

Product datasheet for **SC204896**

CNKSR1 (NM_006314) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	CNKSR1 (NM_006314) Human 3' UTR Clone
Symbol:	CNKSR1
Synonyms:	CNK; CNK1
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_006314
Insert Size:	366 bp
Insert Sequence:	>SC204896 3'UTR clone of NM_006314 The sequence shown below is from the reference sequence of NM_006314. The complete sequence of this clone may contain minor differences, such as SNPs. Blue =Stop Codon Red =Cloning site

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GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC
GAGAGCAGCCTCCGACCTCCTGACCTCTGACCCTGGCCAGCACTCTAGCTCCTGACCTTTGACCCGAGG
GCCACCTCAACCCAGCTTCTGACGTGTCCAGGACAGAGCATCCCTGGATTCTGTTTCAGGGTGGGAAGT
AGTACTGCTAGTCATGGTCTCACCCGAGCTGACCCCTCTGCCTGGGCTTTGTGCCACCTCTCCCTTG
CCAAAGAAGAACTCTCCCCCAATCCTCCAACCTCTGGGGCCACAGCCCTGCCCTCCAGTTCCTTG
GCAGTTCTCCCCAAACCAGGTCTGTACAGGTGTTCTTTATTTTACATGAGGGCTACTTTCCAACCAAA
TAAAGTCAATTTTTCTAAGAA
ACGCGTAAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA
CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
```

Restriction Sites:	Sgfl-MluI
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.



[View online »](#)

RefSeq: [NM_006314.3](#)

Summary: This gene encodes a protein containing several motifs involved in protein-protein interaction, including PDZ, PH (Pleckstrin homology), and SAM (sterile alpha motif) domains. The encoded protein acts as a scaffold component for receptor tyrosine kinase signaling and may mediate crosstalk between different signaling pathways. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Jul 2014]

Locus ID: 10256

MW: 13.4