

## **Product datasheet for SC205026**

## ZFAND2B (NM 138802) Human 3' UTR Clone

**Product data:** 

**Product Type:** 3' UTR Clones

Product Name: ZFAND2B (NM 138802) Human 3' UTR Clone

**Vector:** pMirTarget (PS100062)

Symbol: ZFAND2B
Synonyms: AIRAPL

ACCN: NM\_138802

**Insert Size:** 372 bp

Insert Sequence: >SC205026 3'UTR clone of NM\_138802

The sequence shown below is from the reference sequence of NM\_138802. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

TCGAAGCCGTCCAACTGCAGCCTGTGCTAGGGCCCTGGGCTTGGGAGGAGGAGGTTCACCTGAGGAGGAC TGTGGCCCTCACACCTCTAGGGTACACAGGGAGAGGAGGAGGCCCGGAGCACCCTGGAGGGCAGAGACAAGC GGGAGTGATGTGGAGGTCGCCCTGGGAGGCCTCTGGAAGGCCTTGCTAGTGCTCCAGCTGCATGGAAGAG AGCGGCTAGCAACTGTTCCCTGGTTGGGCCCTCAGTGGATGCTGGCCAGGCCCTACTCTTAGCCCCTTC ATCATGTCATCTCCCTTATGCTGGAGCTGCCCCGATGTGGAGTGGGCAGGAAGGGGCCTGGAAAAAATA

AAGGATCTTGGCAGTTGATAAAACGTA

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 138802.3



**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



## ZFAND2B (NM\_138802) Human 3' UTR Clone - SC205026

Summary: This gene encodes a protein containing AN1-type zinc-fingers and ubiquitin-interacting

motifs. The encoded protein likely associates with the proteosome to stimulate the

degradation of toxic or misfolded proteins. Alternatively spliced transcript variants encoding

multiple isoforms have been observed for this gene. [provided by RefSeq, Aug 2012]

**Locus ID:** 130617

MW: 13.1