

## Product datasheet for **SC205827**

### **RHOD (NM\_014578) Human 3' UTR Clone**

#### Product data:

**Product Type:** 3' UTR Clones  
**Product Name:** RHOD (NM\_014578) Human 3' UTR Clone  
**Vector:** pMirTarget (PS100062)  
**Symbol:** RHOD  
**Synonyms:** ARHD; Rho; RHOHP1; RHOM  
**ACCN:** NM\_014578  
**Insert Size:** 445 bp  
**Insert Sequence:** >SC205827 3'UTR clone of NM\_014578  
The sequence shown below is from the reference sequence of NM\_014578. The complete sequence of this clone may contain minor differences, such as SNPs.  
**Blue**=Stop Codon **Red**=Cloning site

```
GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC
ATTACCCAGGGCTTTTGGTGGTGACCTGAGCGGCTCGGGCGTCCCAGCGACGCGGGAAGGGGCGAGGG
CGCTGACCTGCTGCTGAGCTGGCTGGCTGGACCCGGTCCCTAGGCTGTGACCGCGAATCCACTGCA
ACAGACGGGCGCCACCAAGCCAGGCCCTGAGGCCTGGGAGTCCCTGGACTGAGAAAGGGGTTCTGGG
CCCACCTGCTCTGTAGGGCTCGTCTGCGGTGCCCGAGAATCACTCGCTAACCCCTATGCCCGGTCC
CGGACCGACATCCTGGAGCCGCCTGTGCAGCCTGATGCCCCCTCGTGGCTGCTCCAGGGCTGCACCTG
CCAGGACCTAATGTTCTTAGGTCCCTCTGGCCAGAACCACACCCGGCCCTTCCACCTGTCATACTG
GTAAGTGAACAAGAAAAACGACATCACTTA
ACGCGTAAGCGGCCGCGCATCTAGATTGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA
CGAGATTCGATTCCACCGCCCTTCTATGAAAGG
```

**Restriction Sites:** SgfI-MluI

**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:** [NM\\_014578.4](#)



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**Summary:** Ras homolog, or Rho, proteins interact with protein kinases and may serve as targets for activated GTPase. They play a critical role in muscle differentiation. The protein encoded by this gene binds GTP and is a member of the small GTPase superfamily. It is involved in endosome dynamics and reorganization of the actin cytoskeleton, and it may coordinate membrane transport with the function of the cytoskeleton. Two transcript variants encoding different isoforms have been found for this gene. [provided by RefSeq, Jul 2014]

**Locus ID:** 29984

**MW:** 15.4