

## **Product datasheet for SC204448**

## CDK4 (NM 000075) Human 3' UTR Clone

**Product data:** 

**Product Type:** 3' UTR Clones

Product Name: CDK4 (NM 000075) Human 3' UTR Clone

Symbol: CDK4

**Synonyms:** CMM3; PSK-J3

Mammalian Cell

Selection:

Neomycin

**Vector:** pMirTarget (PS100062)

**ACCN:** NM\_000075

**Insert Size:** 828 bp

Insert Sequence: >SC204448 3'UTR clone of NM\_000075

The sequence shown below is from the reference sequence of NM\_000075. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

 ${\sf TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC}$ 

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

**Restriction Sites:** Sgfl-Mlul



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## CDK4 (NM\_000075) Human 3' UTR Clone - SC204448

**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:** <u>NM 000075.4</u>

**Summary:** The protein encoded by this gene is a member of the Ser/Thr protein kinase family. This

protein is highly similar to the gene products of S. cerevisiae cdc28 and S. pombe cdc2. It is a catalytic subunit of the protein kinase complex that is important for cell cycle G1 phase progression. The activity of this kinase is restricted to the G1-S phase, which is controlled by the regulatory subunits D-type cyclins and CDK inhibitor p16(INK4a). This kinase was shown to be responsible for the phosphorylation of retinoblastoma gene product (Rb). Mutations in this gene as well as in its related proteins including D-type cyclins, p16(INK4a) and Rb were all found to be associated with tumorigenesis of a variety of cancers. Multiple polyadenylation

sites of this gene have been reported. [provided by RefSeq, Jul 2008]

Locus ID: 1019 MW: 31.6