

## Product datasheet for SC126655

### COPA (NM\_004371) Human Untagged Clone

#### Product data:

Product Type:	Expression Plasmids
Product Name:	COPA (NM_004371) Human Untagged Clone
Tag:	Tag Free
Symbol:	COPA
Synonyms:	AIJK; alpha-COP; HEP-COP
Mammalian Cell Selection:	None
Vector:	<u>pCMV6-XL4</u>
E. coli Selection:	Ampicillin (100 ug/mL)
Fully Sequenced ORF:	>OriGene ORF within SC126655 sequence for NM_004371 edited (data generated by NextGen Sequencing)

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ATGTTAACCAAATTCGAGACCAAGAGCGCGGGTCAAAGGGCTCAGCTTTCACCCAAA
AGACCTTGGATCCTGACTAGTTTACATAATGGGGTCATCCAGTTATGGGACTATCGGATG
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TCACATGTAGCACTACTAGCCAAACACGCCATTGTGATCTGTAACCGCAAACCTGGATGCT  
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TGCAGGGTCACCACAGTGACAGAGATTGGCAAAGATGTGATTGGTTTAAGGATCAGTCCT  
CTGCAGTTTCGTAA

Clone variation with respect to NM\_004371.3  
363 c=>t

<b>5' Read Nucleotide Sequence:</b>	>OriGene 5' read for NM_004371 unedited GCCGCGAATTCGGCACGAGCTGACGTGGAGGCGTCCGAAGGGCAGCAGGGTGTGTCGGGG CTCGGATTAAGACATCGGAGTCGGAGACCTGAGAGATGTTAACCAAATTCGAGACCAAGA GCGCGCGGGTCAAAGGGCTCAGCTTTCACCCAAAAGACCTTGGATCCTGACTAGTTTAC ATAATGGGGTCATCCAGTTATGGGACTATCGGATGTGCACTCTCATTGACAAGTTTGATG AACATGATGGTCCAGTGGCAGGCATTGACTTCCATAAGCAGCAGCCACTGTTTCGTCTCTG GAGGAGATGACTATAAGATTAAGGTTTGGAAATTACAAGCTTCGGCGCTGTCTTTTCACAT TGCTTGGGCACCTTAGATTATATTCGCACCACGTTTTTTCATCATGAATATCCCTGGATTC TGAGTGCCTCCGATGATCAGACCATCCGAGTGTGGAATTGGCAATCTAGAACCTGTGTTT GTGTGTTAACAGGGCACAACCATTATGTGATGTGTGCTCAGTTCCACCCACAGAAGACT TGGTAGTATCAGCCAGCCTGGACCAGACTGTGCGCGTTTGGGATATTTCTGGTCTGAGGA AAAAAACCTGTCCCCTGGTGGTGGAAATCGGATGTGAGAGGAATAACTGGGGTTGATC TATNTGGAACACAGATGCAGTGGTGAAGCATGTACTAGAGGGTCACGATCGTGGAGTAA ACTGGGN
<b>Restriction Sites:</b>	NotI-NotI
<b>ACCN:</b>	NM_004371
<b>Insert Size:</b>	4310 bp
<b>OTI Disclaimer:</b>	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>	<a href="#">NM_004371.2</a> , <a href="#">NP_004362.1</a>
<b>RefSeq Size:</b>	5064 bp
<b>RefSeq ORF:</b>	3675 bp
<b>Locus ID:</b>	1314
<b>UniProt ID:</b>	<a href="#">P53621</a>
<b>Cytogenetics:</b>	1q23.2
<b>Domains:</b>	WD40, Coatomer_WDAD

**Gene Summary:**

In eukaryotic cells, protein transport between the endoplasmic reticulum and Golgi compartments is mediated in part by non-clathrin-coated vesicular coat proteins (COPs). Seven coat proteins have been identified, and they represent subunits of a complex known as coatomer. The subunits are designated alpha-COP, beta-COP, beta-prime-COP, gamma-COP, delta-COP, epsilon-COP, and zeta-COP. The alpha-COP, encoded by COPA, shares high sequence similarity with RET1P, the alpha subunit of the coatomer complex in yeast. Also, the N-terminal 25 amino acids of alpha-COP encode the bioactive peptide, xenin, which stimulates exocrine pancreatic secretion and may act as a gastrointestinal hormone. Alternative splicing results in multiple splice forms encoding distinct isoforms. [provided by RefSeq, Jul 2008]

Transcript Variant: This variant (2) uses an alternate in-frame splice site in the 3' coding region, compared to variant 1, resulting in a shorter protein (isoform 2). 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.