

Product datasheet for **SC331166**

ADAR1 (ADAR) (NM_001193495) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	ADAR1 (ADAR) (NM_001193495) Human Untagged Clone
Tag:	Tag Free
Symbol:	ADAR1
Synonyms:	ADAR1; AGS6; DRADA; DSH; DSRAD; G1P1; IFI-4; IFI4; K88DSRBP; P136
Vector:	pCMV6-Entry (PS100001)



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Fully Sequenced ORF: >SC331166 representing NM_001193495.
 Blue=Insert sequence Red=Cloning site Green=Tag(s)

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ATGGCCGAGATCAAGGAGAAAATCTGCGACTATCTTTCAATGTGTCTGACTCCTCTGCCCTGAATTTG
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Restriction Sites: SgfI-MluI
ACCN: NM_001193495
Insert Size: 2796 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:	<ol style="list-style-type: none">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_001193495.1
RefSeq Size:	6730 bp
RefSeq ORF:	2796 bp
Locus ID:	103
UniProt ID:	P55265
Cytogenetics:	1q21.3
Protein Families:	Druggable Genome
Protein Pathways:	Cytosolic DNA-sensing pathway
MW:	103.6 kDa
Gene Summary:	<p>This gene encodes the enzyme responsible for RNA editing by site-specific deamination of adenosines. This enzyme destabilizes double-stranded RNA through conversion of adenosine to inosine. Mutations in this gene have been associated with dyschromatosis symmetrica hereditaria. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Jul 2010]</p> <p>Transcript Variant: This variant (5) differs in the 5' UTR, lacks a portion of the 5' coding region, and uses a downstream start codon, compared to variant 1. The resulting isoform (d) is shorter at the N-terminus, compared to isoform a. Variants 4, 5, 7, 8, and 9, encode the same isoform. 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>