

Product datasheet for SC210057

RBED1 (ELMOD3) (NM_001135022) Human 3' UTR Clone

Product data:

OriGene Technologies, Inc.

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Product Type:	3' UTR Clones
Product Name:	RBED1 (ELMOD3) (NM_001135022) Human 3' UTR Clone
Symbol:	RBED1
Synonyms:	DFNB88; LST3; RBED1; RBM29
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_001135022
Insert Size:	816 bp
Insert Sequence:	<pre>>SC210057 3'UTR clone of NM_001135022 The sequence shown below is from the reference sequence of NM_001135022. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC CACTCATCCGAAGGCGTATGGCTGATCTGACCTCCGAGATGAATGGAGGCTTAAAGGCTGAGCTGCAGG GGCTTTCAGGGGGTCAGTGGAGCCATGTCAGCACCCCCCCGCCACCCCTTGCTGTCTCAGCAGA TGGGATATAGGAAGCTCCTGGGCTTAGCTGGGAAGCCAAGTACCCTCACCGGCATGGGACATGAGGG GCAGCTAGACTTCACCCCCTTCCCGCAGACCTGCCCCCAGGCCAGGCAGG</pre>
Restriction Sites:	Sgfl-Mlul



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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 001135022.2</u>
Summary:	This gene encodes a member of the engulfment and cell motility family of GTPase-activating proteins that regulate Arf GTPase proteins. Members of this family are defined by a conserved engulfment and cell motility domain. In rat cochlea, the encoded protein is found in stereocilia, kinocilia and cuticular plate of developing hair cells suggesting a function for this protein in cochlear sensory cells. An allelic variant of this family has been associated with autosomal recessive nonsyndromic deafness-88 in humans. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Jul 2016]
Locus ID:	84173
MW:	28.8

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