

Product datasheet for SC326968

RASSF8 (NM 001164748) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: RASSF8 (NM_001164748) Human Untagged Clone

Tag: Tag Free Symbol: RASSF8

Synonyms: C12orf2; HOJ1

Mammalian Cell

Selection:

None

Vector: pCMV6-XL5

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

Fully Sequenced ORF: >NCBI ORF sequence for NM_001164748, the custom clone sequence may differ by one or

more nucleotides

ACAACTTGCCAGGAGGTTGTCATAGCCTTAGCTCAAGCAATAGGTCGAACTGGAAGGTAC ACCCTTATAGAGAAATGGAGAGATACTGAAAGACACTTAGCACCTCATGAAAATCCTATC ATATCCTTAAACAAATGGGGGCAGTATGCTAGTGATGTGCAGCTCATTCTACGACGAACT GGGCCGTCTCTCAGTGAGCGACCCACTTCAGACAGTGTGGCTCGAATTCCTGAAAGAACT AAAAGGAGGGAACCGAAAAGGAAATCACTGACATTTACAGGAGGTGCCAAAGGATTAATG GACATTTTTGGAAAAGGTAAAGAAACTGAGTTTAAGCAAAAGGTGCTGAATAACTGCAAA GAGAAACAGCTGGAATCTAATGAAATAGAAATAAGATTTTGGGAGCAAAAGTATAATTCC AACCTTGAAGAGGAAATTGTCCGTCTAGAGCAAAAGATCAAAAGAAACGATGTAGAAATT GAGGAGGAAGAATTCTGGGAAAATGAATTACAGATTGAACAGGAAAATGAAAAACAGCTG AAGGATCAACTTCAAGAAATAAGACAGAAATAACAGAATGTGAAAAACAAATTAAAGGAC TATTTGGCACAGATCCGGACTATGGAAAGTGGTCTTGAAGCAGAAAAATTGCAACGGGAA GTTCAAGAGGCACAGGTCAATGAGGAAGAGGTTAAAGGAAAGATCGGTAAGGTCAAAGGG GAGATTGACATTCAAGGCCAGCAGAGTCTGAGGTTGGAAAATGGCATCAAAGCTGTGGAA AGATCTCTTGGACAAGCCACCAAACGCTTACAGGACAAAGAACAGGAACTGGAGCAGTTG ACTAAGGAGTTGCGGCAAGTCAATCTCCAGCAGTTCATCCAGCAGACAGGGACAAAAGTT ACCGTTTTGCCAGCGGAGCCCATTGAAATAGAGGCCTCACATGCAGACATTGAAAGGGAG GCACCATTCCAGTCTGGGTCCCTGAAGCGACCTGGTTCATCTCGGCAGCTCCCCAGTAAT CTCCGCATTCTGCAGAATCCTATCTCATCTGGTTTTAATCCTGAAGGCATATATGTA

Restriction Sites: Please inquire ACCN: NM_001164748



OriGene Technologies, Inc. 9620 Medical Center Drive, Ste 200

CN: techsupport@origene.cn

Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com

RASSF8 (NM_001164748) Human Untagged Clone - SC326968

OTI Disclaimer: Our molecular clone sequence data has been matched to the reference identifier above as a

point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative

RNA splicing form or single nucleotide polymorphism (SNP).

OTI Annotation: This TrueClone is provided through our Custom Cloning Process that includes sub-cloning

into OriGene's pCMV6 vector and full sequencing to provide a non-variant match to the expected reference without frameshifts, and is delivered as lyophilized plasmid DNA.

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: NM 001164748.1, NP 001158220.1

 RefSeq Size:
 5561 bp

 RefSeq ORF:
 1260 bp

 Locus ID:
 11228

 UniProt ID:
 Q8NHQ8

Cytogenetics: 12p12.1

Gene Summary:

This gene encodes a member of the Ras-assocation domain family (RASSF) of tumor suppressor proteins. This gene is essential for maintaining adherens junction function in epithelial cells and has a role in epithelial cell migration. It is a lung tumor suppressor gene candidate. A chromosomal translocation t(12;22)(p11.2;q13.3) leading to the fusion of this gene and the FBLN1 gene is found in a complex type of synpolydactyly. Multiple alternatively spliced transcript variants encoding different isoforms have been found for this gene.

[provided by RefSeq, May 2011]

Transcript Variant: This variant (3) differs in the 5' UTR compared to variant 1. Variants 1, 2 and 3 encode the same isoform (a). 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.