

Product datasheet for **SC117668**

ARHGEF7 (NM_003899) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	ARHGEF7 (NM_003899) Human Untagged Clone
Tag:	Tag Free
Symbol:	ARHGEF7
Synonyms:	BETA-PIX; COOL-1; COOL1; Nbla10314; P50; P50BP; P85; P85COOL1; P85SPR; PAK3; PIXB
Mammalian Cell Selection:	None
Vector:	<u>pCMV6-XL4</u>
E. coli Selection:	Ampicillin (100 ug/mL)



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Fully Sequenced ORF: >OriGene ORF within SC117668 sequence for NM_003899 edited (data generated by NextGen Sequencing)

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ATGACCGACAATAGCAACAATCAACTGGTAGTAAGAGCAAAGTTAACTCCAGCAGACC
AATGAGGACGAGCTTTCCTTCTCAAAGGAGACGTCATCCATGTCACCCGTGTGGAAGAG
GGAGGCTGGTGGGAGGGCACACTCAACGGCCGGACCGGCTGGTCCCCAGCAACTACGTG
CGCGAGGTCAAGGCCAGCGAGAAGCCTGTGTCTCCAAATCAGGAACACTGAAGAGCCCT
CCCAAAGGATTTGATACGACTGCCATAAACAAAAGCTATTACAATGTGGTGCTACAGAAT
ATTTTAGAAACAGAAAATGAATATTCTAAAGAACTTCAGACTGTGCTTTCACGTACCTA
CGGCCATTGCAGACCAAGTGAAGTTAAGTTCAGCAAACATTTTCATATTTAATGGGAAAT
CTAGAAGAAATATGTTCTTCCAGCAAATGCTCGTACAGTCTTTAGAAGAATGCACCAAG
TTGCCGAAGCTCAGCAGAGAGTCGGAGGCTGCTTTTTAAACCTGATGCCACAGATGAAA
ACCCTGTACCTCACGTATTGTGCCAATCACCTTCTGCAGTGAATGTCCTCACGGAACAC
AGTGAGGAGTTGGGGGAGTTCATGGAGACCAAAGGTGCCAGCAGCCCTGGGATTCTCGTG
CTGACCACGGCCTGAGCAAACCTTTCATGCGCCTGGATAAATACCCTACGCTGCTCAA
GAGCTCGAGAGACACATGGAGGATTATCATAACAGATAGACAAGATATTCAAAAATCCATG
GCTGCCTTCAAAAACCTTTCAGCCCAATGTCAAGAAGTCCGGAAGAGGAAAGAGCTTGAG
CTGCAGATCCTGACGGAAGCCATCCGGAAGTGGGAGGGCGATGACATTAACCTCTGGGC
AACGTCACTTACATGTCCAGGTCCTGATTCAAGTGTGCCGGAAGTGAAGAAAAGAATGAA
AGATATCTTCTACTCTTCCCAAATGTTTTGCTAATGTTGTCTGCCAGTCTTAGGATGAGT
GGCTTTATCTATCAGGGAAAGCTTCCAACGACAGGAATGACAATCACAAAGCTTGAGGAC
AGTGAAAATCATAGAAATGCATTTGAAATATCAGGGAGCATGATTGAGCGGATATTAGTG
TCGTGCAACAACCAGCAGGATCTGCAGGAATGGGTGGAGCACCTACAGAAGCAAACGAAG
GTCACGTCTGTGGGAAACCCCAACATAAAGCCTCATTCAAGTCCATCTCATACCCTCCCC
TCCCACCCGGTCACTCCGTCCAGCAAGCACGCAGACAGCAAGCCCGCGCCGCTGACGCC
GCCTACCAACAGCTGCCCAACCCCTCCCACACGGCACCCCGCACACCACCATCAACTGG
GGACCCCTGGAGCCTCCGAAAACACCAAGCCCTGGAGCCTGAGCTGCCTGCGGCCCGCG
CCTCCCTCCGGCCCTCAGCTGCTCTGCTACAAGGAGGATCTTAGTAAGAGCCCTAAG
ACCATGAAAAGCTGCTGCCAAGCGCAAACCTGAACGGAAGCCTTCAAGTGAAGGAGTTC
GCGTCCCAGAAAAGCACAGCTGCTTTGGAAGAAGATGCTCAGATTCTGAAAGTCATTGAA
GCTTACTGCACCAGCGCCAAAACAAGGCAAACACTCAATTCAAGTTCACGCAAAGATCT
GCTCCACAAGTTTTGCTTCCAGAAGAAGAGAAAATTATAGTGGAAGAACTAAAAGTAAT
GGTCAGACAGTGATAGAAGAAAAGAGTCTTGTGGATACCGTATATGCATTAAGGATGAA
GTTCAAGAATTAAGACAGGACAACAAAAGATGAAGAAATCTCTAGAGGAAGAACAGAGA
GCCCGCAAAGACCTGGAGAAGCTGGTGAGGAAAGTCTGAAGAACATGAATGATCCTGCC
TGGGATGAGACCAATCTATAA
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Clone variation with respect to NM_003899.3

9 t=>c

5' Read Nucleotide Sequence:

