

Product datasheet for SC207914

RHOG (NM 001665) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: RHOG (NM_001665) Human 3' UTR Clone

Symbol: RHOG Synonyms: ARHG

Mammalian Cell Neomycin

Selection:

Neomychi

Vector: pMirTarget (PS100062)

ACCN: NM_001665

Insert Size: 615 bp

Insert Sequence: >SC207914 3'UTR clone of NM_001665

The sequence shown below is from the reference sequence of NM_001665. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

ACGCGTAAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA

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).



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RHOG (NM_001665) Human 3' UTR Clone - SC207914

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 001665.4</u>

Summary: This gene encodes a member of the Rho family of small GTPases, which cycle between

inactive GDP-bound and active GTP-bound states and function as molecular switches in signal transduction cascades. Rho proteins promote reorganization of the actin cytoskeleton and regulate cell shape, attachment, and motility. The encoded protein facilitates translocation of a functional guanine nucleotide exchange factor (GEF) complex from the cytoplasm to the plasma membrane where ras-related C3 botulinum toxin substrate 1 is activated to promote lamellipodium formation and cell migration. Two related pseudogene have been identified on

chromosomes 20 and X. [provided by RefSeq, Aug 2011]

Locus ID: 391

MW: 22.1