTACC 1567 Y515*
AACAGCAGCC ATGATAGGGA GGACCAGGGC AGGGCCAGGG GGTGCCCAGG *36
TGGTGGCTGC AAGGTGGCTC CAGCACCATC CGCCAGGGCC CACACCCCCT *86
TCCTACTCCC AGACACCCAC CCTCGCTTCA GCCACAGTTT CCTCATCTGT *136
CCAGTGGGTA GGTGGACTG GAAAATCTCT TTTTGGACTCT TGCAATCCAC *186
AATCTGACAT TCTCAGGAAG CCCCCAAGTT GATATTTCTA TTTCCTGGAA *236
TGGTTGGATT TTAGTTACAG CTGTGATTTG GAAGGAAAC TTTCAAAATA *286
GTGAAATGAA TATTTAAATA AAAGATATAA ATGCCAAAGT CTTTACCAA *336

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