>OriGene 5' read for NM_003899 unedited
 CGTTANCATTTGTATACGACTCATATAGGCGGCCGCAATTCGGCACGAGGGCAAACCGT
 TACGTGGCTCATCACTCTGGGGTGTGGAGTCGCCCAAAAAACCATCTCGGACCCGGA
 GGGCTTTCTGCAGGCGTCGCTGAAGGATGGGGTGGTCCTCTGCAGGCTGCTGGAGCGCT
 GCTCCCCGGGACCATCGAGAAAGTCTACCCCGAGCCCGGAGCGAGAGCGAGTGCCTGAG
 CAACATCCGCGAGTTCCTGCGCGGCTGCGGGGCTTCCCTGCGGCTGGAGACGTTTGATGC
 AAATGATTTGTATCAGGGCAGAATTTTAAACAAGTCTCAGTTCCTTAGTGACTCTAAA
 TAAAGTAACAGCAGACATCGGGCTGGGGAGTGACTCCGTGTGTGCCCGCCCTCGTCTCA
 CCGCATAAAGTCTTTTACTCCCTTGGATCACAGTCTTTGCACACTCGGACTTCAAAACT
 GTTCCAGGGCCAGTATCGGAGTTTGACATGACCGACAATAGCAACAATCAACTGGTAGT
 AAGAGCAAAGTTTAACTCCAGCAGACCAATGAGGACGAGCTTTCCTTCTCAAAGGAGAC
 GTCATCCATGTCACCCGTGTGGAATAGGAGGCTGTGGNGGAAGGGCACACTCAACGGCCG
 GACCGGCTGGTCCCAGCAACTACGTGCGCGAGGTCAAGGCCAGCGAGAAGCCTGTGTCT
 CCCATATCATGACTGAAGAGCCCTCCAAGGATTTGATACGACTGCCATANNACAAC
 ATTACAATGTGGTGTACAGAATATTTTAGAAACAGATNATGAATATTCTAAAGAATTNC
 AGACTGTGCTTTTACGTACCTACGGCCATTGCAGACANTGGAAAAAGTTAAGTTCAGC
 ANACATTTCTNATTTAATGGNGAATNCTAGAAGAATTAATGCCCTT

3' Read Nucleotide Sequence:

>OriGene 3' read for NM_003899 unedited
 TGCTCTGGACCGCGCCGAATCTAGGGTCGAGTTTTTTTTTTTTTTTTTTTTTTTTTTT
 ACATAAGAAAAAATCAGTTTAATAATTACAACAGTGTAGCATCAACATTGTAGTAGTCA
 AACTAGGACATTGTCTGTGACATTTAAATTAGGAGTTTCTCCACTTGTCAACAGGCG
 CCCTGGGTGTCCAGAGAAGTGAAGAAATGATTGGCATTAAAGGAGTCCACACAACAGGG
 CAGGCATTCAAGGTCCCAGACGCCAGTGCATGTGGACAAGCACAGACGCCACTGAGTGT
 GAAGGAACGAGCTTCTGATGCTACATTTCAACATCGCTGAAAAGTAAAGTGTTCACAT
 CTACAGGATAAGAAATGGCTTCTCCTCATCCATACAAGGTGAGAGCAAAGGAGTAAC
 CCTCGGGCAAATCGTGGCCCTTTAAACGTTTATCAAGTTCACCCCTTGGGAGGT
 CACACGGTGAAACAGACAGTTATATAACAACAGGGCAGTTTTTAAAAAGAGTTGCTCT
 CAGACGCATTTTCTGCTCCCTAAAAAGCCGAGGAAGATACTGGATCCACAGAAAGAAA
 AGGCAATGCCGTAACATGAGCCCTCATGGCCGACCGTCCAGGGGAAGGGCTGNTAAAA
 ACACAAGTATTCTGTNGAATACTTCGATCTGAGCATTAAAGGCAGTCTGCAGGAGATCC
 GTCTGGGGACTCCGACAGCAACGCTACCGCTTCGAGAGGACAGTTAATGTCGCCCTCC
 CGCAGAGGGGCCGAGAGATCAAACAAGAGTTGTTCTGAGTTAAAAACTGCTACACAGCA
 AACTCCCATAACTCAAGGTGAATGTTTTCCACTGGCATGCTTTACGCATACCAAAAATAA
 CTTGTAAACTCAGCTACCAAAAGAGTGTGACTCTGCTGGGCCCTGATGAATAACC
 GCTAN

Restriction Sites:

NotI-NotI

ACCN:

NM_003899

Insert Size:

5130 bp

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).

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_003899.2](#), [NP_003890.1](#)

RefSeq Size: 5032 bp

RefSeq ORF: 1941 bp

Locus ID: 8874

UniProt ID: [Q14155](#)

Cytogenetics: 13q34

Domains: RhoGEF, SH3, PH

Protein Pathways: Regulation of actin cytoskeleton

Gene Summary: This gene encodes a protein that belongs to a family of cytoplasmic proteins that activate the Ras-like family of Rho proteins by exchanging bound GDP for GTP. It forms a complex with the small GTP binding protein Rac1 and recruits Rac1 to membrane ruffles and to focal adhesions. Multiple alternatively spliced transcript variants encoding different isoforms have been observed for this gene. [provided by RefSeq, Mar 2016]

Transcript Variant: This variant (1) differs in the 5' UTR, lacks a portion of the 5' coding region, and contains a different 3' coding region and 3' UTR, compared to variant 1. It initiates translation at a downstream start codon. Variants 1, 5, 6, 18, and 19 all encode isoform a (also known as p85Cool-1), which has a shorter N-terminus and a longer and distinct C-terminus, compared to isoform c. Sequence Note: The RefSeq transcript and protein were derived from genomic sequence to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on alignments.