

Product datasheet for SC210126

ACK1 (TNK2) (NM 001010938) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: ACK1 (TNK2) (NM_001010938) Human 3' UTR Clone

Symbol: ACK1

Synonyms: ACK; ACK-1; ACK1; p21cdc42Hs

Mammalian Cell

Selection:

Neomycin

Vector: pMirTarget (PS100062)

ACCN: NM_001010938

Insert Size: 847 bp

Insert Sequence: >SC210126 3'UTR clone of NM_001010938

The sequence shown below is from the reference sequence of NM_001010938. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

 ${\sf TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC}$

TAGTTTCTATTTTATGTTA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul



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



ACK1 (TNK2) (NM_001010938) Human 3' UTR Clone - SC210126

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 001010938.2</u>

Summary: This gene encodes a tyrosine kinase that binds Cdc42Hs in its GTP-bound form and inhibits

both the intrinsic and GTPase-activating protein (GAP)-stimulated GTPase activity of Cdc42Hs.

This binding is mediated by a unique sequence of 47 amino acids C-terminal to an SH3 domain. The protein may be involved in a regulatory mechanism that sustains the GTP-bound active form of Cdc42Hs and which is directly linked to a tyrosine phosphorylation signal transduction pathway. Several alternatively spliced transcript variants have been identified from this gene, but the full-length nature of only two transcript variants has been

determined. [provided by RefSeq, Jul 2008]

Locus ID: 10188

MW: 29.5