

## **Product datasheet for SC115778**

## RIT1 (NM\_006912) Human Untagged Clone

## **Product data:**

**Product Type:** Expression Plasmids

Product Name: RIT1 (NM\_006912) Human Untagged Clone

Tag: Tag Free

Symbol: RIT1

Synonyms: NS8; RIBB; RIT; ROC1

Mammalian Cell None

Selection:

Vector: pCMV6-XL5

E. coli Selection: Ampicillin (100 ug/mL)

Fully Sequenced ORF: >OriGene ORF sequence for NM\_006912 edited

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5' Read Nucleotide Sequence: >OriGene 5' read for NM\_006912 unedited

TTTGTAATACGACTCACTATAGGGCGGCCGCGAATTCGGCACGAGGGGGAAGACGAAGTG CGTGACCCGACCGGCTGTGGTGTTCCAGTCCCCACTGACCAGTAGGAGCAGCAGGGCGTC GGCTTGTGAGGTGGCTTTTCCTCGGGGCAACCCAGGAAGGCCCCAAGAGGACAATGGATT CTGGAACTCGCCCAGTTGGTAGCTGCTGTAGCAGCCCCGCTGGGCTCTCACGGGAGTACA AACTAGTGATGCTGGGTGCTGGTGGTGTAGGGAAGAGTGCCATGACCATGCAGTTCATCA GCCACCGATTCCCAGAAGATCATGATCCCACCATTGAAGATGCTTATAAGATCAGGATCC GTATTGATGATGAGCCTGCCAATCTGGACATTTTGGATACAGCTGGACAGGCAGAGTTTA CAGCCATGCGGGACCAGTATATGAGGGCAGGAGAAGGGTTTATCATCTGTTACTCTATCA GTACTGACGATACACCTGTGGTTCTTGTGGGAAACAAGTCAGACCTCAAACAGCTAAGAC AGGTCACCAAGGAAGAAGGATTGGCCTTGGCCCGAGAATTCAGCTGTCCCTTTTTTGAGA CATCTGCTGCATACCGCTACTATATTGATGATGTTTTCCATGCCCTTGTACGGGAGATAC GTANGAAAGAAAAGGAGCAGTACTGGCCATGGAGAANAAATCTAAGCCAAAACAGTGTAT GGAAGAAGCTAANATCACCATTCCGGAAGAAGAAGATTCAGTAACTTGAAGAGAGATGT GAAGTGNTTATCTGTGAACTGCAGTGCTGTATCAAAGCAGTCCAGTAACCTGCAGTACTG AGTATGGTGCT

Restriction Sites: Notl-Notl
ACCN: NM\_006912
Insert Size: 2640 bp

OTI Disclaimer: Due to the inherent nature of this plasmid, standard methods to replicate additional amounts

of DNA in E. coli are highly likely to result in mutations and/or rearrangements. Therefore, OriGene does not guarantee the capability to replicate this plasmid DNA. Additional amounts of DNA can be purchased from OriGene with batch-specific, full-sequence verification at a reduced cost. Please contact our customer care team at <a href="mailto:customercom">customercom</a> or by

calling 301.340.3188 option 3 for pricing and delivery.

The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing

variants is recommended prior to use. <u>More info</u>

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: <u>NM 006912.3, NP 008843.1</u>

RefSeq Size: 1104 bp



## RIT1 (NM\_006912) Human Untagged Clone - SC115778

 RefSeq ORF:
 660 bp

 Locus ID:
 6016

 UniProt ID:
 Q92963

 Cytogenetics:
 1q22

**Domains:** ras, RAN, RAS, RHO, RAB

**Gene Summary:** This gene encodes a member of a subfamily of Ras-related GTPases. The encoded protein is

involved in regulating p38 MAPK-dependent signaling cascades related to cellular stress. This protein also cooperates with nerve growth factor to promote neuronal development and regeneration. Alternate splicing results in multiple transcript variants. [provided by RefSeq,

Feb 2012]

Transcript Variant: This variant (2) differs in the 5' UTR and has multiple coding region differences. These difference cause translation at a downstream start codon, compared to variant 1. The encoded isoform (2) has a shorter N-terminus, compared to isoform 2.

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.