

Product datasheet for SC202401

CAMK2D (NM 172114) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: CAMK2D (NM 172114) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: CAMK2D Synonyms: CAMKD

ACCN: NM_172114

Insert Size: 173 bp

Insert Sequence: >SC202401 3'UTR clone of NM_172114

The sequence shown below is from the reference sequence of NM_172114. The complete

sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

TCGGGGTCACCAACAGTACCCATCAAGTAAATATTTCCAGGCTGTCAGCTTCTTTGTTAATACACCCAT GGTCAGCTCCTTCTACTTATTCCATTGTTAATAGCATGGTATATGTTATTTAACGCTAGTAGTTGGTTA

CACTGATGAAAATAAATGCCTTCACGGGAAAGGTT

ACGCGTAAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul

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.

RefSeg: NM 172114.2



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CAMK2D (NM_172114) Human 3' UTR Clone - SC202401

Summary: The product of this gene belongs to the serine/threonine protein kinase family and to the

Ca(2+)/calmodulin-dependent protein kinase subfamily. Calcium signaling is crucial for several aspects of plasticity at glutamatergic synapses. In mammalian cells, the enzyme is composed of four different chains: alpha, beta, gamma, and delta. The product of this gene is a delta chain. Alternative splicing results in multiple transcript variants encoding distinct isoforms. Distinct isoforms of this chain have different expression patterns.[provided by

RefSeq, Nov 2008]

Locus ID: 817

MW: 6.5