

Product datasheet for **SC333558**

RABL5 (IFT22) (NM_001287525) Human Untagged Clone

Product data:

Product Type: Expression Plasmids
Product Name: RABL5 (IFT22) (NM_001287525) Human Untagged Clone
Tag: Tag Free
Symbol: RABL5
Synonyms: FAP9; RABL5
Vector: pCMV6-Entry (PS100001)
Fully Sequenced ORF: >SC333558 representing NM_001287525.
Blue=Insert sequence **Red**=Cloning site **Green**=Tag(s)

ATGAAGGATGCTCATGGAGTGGTGATCGTCTTCAATGCTGACATCCCAAGCCACCGGAAGGAAATGGAG
ATGTGGTATTCTGCTTTGTCCAACAGCCGCTTACAGGACACACAGTGTATGCTAATTGCACACCAC
AAACCAGGCTCTGGAGATGATAAAGGAAGCCTGTCTTTGTCGCCACCCTTGAACAAGCTGAAGCTGGTG
CACTCAAACCTGGAAGATGACCCTGAGGAGATCCGGATGGAATTCATAAAGTATTTAAAAAGCATAATC
AACTCCATGTCTGAGAGCAGAGACAGGGAGGAGATGTCAATTATGACCTAG

Restriction Sites: SgfI-MluI
ACCN: NM_001287525
Insert Size: 327 bp
OTI Disclaimer: Our molecular clone sequence data has been matched to the reference identifier above as a point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative RNA splicing form or single nucleotide polymorphism (SNP).
Components: The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).
Reconstitution Method:

1. Centrifuge at 5,000xg for 5min.
2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.
3. Close the tube and incubate for 10 minutes at room temperature.
4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.
5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.

RefSeq: [NM_001287525.1](#)



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RefSeq Size:	2512 bp
RefSeq ORF:	327 bp
Locus ID:	64792
UniProt ID:	Q9H7X7
Cytogenetics:	7q22.1
Protein Families:	Druggable Genome
MW:	12.4 kDa
Gene Summary:	<p>Small GTPase-like component of the intraflagellar transport (IFT) complex B.[UniProtKB/Swiss-Prot Function]</p> <p>Transcript Variant: This variant (5) differs in the 5' UTR, lacks a portion of the 5' coding region and initiates translation at a downstream start codon, compared to variant 1. Variants 3 through 6 encode the same isoform (c), which is shorter at the N-terminus, compared to isoform a. Sequence Note: This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on transcript alignments.</p>