

## Product datasheet for **SC332696**

### ARHGAP20 (NM\_001258416) Human Untagged Clone

#### Product data:

**Product Type:** Expression Plasmids  
**Product Name:** ARHGAP20 (NM\_001258416) Human Untagged Clone  
**Tag:** Tag Free  
**Symbol:** ARHGAP20  
**Synonyms:** RARHOGAP  
**Vector:** pCMV6-Entry (PS100001)  
**Fully Sequenced ORF:** >SC332696 representing NM\_001258416.  
Blue=Insert sequence Red=Cloning site Green=Tag(s)

```
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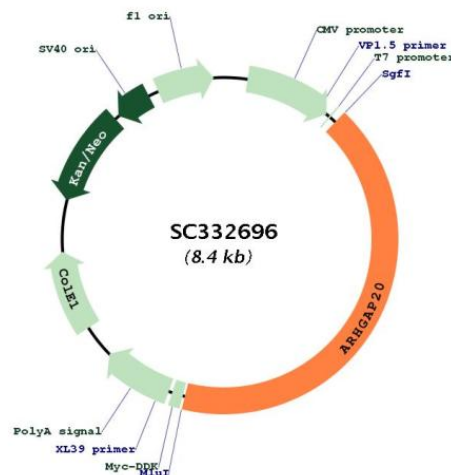
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```

Restriction Sites:

SgfI-MluI

Plasmid Map:



ACCN:

NM\_001258416

Insert Size:

3498 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).

<b>Components:</b>	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).
<b>Reconstitution Method:</b>	<ol style="list-style-type: none"><li>1. Centrifuge at 5,000xg for 5min.</li><li>2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.</li><li>3. Close the tube and incubate for 10 minutes at room temperature.</li><li>4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.</li><li>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.</li></ol>
<b>RefSeq:</b>	<u>NM_001258416.1</u>
<b>RefSeq Size:</b>	6085 bp
<b>RefSeq ORF:</b>	3498 bp
<b>Locus ID:</b>	57569
<b>UniProt ID:</b>	<u>Q9P2F6</u>
<b>Cytogenetics:</b>	11q22.3-q23.1
<b>MW:</b>	130.2 kDa
<b>Gene Summary:</b>	<p>The protein encoded by this gene is an activator of RHO-type GTPases, transducing a signal from RAP1 to RHO and impacting neurite outgrowth. [provided by RefSeq, Sep 2016]</p> <p>Transcript Variant: This variant (3) differs in the 5' UTR and initiates translation at an alternate start codon, compared to variant 1. The encoded isoform (3) is shorter and has a distinct N-terminus, compared to isoform 1. 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.</p>