

## **Product datasheet for SC209217**

## p57 Kip2 (CDKN1C) (NM 000076) Human 3' UTR Clone

**Product data:** 

**Product Type:** 3' UTR Clones

Product Name: p57 Kip2 (CDKN1C) (NM\_000076) Human 3' UTR Clone

**Vector:** pMirTarget (PS100062)

Symbol: CDKN1C

Synonyms: BWCR; BWS; KIP2; p57; p57Kip2; WBS

**ACCN:** NM\_000076

**Insert Size:** 746 bp

Insert Sequence: >SC209217 3'UTR clone of NM\_000076

The sequence shown below is from the reference sequence of NM\_000076. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

**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).



**OriGene Technologies, Inc.** 9620 Medical Center Drive, Ste 200

CN: techsupport@origene.cn

Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com



## p57 Kip2 (CDKN1C) (NM\_000076) Human 3' UTR Clone - SC209217

**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 000076.2</u>

Summary: This gene is imprinted, with preferential expression of the maternal allele. The encoded

protein is a tight-binding, strong inhibitor of several G1 cyclin/Cdk complexes and a negative regulator of cell proliferation. Mutations in this gene are implicated in sporadic cancers and Beckwith-Wiedemann syndorome, suggesting that this gene is a tumor suppressor candidate. Three transcript variants encoding two different isoforms have been found for this gene.

Three transcript variants encouning two uniferent isolornis have been found for the

[provided by RefSeq, Oct 2010]

Locus ID: 1028 MW: 28.1