HCK (NM_002110.5) - cDNA + Protein - 2025-10-30

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ACCACGTCCC TGGTCCCAGC TCGGGAGCAC ATCAGAGGCT TAGAGGCGAG -134
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                                   MetGlyG lyArgSerSe 6

CTGCGAGGAT CCGGGCTGCC CGCGAGACGA GGAGCGGGCG CCCAGGATGG 67
rCysGluAsp ProGlyCysP roArgAspGl uGluArgAla ProArgMetG 23

GGTGCATGAA GTCCAAGTTC CTCCAGGTCG GAGGCAATAC ATTCTCAAAA 117
lyCysMetLy sSerLysPhe LeuGlnValG lyGlyAsnTh rPheSerLys 39

ACTGAAACCA GCGCCAGCCC ACACTGTCCT GTGTACGTGC CGGATCCCAC 167
ThrGluThrS erAlaSerPr oHisCysPro ValTyrValP roAspProTh 56

ATCCACCATC AAGCCGGGGC CTAATAGCCA CAACAGCAAC ACACCAGGAA 217
rSerThrIle LysProGlyP roAsnSerHi sAsnSerAsn ThrProGlyI 73

TCAGGGAGGC AGGCTCTGAG GACATCATCG TGGTTGCCCT GTATGATTAC 267
leArgGluAl aGlySerGlu AspIleIleV alValAlaLe uTyrAspTyr 89

GAGGCCATTC ACCACGAAGA CCTCAGCTTC CAGAAGGGGG ACCAGATGGT 317
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GluAlaIleH isHisGluAs pLeuSerPhe GlnLysGlyA spGlnMetVa 106

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rgLysGluGl yTyrIlePro SerAsnTyrV alAlaArgVa lAspSerLeu 139

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LysThrMetL ysProGlySe rMetSerVal GluAlaPheL euAlaGluAl 306

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AGCAGAGGAA CTACATCCAC CGAGACCTCC GAGCTGCCAA CATCTTGGTC 1167
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SerAlaSerL euValCysLy sIleAlaAsp PheGlyLeuA laArgValIl 406

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ValTrpSerP heGlyIleLe uLeuMetGlu IleValThrT yrGlyArgIl 456

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eProTyrPro GlyMetSerA snProGluVa lIleArgAla LeuGluArgG 473

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lyTyrArgMe tProArgPro GluAsnCysP roGluGluLe uTyrAsnIle 489

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MetMetArgC ysTrpLysAs nArgProGlu GluArgProT hrPheGluTy 506

CATCCAGAGT GTGCTGGATG ACTTCTACAC GGCCACAGAG AGCCAGTACC 1567 Y515*

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lnGlnGlnPr oStop

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GTGAAATGAA TATTTAAATA AAAGATATAA ATGCCAAAGT CTTTACCAAA *336

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HCK (NM_002110.5) - cDNA + Protein - 2025-10-30

nfevers