

Product datasheet for SC209062

CSK (NM 001127190) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: CSK (NM_001127190) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: CSK

ACCN: NM 001127190

Insert Size: 714 bp

Insert Sequence: >SC209062 3'UTR clone of NM_001127190

The sequence shown below is from the reference sequence of NM_001127190. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

 ${\tt CCGCCTCGGCGCTTCCTCCACCGA}$

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.



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CSK (NM_001127190) Human 3' UTR Clone - SC209062

RefSeq: <u>NM 001127190.2</u>

Summary: The protein encoded by this gene is involved in multiple pathways, including the regulation of

Src family kinases. It plays an important role in T-cell activation through its association with the protein encoded by the protein tyrosine phosphatase, non-receptor type 22 (PTPN22) gene. This protein also phosphorylates C-terminal tyrosine residues on multiple substrates, including the protein encoded by the SRC proto-oncogene, non-receptor tyrosine kinase gene. Phosphorylation suppresses the kinase activity of the Src family tyrosine kinases. An intronic polymorphism (rs34933034) in this gene has been found to affect B-cell activation and is associated with systemic lupus erythematosus (SLE). Alternative splicing results in multiple

transcript variants. [provided by RefSeq, Aug 2017]

Locus ID: 1445 MW: 26.4