

## Product datasheet for SC206990

### RGL2 (NM\_004761) Human 3' UTR Clone

#### Product data:

**Product Type:** 3' UTR Clones  
**Product Name:** RGL2 (NM\_004761) Human 3' UTR Clone  
**Symbol:** RGL2  
**Synonyms:** HKE1.5; KE1.5; RAB2L  
**Mammalian Cell Selection:** Neomycin  
**Vector:** pMirTarget (PS100062)  
**ACCN:** NM\_004761  
**Insert Size:** 478 bp

**Insert Sequence:** >SC206990 3'UTR clone of NM\_004761  
 The sequence shown below is from the reference sequence of NM\_004761. The complete sequence of this clone may contain minor differences, such as SNPs.  
 Blue=Stop Codon Red=Cloning site

```
GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC
GGGAGGAAGATTGCACGGGCACTGTTCTGAGGAGGAAGCCCCGTTGGCTTACAGAAGTCATGGTGTTC
TACCAGATGTGGGTAGCCATCCTGAATGGTGGCAATTATACATTGAGACAGAAATTCAGAAAGGGA
GCCAGCCACCCTGGGCGAGTGAAGTGCCACTGGTTTACCAGACAGCTGAGAAATCCAGCCCTGTGGGAA
CTGGTGTCTTATAACCAAGTTGGATACCTGTGTATAGCTTCCCACCTCCATGAGTGCAGCACACAGGT
AGTGCTGGAAAAACGCATCAGTTTCTGATTCTTGGCCATATCCTAACATGCAAGGCCAAGCAAAGGCT
TCAAGGCTCTGAGCCCCAGGGCAGAGGGGAATGGCAAAATGTAGGTCTCGCAGGAGCTCTTCTCCCA
CTCTGGGGTTTCTATCACTGTGACAACACTAAGATAATAAACAAAAACTACCTGAATTCTA
ACGCGTAAGCGGCCGCGCATCTAGATTGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA
CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
```

**Restriction Sites:** Sgfl-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\\_004761.5](#)

Summary: Probable guanine nucleotide exchange factor. Putative effector of Ras and/or Rap. Associates with the GTP-bound form of Rap 1A and H-Ras in vitro (By similarity).[UniProtKB/Swiss-Prot Function]

Locus ID: 5863

MW: 17.9