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## Product datasheet for SC108887

## ACOX1 (NM_007292) Human Untagged Clone

## Product data:

Product Type: Expression Plasmids
Product Name: ACOX1 (NM_007292) Human Untagged Clone
Tag:
Symbol:
Synonyms:
Mammalian Cell
Selection:
Vector:
E. coli Selection:

Tag Free
ACOX1
ACOX; MITCH; PALMCOX; SCOX
None

## pCMV6-XL4

Ampicillin ( $100 \mathrm{ug} / \mathrm{mL}$ )

Fully Sequenced ORF: >NCBI ORF sequence for NM_007292, the custom clone sequence may differ by one or more nucleotides

ATGAACCCGGACCTGCGCAGGGAGCGGGATTCCGCCAGCTTCAACCCGGAGCTGCTTACACACATCCTGG ACGGCAGCCCCGAGAAAACCCGGCGCCGCCGAGAGATCGAGAACATGATCCTGAACGACCCAGACTTCCA GCATGAGGACTTGAACTTCCTCACTCGCAGCCAGCGTTATGAGGTGGCTGTCAGGAAAAGTGCCATCATG GTGAAGAAGATGAGGGAGTTTGGCATCGCTGACCCTGATGAAATTATGTGGTTTAAAAAACTACATTTGG TCAATTTTGTGGAACCTGTGGGCCTCAATTACTCCATGTTTATTCCTACCTTGCTGAATCAGGGCACCAC TGCTCAGAAAGAGAAATGGCTGCTTTCATCCAAAGGACTCCAGATAATTGGCACCTACGCCCAGACGGAA ATGGGCCACGGAACTCACCTTCGAGGCTTGGAAACCACAGCCACGTATGACCCTGAAACCCAGGAGTTCA TTCTCAACAGTCCTACTGTGACCTCCATTAAATGGTGGCCTGGTGGGCTTGGAAAGACTTCAAATCATGC AATAGTTCTTGCCCAGCTCATCACTAAGGGGAAATGCTATGGATTACATGCCTTTATCGTACCTATTCGT GAAATCGGGACCCATAAGCCTTTGCCAGGAATTACCGTTGGTGACATCGGCCCCAAATTTGGTTATGATG AGATAGACAATGGCTACCTCAAAATGGACAACCATCGTATTCCCAGAGAAAACATGCTGATGAAGTATGC CCAGGTGAAGCCTGATGGCACATACGTGAAACCGCTGAGTAACAAGCTGACTTACGGGACCATGGTGTTT GTCAGGTCCTTCCTTGTGGGAGAAGCTGCTCGGGCTCTGTCTAAGGCGTGCACCATTGCCATCCGATACA GCGCTGTGAGGCACCAGTCTGAAATCAAGCCAGGTGAACCAGAACCACAGATTTTGGATTTTCAAACCCA GCAGTATAAACTCTTTCCACTCCTGGCCACTGCCTATGCCTTCCAGTTTGTGGGCGCATACATGAAGGAG ACCTATCACCGGATTAACGAAGGCATTGGTCAAGGGGACCTGAGTGAACTGCCTGAGCTTCATGCCCTCA CCGCTGGACTGAAGGCTTTCACCTCCTGGACTGCAAACACTGGCATTGAAGCATGTCGGATGGCTTGTGG TGGGCATGGCTATTCTCATTGCAGTGGTCTTCCAAATATTTATGTCAATTTCACCCCAAGCTGTACCTTT GAGGGAGAAAACACTGTCATGATGCTCCAGACGGCTAGGTTCCTGATGAAAAGTTATGATCAGGTGCACT CAGGAAAGTTGGTGTGTGGCATGGTGTCCTATTTGAACGACCTGCCCAGTCAGCGCATCCAGCCACAGCA GGTAGCAGTCTGGCCAACCATGGTGGATATCAACAGCCCCGAAAGCCTAACCGAAGCATATAAACTCCGT GCAGCCAGATTAGTAGAAATTGCTGCAAAAAACCTTCAAAAAGAAGTGATTCACAGAAAAAGCAAGGAGG TAGCTTGGAACCTAACTTCTGTTGACCTTGTTCGAGCAAGTGAGGCACATTGCCACTATGTGGTAGTTAA GCTCTTTTCAGAAAAACTCCTCAAAATTCAAGATAAAGCCATTCAAGCTGTCTTAAGGAGTTTATGTCTG CTGTATTCTCTGTATGGAATCAGTCAGAACGCGGGGGATTTCCTTCAGGGGAGCATCATGACAGAGCCTC AGATTACACAAGTAAACCAGCGTGTAAAGGAGTTACTCACTCTGATTCGCTCAGATGCTGTTGCTTTGGT TGATGCATTTGATTTTCAGGATGTGACACTTGGCTCTGTGCTTGGCCGCTATGATGGGAATGTGTATGAA AACTTGTTTGAGTGGGCTAAGAACTCCCCACTGAACAAAGCAGAGGTCCACGAATCTTACAAGCACCTGA AGTCACTGCAGTCCAAGCTCTGA

