

Product datasheet for MC207618

Cinp (NM_026048) Mouse Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: Cinp (NM_026048) Mouse Untagged Clone

Tag: Tag Free
Symbol: Cinp

Synonyms: 1810047K05Rik; 2810452K22Rik

Mammalian Cell

Selection:

Neomycin

Vector:pCMV6-Entry (PS100001)E. coli Selection:Kanamycin (25 ug/mL)

Fully Sequenced ORF: >MC207618 representing NM_026048

Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC

GCCGCGATCGCC

GCCCTGTGA

 ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT

ACAAGGATGACGACGATAAGGTTTAA

Restriction Sites: Sgfl-Mlul ACCN: NM 026048

Insert Size: 639 bp



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OTI Disclaimer: Our molecular clone sequence data has been matched to the reference identifier above as a

point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative

RNA splicing form or single nucleotide polymorphism (SNP).

Components: The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube

containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

Reconstitution Method: 1. Centrifuge at 5,000xg for 5min.

2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.

3. Close the tube and incubate for 10 minutes at room temperature.

4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid

at the bottom.

5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of

shipping when stored at -20°C.

RefSeq: <u>NM 026048.4</u>, <u>NP 080324.1</u>

12 F1

 RefSeq Size:
 2029 bp

 RefSeq ORF:
 639 bp

 Locus ID:
 67236

 UniProt ID:
 Q9D0V8

Cytogenetics:

Gene Summary: Interacts with the components of the replication complex and 2 kinases, CDK2 and CDC7,

thereby providing a functional and physical link between CDK2 and CDC7 during firing of the

origins of replication. Regulates ATR-mediated checkpoint signaling (By similarity).

[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (1) represents the predominant transcript. It encodes isoform 1. Sequence Note: This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on transcript alignments.