

Product datasheet for SC205179

RbAp46 (RBBP7) (NM 002893) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: RbAp46 (RBBP7) (NM_002893) Human 3' UTR Clone

Symbol: RbAp46
Synonyms: RbAp46

Mammalian Cell

Selection:

Neomycin

Vector: pMirTarget (PS100062)

ACCN: NM_002893

Insert Size: 724 bp

Insert Sequence: >SC205179 3'UTR clone of NM_002893

The sequence shown below is from the reference sequence of NM_002893. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

GGCTGTAAAAATAAAGCAGTGTGTTTAGCACAAA

ACGCGTAAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA

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).



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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 002893.4</u>

Summary: This protein is a ubiquitously expressed nuclear protein and belongs to a highly conserved

subfamily of WD-repeat proteins. It is found among several proteins that binds directly to retinoblastoma protein, which regulates cell proliferation. The encoded protein is found in many histone deacetylase complexes, including mSin3 co-repressor complex. It is also present in protein complexes involved in chromatin assembly. This protein can interact with BRCA1 tumor-suppressor gene and may have a role in the regulation of cell proliferation and differentiation. Two transcript variants encoding different isoforms have been found for this

gene. [provided by RefSeq, Nov 2010]

Locus ID: 5931 **MW:** 27.4