

Product datasheet for SC205646

OriGene Technologies, Inc.

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PKC zeta (PRKCZ) (NM_002744) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: PKC zeta (PRKCZ) (NM_002744) Human 3' UTR Clone

Symbol: PKC zeta

Synonyms: PKC-ZETA; PKC2

Mammalian Cell

Selection:

Neomycin

Vector: pMirTarget (PS100062)

ACCN: NM_002744

Insert Size: 416 bp

Insert Sequence: >SC205646 3'UTR clone of NM_002744

The sequence shown below is from the reference sequence of NM_002744. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

AA

CAACCTGCCATCACGAGATTTCGATTCCACCGCCGC

Restriction Sites: Sgfl-Rsrll

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 002744.6</u>

Summary: Protein kinase C (PKC) zeta is a member of the PKC family of serine/threonine kinases which

are involved in a variety of cellular processes such as proliferation, differentiation and secretion. Unlike the classical PKC isoenzymes which are calcium-dependent, PKC zeta exhibits a kinase activity which is independent of calcium and diacylglycerol but not of phosphatidylserine. Furthermore, it is insensitive to typical PKC inhibitors and cannot be activated by phorbol ester. Unlike the classical PKC isoenzymes, it has only a single zinc finger module. These structural and biochemical properties indicate that the zeta subspecies is related to, but distinct from other isoenzymes of PKC. Alternative splicing results in multiple

transcript variants encoding different isoforms. [provided by RefSeq, Jul 2008]

Locus ID: 5590 **MW:** 15.4