

Product datasheet for SC118041

DAP12 (TYROBP) (NM_003332) Human Untagged Clone

Product data:

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Product Type:	Expression Plasmids
Product Name:	DAP12 (TYROBP) (NM_003332) Human Untagged Clone
Tag:	Tag Free
Symbol:	DAP12
Synonyms:	DAP12; KARAP; PLOSL; PLOSL1
Mammalian Cell Selection:	None
Vector:	pCMV6-XL4
E. coli Selection:	Ampicillin (100 ug/mL)
Fully Sequenced ORF:	<pre>>OriGene ORF within SC118041 sequence for NM_003332 edited (data generated by NextGen Sequencing) ATGGGGGGACTTGAACCCTGCAGCAGGCTCCTGCTCCTGCTCTCCTGCTGGCTG</pre>
5' Read Nucleotide Sequence:	>OriGene 5' read for NM_003332 unedited CCCACCGGCCCTTACACTGTGGTGTCCAGCAGCATCCGGCTTCATGGGGGGGACTTGAACC CTGCAGCAGGCTCCTGCTCCTGCCTGCTGGCTGTAAGTGGTCTCCGTCCTGTCCA GGCCCAGGCCCAGAGCGATTGCAGTTGCTCTACGGTGAGCCCGGGCGTGCTGGCAGGGAT CGTGATGGGAGACCTGGTGCTGACAGTGCTCATTGCCCTGGCCGTGTACTTCCTGGGCCG GCTGGTCCCTCGGGGGCGAGGGGCTGCGGAGGCAGCGACCCGGAAACAGCGTATCACTGA GACCGAGTCGCCTTATCAGGAGGCTCCAGGGTCAGAGGTCGGATGTCTACAGCGACCTCAA CACACAGAGGCCGTATTACAAATGAGCCCGAATCATGACAGTCAGCAACATGATACCTGG ATCCAGCCATTCCTGAAGCCCCGCACCGCA



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3' Read Nucleotide Sequence:	<pre>>OriGene 3' read for NM_003332 unedited GGGGGGGNNNGGGGGGNNNNNNNNNNNNNTTTTNNNNNAANTTTTACTTAGNACCGNCGN CCGCATACTANGATCGNGTTTTTTTTTTTTTTTTTTTTTT</pre>
Restriction Sites:	Notl-Notl
ACCN:	NM_003332
Insert Size:	630 bp
OTI Disclaimer:	Due to the inherent nature of this plasmid, standard methods to replicate additional amounts of DNA in E. coli are highly likely to result in mutations and/or rearrangements. Therefore, OriGene does not guarantee the capability to replicate this plasmid DNA. Additional amounts of DNA can be purchased from OriGene with batch-specific, full-sequence verification at a reduced cost. Please contact our customer care team at <u>custsupport@origene.com</u> or by calling 301.340.3188 option 3 for pricing and delivery. The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing variants is recommended prior to use. <u>More info</u>
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:	 Centrifuge at 5,000xg for 5min. Carefully open the tube and add 100ul of sterile water to dissolve the DNA. Close the tube and incubate for 10 minutes at room temperature. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom. 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:	<u>NM 003332.2, NP 003323.1</u>
RefSeq Size:	583 bp

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RefSeq ORF: Locus ID: UniProt ID: Cytogenetics: Protein Families:	342 bp 7305 <u>O43914</u> 19q13.12 Druggable Genome, Transmembrane
Protein Pathways:	Natural killer cell mediated cytotoxicity
Gene Summary:	This gene encodes a transmembrane signaling polypeptide which contains an immunoreceptor tyrosine-based activation motif (ITAM) in its cytoplasmic domain. The encoded protein may associate with the killer-cell inhibitory receptor (KIR) family of membrane glycoproteins and may act as an activating signal transduction element. This protein may bind zeta-chain (TCR) associated protein kinase 70kDa (ZAP-70) and spleen tyrosine kinase (SYK) and play a role in signal transduction, bone modeling, brain myelination, and inflammation. Mutations within this gene have been associated with polycystic lipomembranous osteodysplasia with sclerosing leukoencephalopathy (PLOSL), also known as Nasu-Hakola disease. Its putative receptor, triggering receptor expressed on myeloid cells 2 (TREM2), also causes PLOSL. Multiple alternative transcript variants encoding distinct isoforms have been identified for this gene. [provided by RefSeq, Mar 2010] Transcript Variant: This variant (1) represents the longest transcript and encodes the longest isoform (1).