

Product datasheet for MC229461

Arap1 (NM_027180) Mouse Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	Arap1 (NM_027180) Mouse Untagged Clone
Tag:	Tag Free
Symbol:	Arap1
Synonyms:	2410002L19Rik; Centd2; mKIAA0782
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Cell Selection:	Neomycin
Fully Sequenced ORF:	>MC229461 representing NM_027180 Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCCGCGATCGCC

ATGACCAAGGAGGAGCCCTGCCGAGCCGAGTCCCACGGGCTGTGCGTGTGGCTAGTCTGCTGAGCGAGG
GGGAGGAGCTGTCTGGGGATGATTCGGAAGATGACGATGACCATGCCTATGAGGGTATCCCAATGGTGG
GTGGCCAAACGAGTGGCCTGAATCCACCCTTACGCAGCCTGATCCCTGATCTCCACTGCACCCCATGGAT
GAGTTGCCTGGGGTCCCACCCCATCACCTGTCAAGGCTGGCTGGTGGACAAGAACCACCCAC
AGGGGTCTTATCTATCAGAAGCGATGGGTGAGACTGGACGCTGATTACCTGCGATACTTTGACAGTAA
CAAGGACGCCTACTCTAAGCGCTTTGTTCTGTGGCCTGCATCTGCCGAGTAGCTCCTATTGGAGACCAG
AAGTTTGAAGTGATCACAAATAATCGGACCTTCGCCTTCGGGCGAGAGAGTGATGTGGAGCGGAACGAGT
GGATGCAGGCCCTGCAGCAGGCAGTAGTTGAGCATCGAGCCGTTTTTCGGCTTTCTAGTGCTTCTGTGTT
GGGAGTCCGAGGCTCCGAGCAGCCTGACCGTGTGGCAGCCTGGAGCTACGAGGCTTCAAGAATAAGCTT
TACGTGGCTGTGACTGGGACAAAGTACAGCTCTATAAGAATCTGGAGGAATCCATCTGGGCAATTGGTA
TCACCTTTATCGACATGAATGTGGCAATGTCAAGGAAGTGGATCGGCGCAGCTTTGACCTCACTACCCC
CTACCGCATCTTCAGCTTCTCGGCTGATTCAGAGTTAGAGAAGGAACAGTGGCTGGAGGCCATGCAGGGA
GCCATCGCAGAGGCCCTATCTACCTCAGAGGTGGCTGAGCGCATCTGGGCCGAGCCCCAACAGTTCT
GTGCCGACTGTGGGGCCGCCAGCCTGACTGGGCTCCATCAACCTCTGCGTTGTGATCTGCAAGCGCTG
TGCAGGGGAGCACCGCGCCTGGGTGCAGGAGTGTCCAAGGTGCGGAGTTTGAAGATGGACAGGAAGGTG
TGGACAGAAGCACTCATCCAGCTCTTCTACATCTGGGCAATGGCCCCGGAACCACTTCTGGGCTGCTA
ATGTGCCTCCTAGTGAGGCCCTGGAGCCAGCAGCAGCCCTGGTCCCGCGGTATCACCTGGAGGCCAA
GTACAGAGAGGGCAAGTACCGCCGCTACCATCCGCTCTTTGGCAACCAAGAGGAAGTGGACAAGGCCCTG
TGTGCTGCAGTTACTACCACTGACCTGGCTGAGACCCAGGCACTCCTGGGCTGTGGGGCTGGGGTCACT
GCTTCTCAGGGGACCCAGCAGCTCCACACCCCTGGCTTTGCTGAGCAGGCCGACAGACTCTGCAGAT
GGAGTTTCTACGGAATAATCAGAGTACAGAGGTCCCTCGTTGGACTCAGTGAAGCCCTTGGAAAAGCAC
TACTCCGTTACCTGCCAACTGTGAGCCACAGCGGCTTCTGTACAAAAGTCTTCTGCCGGCAAACCTC



