

Product datasheet for MC216857

Steap2 (NM_001103156) Mouse Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	Steap2 (NM_001103156) Mouse Untagged Clone
Tag:	Tag Free
Symbol:	Steap2
Synonyms:	4921538B17Rik; AI930049; AW045895; IPCA-1; IPCA1; PCANAP1; STAMP1; STMP
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Fully Sequenced ORF:	>NCBI ORF sequence for NM_001103156, the custom clone sequence may differ by one or more nucleotides

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ATGGAATCCATCTCTATGATGGGAAGCCCTAAGAGCCTGGAGACGTTTTTGCCTAATGGCATAAATGGTA
TCAAAGACGCAAGGCAGGTACCAGTGGGGGTGATAGGAAGTGGGGATTTTGCCAAGTCTCTGACCATTCC
GCTTATTAGATGTGGCTACCACGTGGTCATAGGAAGCAGAAAATCCAAGTTTGCATCAGAATTTTTTCT
CACGTGGTAGACGTACCCACCATGAAGATGCTTAAACAAAAACAATAAATATTCGTGGCTATCCATA
GAGAACATTACACCTCCTTGTGGGACCTGAGACATCTGCTTGTGGGCAAATCCTCATTGATGTGAGCAA
CAACATGAGAGTAAACCAGTACCAGAATCCAATGCAGAGTACCTGGCTCATTATCCCGGACTCCTTG
ATTGTCAAAGGATTTAATGTGATCTCAGCTTGGGCACTTCAGCTAGGTCCAAGGATGCCAGCCGCCAGG
TTTATATATGCAGCAACAATATCCAAGCTCGACAGCAGGTTATCGAGCTCGCCCGCCAGCTGAATTTAT
TCCTGTTGACTTGGGATCTTTGTCGTCAGCCAAGGAGATTGAAAACCTTACCTCTGCGACTGTTTACTCTC
TGGAGGGGGCCAGTGGTAGTAGCCATAAGCTTGGCCACATTTTTCTTTTATTCTTTTGTGAGAGATG
TGATACATCCATATGCCAGAAACCAGCAGAGTGACTTTTACAAGATCCCATTGAGATTGTGAACAAAAC
CTTGCCGATCGTCGCCATCACCTGCTGTCTCTGGTGTACCTGGCTGGCCTCCTGGCAGCTGCGTATCAG
CTTTATTATGGCACTAAGTACCGCCGATTTCCCCCGTGGCTGGATACTTGGCTGCAGTGCAGGAAACAGC
TGGGATTGCTGAGCTTCTTCTTTCAGTGTTCACGTAGCCTACAGCCTCTGCTTACCAATGAGGAGGTC
GGAAAGATACCTGTTCCCAACATGGCTTATCAGCAGGTTCAATGCAATATTGAGAACCGGTGGAACGAG
GAGGAGGTCTGGAGGATTGAGATGTACATTTTCTTTGGCATCATGAGCCTGGGCTTGTGTCCCTGCTGG
CGGTCACTTCCATCCCATCAGTGAAGCAACGCTTTGAACTGGAGAGAGTTAGTTTTCATCCAGTCTACGCT
TGGCTACGTCGCCCTGCTCATCACGACCTTCCACGTGTTAATTTACGGATGGAAGCGTGCCTTTGCAGAA
GAGTACTACCGCTTTTACACACCACAACTTCGTTCTTGGCCCTCGTTTTGCCCTCCATTGTAATTTCTGG
GTAAGATGATATTACTCTCCCATGCATAAGCCGAAAGCTAAAACGAATAAAAAGGGCTGGGAAAAGAG
CCAGTTTCTAGACGAAGGCATGGGAGGAGCGGTTCTCATCTGTCCCGAGAGAGGTCACAGTGTGTGA

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Restriction Sites:	Sgfl-RsrII
ACCN:	NM_001103156
Insert Size:	1470 bp
OTI Disclaimer:	<p>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 custsupport@origene.com or by calling 301.340.3188 option 3 for pricing and delivery.</p> <p>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).</p>
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:	<u>BC150997</u> , <u>AAI50998</u>
RefSeq Size:	6320 bp
RefSeq ORF:	1470 bp
Locus ID:	74051
UniProt ID:	<u>Q8BWB6</u>
Cytogenetics:	5 A1
Gene Summary:	<p>Metalloreductase that has the ability to reduce both Fe(3+) to Fe(2+) and Cu(2+) to Cu(1+). Uses NAD(+) as acceptor.[UniProtKB/Swiss-Prot Function]</p> <p>Transcript Variant: This variant (3) differs in the 5' UTR compared to variant 1. Variants 1, 2, 3, 4, and 5 encode the same 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>