

## **Product datasheet for SC202624**

## OriGene Technologies, Inc.

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## Protein Kinase D2 (PRKD2) (NM\_001079881) Human 3' UTR Clone

**Product data:** 

**Product Type:** 3' UTR Clones

Product Name: Protein Kinase D2 (PRKD2) (NM 001079881) Human 3' UTR Clone

Symbol: Protein Kinase D2

**Synonyms:** HSPC187; nPKC-D2; PKD2

**Mammalian Cell** 

Selection:

Neomycin

**Vector:** pMirTarget (PS100062)

**ACCN:** NM\_001079881

**Insert Size:** 237 bp

Insert Sequence: >SC202624 3'UTR clone of NM\_001079881

The sequence shown below is from the reference sequence of NM\_001079881. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

ATCTCTGGCCCCATGGCCTTGATCTCAGCA

**ACGCGT**AAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA

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.

**RefSeq:** <u>NM 001079881.2</u>





## Protein Kinase D2 (PRKD2) (NM\_001079881) Human 3' UTR Clone - SC202624

Summary: The protein encoded by this gene belongs to the protein kinase D (PKD) family of

serine/threonine protein kinases. This kinase can be activated by phorbol esters as well as by gastrin via the cholecystokinin B receptor (CCKBR) in gastric cancer cells. It can bind to diacylglycerol (DAG) in the trans-Golgi network (TGN) and may regulate basolateral membrane protein exit from TGN. Alternative splicing results in multiple transcript variants

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

**Locus ID:** 25865 **MW:** 7.9