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TGCAGGACCGCCGTGCCCGGAAGAGTTCAGCCGGCGCTGGTGTGTCCTGAGTGATGGGGTCTGAGCTA
 CTATGAAAACGAACGGGCAGTGACACCCAATGGGGAGATTTCGGCCAGCGAGATAGTATGCCTAGCAGTT
 TCCCCTCTGGACACCCATGGCTTTGAGCACACCTTTGAGGTGTACACAGAGGGAGAACGGCTGTACCTGT
 TTGGGCTGGAGAATGCAGAGCTGGCTCATGAGTGGTCAAGTGCATTGCCAAGGCGTTCGTGCCTCCCCT
 GGCTGAGGACCTGCTAGCCCGGACTTTGAACGTCTTGGGCGCCTACCCTGTAAGCTGGCTGAGCCTT
 CAGCAGGCTCAGGAAGGCTGGTTTGCCTTGACTGGCTCTGAGCTCCGGGCTGTCTCCCAGAGGGGCCCT
 GGAAGAGCCGCTGCAGCTCCGAAACTGCAAGAGCTTCTATCCAAGGAGACAGCGAGAACAAGTGCT
 GGTGCTGGTGGAGCGGAGGAGGACACTGTACATCCAGGGTGAGCGGCGGCTGGACTTCATGGCTTGGCTA
 GGGGTCATCCAGAAAGCAGCAGCCAGCTTGGGAGACACTATCAGAACAACAGCTTGGGGACTCGGACA
 TCCAGTGATTGTGTACCCTGTGTGGACTACATCACGCAGTGTGGCTTACCTCAGAGGGTATCTATCG
 AAAGTGTGGTCAGACCTCAAGACTCAGAGACTGCTAGACAGCCTCCGGCAGGACGCACGCTCTGTGCAC
 CTAAGGAGGGAGAACAGCACGTGGACGACGTCTCTCTGCACCTCAAACGCTTCCCTCAGGACCTGCCCC
 ATGGGCTCTTACGCGTGCAGCGCCTGGCCTGGCTGGAGGCCCTCTGAGATCGAGGATGAGGAAGAAAA
 GATCTCCAGGTATCGAGAGCTCTTGGTGCATCTGCCCCCTGTCAACCGGGCCACTGTGAAGGCCCTTATC
 AGCCATCTGTACTGTGTACAGTCTTCTCAGACACAAACCAAATGAACACACACAACCTGGCTATCGTGT
 TTGGGCTACACTCTTCCAGACAGACGGCAGGACTACAAGGCCGGAAGGTGGTGAAGACCTCATCAA
 CCACTACGTGGTGGTGTTCAGTGTGGACGAGGAGGAGCTGAGGAAGCAGAGGGAGGAAGTACCGCCATC
 GTGAAGATGCGAGTGGCTGGCACTGCCAGTGGGACCCAGCATGCTGGCGACTTTCATCTGCACAGTCTACC
 TGGAGGAGAAGAAGGTGGAGACTGAACAGCATGTTAAGATCCCAGCATCCATGACTGCAGAGGAGCTTAC
 TCTGGAGATTCTGGACCGCCGCAATGTGAGCATCAGGGAGAAGGACTACTGGACTTGCTTTGAGGTCAAC
 GAGAAGGAGGAGGAGCAGCGCCCGTGCACCTTGCAGAGAAGGTGCTGCCATTGTCCATGGGCTGGGCA
 TAGACAGCCATCTGGTGGTGAAGAAGTACCAGTCCATGGAGGCCATGCTGTTGTACTTGGCCAGCCGTGT
 GGGTGACACCAAGCATGGTATGATGAAGTTCCTGGAAGACCGCAGCCTCCTGGGCTGGGCTACCTTCG
 GGTGGCTTCCAGCATCGCTACTTCACTTCAACAGCAGCTGCCTACGGCTCTACAAGGAGGTCCGAGTCC
 ACCGGCCTGAGAAGGAGTGGCCTGTCAAGAGCCTCAAAGTCTACCTGGGTGTAAGAAAGAAACTGCGGCC
 ACCTACTTGCTGGGGCTTACGGTGGTGCAGAGACAGAAAAGCAGAGAAGCAGCAGTGGTACCTCTGC
 TGTGACACGCAGATGAACTTCGAGAGTGGTTTGCACCTTCTCTGTGCAGCAGCATGGCCTGGTGT
 GGCCCTCCGAGCCATCTCGAGTGTCCCGGCGAGTGCCTGAGGTCCGGATGGGCAGTGTATCGCTGATCCC
 TCTACGAGGCAGTAAAATGAAATGCGCCGGAGTGTGGCAGCCTTCACTGCTGACCCCTTCCCTTCTC
 CGCCATGCTGA

AGCGGACCGACGCGTACGCGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCC
 TGGATTACAAGGATGACGACGATAAGGTTTAA

- Restriction Sites:** SgfI-RsrII
- ACCN:** NM_027180
- Insert Size:** 3582 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).
- OTI Annotation:** Clone contains native stop codon, and expresses the complete ORF without any c-terminal tag.
- 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_027180.3](#), [NP_081456.2](#)

RefSeq Size: 4929 bp

RefSeq ORF: 3582 bp

Locus ID: 69710

UniProt ID: [Q4LDD4](#)

Cytogenetics: 7 E2

Gene Summary: Phosphatidylinositol 3,4,5-trisphosphate-dependent GTPase-activating protein that modulates actin cytoskeleton remodeling by regulating ARF and RHO family members. Is activated by phosphatidylinositol 3,4,5-trisphosphate (PtdIns(3,4,5)P3) binding. Can be activated by phosphatidylinositol 3,4-bisphosphate (PtdIns(3,4,5)P2) binding, albeit with lower efficiency. Has a preference for ARF1 and ARF5 (By similarity).[UniProtKB/Swiss-Prot Function]
Transcript Variant: This variant (1) differs in the 5' UTR and lacks some 5' coding sequence, compared to variant 3. These differences cause translation initiation at a downstream AUG and an isoform (1) with a shorter N-terminus compared to isoform 3. Variants 1 and 4 encode the same protein.