

Product datasheet for SC209326

RAB38 (NM 022337) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: RAB38 (NM_022337) Human 3' UTR Clone

Symbol: RAB38

Synonyms: NY-MEL-1; rrGTPbp

Mammalian Cell

Selection:

Neomycin

Vector: pMirTarget (PS100062)

ACCN: NM_022337

Insert Size: 767 bp

Insert Sequence: >SC209326 3'UTR clone of NM_022337

The sequence shown below is from the reference sequence of NM_022337. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

 ${\sf TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC}$

TTTCCCTA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul



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RAB38 (NM_022337) Human 3' UTR Clone - SC209326

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 022337.3</u>

Summary: May be involved in melanosomal transport and docking. Involved in the proper sorting of

TYRP1. Involved in peripheral melanosomal distribution of TYRP1 in melanocytes; the

function, which probably is implicating vesicle-trafficking, includes cooperation with ANKRD27

and VAMP7 (By similarity). Plays a role in the maturation of phagosomes that engulf

pathogens, such as S.aureus and M.tuberculosis (PubMed:21255211). Plays an important role in the control of melanin production and melanosome biogenesis (PubMed:23084991). In concert with RAB32, regulates the proper trafficking of melanogenic enzymes TYR, TYRP1 and DCT/TYRP2 to melanosomes in melanocytes (By similarity).[UniProtKB/Swiss-Prot Function]

Locus ID: 23682

MW: 29.1