

Product datasheet for **SC202225**

AK2 (NM_001625) Human 3' UTR Clone

Product data:

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|---------------------------|------------------------------------|
| Product Type: | 3' UTR Clones |
| Product Name: | AK2 (NM_001625) Human 3' UTR Clone |
| Symbol: | AK2 |
| Synonyms: | ADK2 |
| Mammalian Cell Selection: | Neomycin |
| Vector: | pMirTarget (PS100062) |
| ACCN: | NM_001625 |
| Insert Size: | 2000 bp |



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Insert Sequence: >SC202225 3'UTR clone of NM_001625
 The sequence shown below is from the reference sequence of NM_001625. The complete sequence of this clone may contain minor differences, such as SNPs.
 Blue=Stop Codon Red=Cloning site

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GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAGCGATCGCC
ACATGTAAGACTTGTTATGTTTATCTAATGTTGGGTCCAAGAAGGAATTTCTTTCCATCCCTGTGAG
GCAATGGGTGGGAATGATAGGACAGGCAAGAGAAGCTTCTCAGGCTAGCAAAAATATCATTGATGT
ATTGATTAATAAGCACTTGCTTGATGATCTTTGGCGTGTGTGCTACTCTCATCTGTGTATGTGTG
TTGTGTGTGTGTGTGTGCATGCACATATGTGTTCACTCTGCTACTTTGTAAGTTTTAGGCTAGGTTG
CTTTACCAGCTGTTACTTCTTTTTTGTGTTGTTTTGAGACAAGGTTTCGCTCTGCCACCCTGGCTGG
AGTGCAGTGGCGTGATCTGGCTCACGGCAACCTCTGCCTCTGGGGCTCAAGCAATTATCCACCTCA
GCCTCCTGAGCAGCTGGGACTACAGGTGCATGCCACAACACCTGGCTGATATTTGATTTTTTTAGAG
ACAGGATTTTCCAAGTTGCCAGGCTGGTCTTGAAGCTTAGGCTTAAGCAATCCACCCACCTTGCC
TCCTGAAGTGCCAGGATCACAGAGCTGAGCCACTACACCCAGCCAGCTGTTACTTCTTTAACCATAC
TTTTGATTTTATTTTTGACCAAAATGAACTAACCCAGGTAATCTTCCAGGGACCGAATTCAGAACC
TCATAGTATTTCTCCATTTCCAGCAGCTGATTAGAAGTCCAGGATCATGTGAAGTCAGGCAGGGTAC
AGTTCTGATGGCACATTATGGACAGAGAATTCATTTTGTCTAACCATGATGAAAACCCACGTG
AGTCAGTGTGTGAACAGGGATCATTAAATTTTTCCCCCTAGGTGGAAGGAAAAAGGCACTTACTTTGCA
GGTTACAGAAATTAAGGAGAGATATCGTCATAAAAAGAGCCAGGCCAAATTTGGAATTTTTGTGA
TCTGCATCATGATGCTGAAAATAGCAATATTTGGGAATTGGGTTTGAAGTGAATTTGTTGCCAGAGA
ATTAACCAGGTGAAAGGTCCTTTTGAATTCAGATTGTCTTGAACATCCAGGCTGATCATCTGAGAG
CAGTCAAATCTACTTCCCAAAAAGAGACCAGGGTAGGTTTATTTGCTTTTATTTTTAATGTTTGCCTG
TGTTTCCAAGTGTGAACAAAACAGTGTGTGATCTATTCTTGATTCAATTTGATCAGTATTTATTCAA
CCCAGTCTCTCTCCAGGACATAAACTGAAATCAGATATGTTCTTTTTAAGCCAAAACCTCTCCTTTC
TAGATCCAACCCTTACCCTAATTTTATGATGGCTATAGCCATGGACTTCCCAAGAAAAGATCACCC
AGAAATAAGACCACCTGTGACAGTTACCAGCTTTTATTACATAACCTTAGCTTCCCAACTATTGAGCATT
TTCTAAGTCCCTGCTGTCTTTTGGTCTCTGGTTGATTTGTGGCAAACAGATGAAGTAACAGACTGCT
ATGAAGGACCACAAAACGGCAGCCTCTGAAAAACCATTAGAAAGTCAGTGGCAGATCCAGTAAATAA
TATCGCCAGCCTCAGCATAATCTGCTGCTGACTCGATTAGTGGACTCTAAAGTGCCAGCCTCCTGAC
CTGAGCTCTCCTGCCATCTGTGAGACTACCAGAGGCTTATCTGCTGTCCACATGGCAACTGGGCATGA
GTACCTGGCCACCTTGCTTCCCTCTTGGCTGGTCCAAGTGAGTGTCTGCTGCCTCTGCTCTGCCTGTG
TTTCTGGCTCTAAACCAACTCCACCCACTCTTAATGGAACTCAGTCTGGCTTTGTGTGTTTCTGGGA
AGCACATGACTTCTGGGAATGGGCAAGGAAGAGGAGTGAACAAAACCTGTGAGTATGTGTGCCTGGT
CTGGGATCCTTCTCTGGGTGACAGTGGCATCATGAATCTTAGAATCAGCTCCCCTTAACTAGGGACA
ACGCGTAAGCGGCCGCGGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCAACCTGCCATCA
CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
  
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Restriction Sites: SgfI-MluI

OTI Disclaimer: Our molecular clone sequence data has been matched to the sequence identifier above as a point of reference. Note that the complete sequence of this clone is largely the same as the reference sequence but may contain minor differences , e.g., single nucleotide polymorphisms (SNPs).

Components: The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The package also includes 100 pmols of both the corresponding 5' and 3' vector primers in separate vials.

RefSeq: [NM_001625.4](#)

Summary:

Adenylate kinases are involved in regulating the adenine nucleotide composition within a cell by catalyzing the reversible transfer of phosphate groups among adenine nucleotides. Three isozymes of adenylate kinase, namely 1, 2, and 3, have been identified in vertebrates; this gene encodes isozyme 2. Expression of these isozymes is tissue-specific and developmentally regulated. Isozyme 2 is localized in the mitochondrial intermembrane space and may play a role in apoptosis. Mutations in this gene are the cause of reticular dysgenesis. Alternate splicing results in multiple transcript variants. Pseudogenes of this gene are found on chromosomes 1 and 2.[provided by RefSeq, Nov 2010]

Locus ID:

204

MW:

75.4