

Product datasheet for **SC203466**

PCK2 (NM_001018073) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones
Product Name: PCK2 (NM_001018073) Human 3' UTR Clone
Vector: pMirTarget (PS100062)
Symbol: PCK2
Synonyms: PEPCK; PEPCK-M; PEPCK2
ACCN: NM_001018073
Insert Size: 478 bp
Insert Sequence: >SC203466 3'UTR clone of NM_001018073

The sequence shown below is from the reference sequence of NM_001018073. The complete sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

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GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC
CTTTCTCCACAACCTCCAACCATCTTCTAGGACTGCCAGGAGGCACAGAAGTCATGAACGTTTGCAGTT
TCCAGTCCCAGGCAAAATCTCAGTTCATGTCCCACTCCACCAGTCACTGGTTTTGTGATCTGGCTAAG
TTGCTCAACTTCCCTAAGCTTTAGTTTCCACATCAGTTGAATGAGGGTAGTTGTGATAGTACCTATCTC
ATGAGATTGTTGGAGGATTAATAGTGCATAAAAAGGGTTTATCACACTGACAAATACACAGTAAATTC
TCAATAATAAATACAGGCTGGATTTTTTTTTAATGAAAGGAAAAGGAAGGACTTTTGAACATTCTTACA
GAAGGTATTGGGCTCCAAGCACTATCCATAAAGTTTGGCCATTAGGAAAAGAGGAAAGCTGCCTCCTC
TGCTCCAACCTCCTCCTGCCACTTGGCTCCCCTGTCCCTGTATAATAACCACTGTCTAAAG
ACGCGTAAGCGGCCGCGGCATCTAGATTGGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA
CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
```

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_001018073.3](#)



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Summary: This gene encodes a mitochondrial enzyme that catalyzes the conversion of oxaloacetate to phosphoenolpyruvate in the presence of guanosine triphosphate (GTP). A cytosolic form of this protein is encoded by a different gene and is the key enzyme of gluconeogenesis in the liver. Alternatively spliced transcript variants have been described. [provided by RefSeq, Apr 2014]

Locus ID: 5106

MW: 17.7