

Product datasheet for SC205892

RNF22 (TRIM3) (NM 033278) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: RNF22 (TRIM3) (NM_033278) Human 3' UTR Clone

Symbol: RNF22

Synonyms: BERP; HAC1; RNF22; RNF97

Mammalian Cell

Selection:

Neomycin

Vector: pMirTarget (PS100062)

ACCN: NM_033278

Insert Size: 445 bp

The sequence shown below is from the reference sequence of NM_033278. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

CCCCCATATTAAATAAATGTCTTCACCAAGA

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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RefSeq: <u>NM 033278.4</u>

Summary: The protein encoded by this gene is a member of the tripartite motif (TRIM) family, also called

the 'RING-B-box-coiled-coil' (RBCC) subgroup of RING finger proteins. The TRIM motif includes three zinc-binding domains, a RING, a B-box type 1 and a B-box type 2, and a coiled-coil region. This protein localizes to cytoplasmic filaments. It is similar to a rat protein which is a specific partner for the tail domain of myosin V, a class of myosins which are involved in the targeted transport of organelles. The rat protein can also interact with alpha-actinin-4. Thus it is suggested that this human protein may play a role in myosin V-mediated cargo transport.

Alternatively spliced transcript variants encoding the same isoform have been identified.

[provided by RefSeq, Jul 2008]

Locus ID: 10612 MW: 16.1