

Product datasheet for **SC202041**

ARHGAP8 (NM_001017526) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	ARHGAP8 (NM_001017526) Human 3' UTR Clone
Symbol:	ARHGAP8
Synonyms:	BPGAP1; PP610
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_001017526
Insert Size:	219 bp
Insert Sequence:	<p>>SC202041 3'UTR clone of NM_001017526</p> <p>The sequence shown below is from the reference sequence of NM_001017526. The complete sequence of this clone may contain minor differences, such as SNPs.</p> <p>Blue=Stop Codon Red=Cloning site</p> <pre> GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAACGATCGCC CCCCTGATGGCAGCCAGAAGACGTCTCTAGTGTTCGGAACACTCTGTATATTTTCGAGCTACCTCCCACA CCTGTCTGTGCACTTGTATGTTTGTAACTTGGCATCTGTAAAAATAACCAGCCATTAGATGAATTCA GAACCTTCTAATGAAACTCCATGCCTCTGGTCTTGACTCTTGTCATGGTTCTGAGCTGTGGACC GGGATAGAATAA ACGCGTAAGCGGCCGCGCATCTAGATTGAAGAAAATGACCGACCAAGCGACGCCAACCTGCCATCA CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG </pre>
Restriction Sites:	SgfI-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:	<u>NM_001017526.2</u>


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Summary:

This gene encodes a member of the RHOGAP family. GAP (GTPase-activating) family proteins participate in signaling pathways that regulate cell processes involved in cytoskeletal changes. GAP proteins alternate between an active (GTP-bound) and inactive (GDP-bound) state based on the GTP:GDP ratio in the cell. This family member is a multidomain protein that functions to promote Erk activation and cell motility. Alternative splicing results in multiple transcript variants. Read-through transcripts from the upstream proline rich 5, renal (PRR5) gene into this gene also exist, which led to the original description of PRR5 and ARHGAP8 being a single gene. [provided by RefSeq, Nov 2010]

Locus ID:

23779

MW:

8