

## **Product datasheet for SC201052**

## RANBP1 (NM 002882) Human 3' UTR Clone

**Product data:** 

**Product Type:** 3' UTR Clones

Product Name: RANBP1 (NM\_002882) Human 3' UTR Clone

**Vector:** pMirTarget (PS100062)

Symbol: RANBP1
Synonyms: HTF9A

**ACCN:** NM\_002882

**Insert Size:** 333 bp

Insert Sequence: >SC201052 3'UTR clone of NM\_002882

The sequence shown below is from the reference sequence of NM\_002882. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

AAACACTTATTTATAGTCATCAAAAATAGTGAATAAAAAACACATTTGGAACCTGGG

**ACGCGT**AAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA

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.

**RefSeg:** NM 002882.4



**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



## RANBP1 (NM\_002882) Human 3' UTR Clone - SC201052

Summary: This gene encodes a protein that forms a complex with Ras-related nuclear protein (Ran) and

metabolizes guanoside triphosphate (GTP). This complex participates in the regulation of the cell cycle by controlling transport of proteins and nucleic acids into the nucleus. There are multiple pseudogenes for this gene on chromosomes 9, 12, 17, and X. Alternative splicing

results in multiple transcript variants. [provided by RefSeq, Jul 2013]

**Locus ID:** 5902

MW: 13.2