5' Read Nucleotide Sequence:

| 3' Read Nucleotide | >OriGene 3' read for NM_007292 unedited |
| :---: | :---: |
| Sequence: | TGCTTTGGACCGGCGGCCGCAATCTAGGGTCGAGTTTTTTTTTTTTTTTTTTTTTTTTTT |
|  | TTTTTTTTTTTTGACTTTAAAAAAAAATAATTAAGATGTATTTAACCTTTAAGTATAGAAT |
|  | GATTGGATAAATTAGATTTCAACTGAAAGGACAATTAGCGATTCAGAGAAAGCTCCTAAA |
|  | AATAGACTGAAAACAAAGCCCTACATAGAGTAAGAAATAGAACCTATGGAACTTCCATTT |
|  | TTCTAACTATAGTTTTTTGGTTTTGGTTTTTCCCAGTTGTTAAAAAAAAAACAACTCACA |
|  | GTTTCAATATTTTCTGAACCAAAAATCAGAACACATGGTTTGGGTCAAAGGGTTTTTTCTG |
|  | GGCTACGTCTGGGAGGGCTGACCTAAGGGACATTTTTTTTTAATGCCAAATACAGTAATC |
|  | TCCAAGCTTTTAATGGCTTATGCCAAGATGACAGAATATGTGAAATCTGATTGTCCCAAA |
|  | GTTACACTCTGCACTCCAAAGCTACAACAGTGCCACAGCTGAGAGGGTTCCCTATACTTC |
|  | CTACTACTGGGACAATTTAGCGATCCTTCAAATGGAAAATTCCTAATTACACGAGACAAT |
|  | GGTCCTACAGTAGGCCCGTGCGGAATAAGTTCCCTCGTTGGAAAAATGCTAGCATGAATA |
|  | GTTTGATAAATGCCATCCATCAGCAAAGGAGAGCAAGCAGGCAGAGACTGTGCCAAGCAG |
|  | AAACTCATTCGCATCAGGTTTCCAATAAGTTTCCTTTCCCCAGCCCCTTTTTTTAATCCT |
|  | GCTTTAAGCCCAGCCCCAGGGTAAGTCTGTGAAACACTGAGCAGAAAGCTTGGAAGGTTC |
|  | TAACACATTTAAAATACCCTTGCTAAAACACTTAATAGCCCGGGATAGCAATTAAATGGG |
|  | AAATATCACCCCTCCGGGGTTAAATTCTGCTATATCAAAGGCATACCTACTGGAACAN |
| Restriction Sites: | Notl-Notl |
| ACCN: | NM_007292 |
| Insert Size: | 3420 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: | 1. Centrifuge at 5,000xg for 5 min . |
|  | 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 5000 xg ) to concentrate the liquid at the bottom. |
|  | 5. Store the suspended plasmid at $-20^{\circ} \mathrm{C}$. The DNA is stable for at least one year from date of shipping when stored at $-20^{\circ} \mathrm{C}$. |
| RefSeq: | NM 007292.3 NP 009223.2 |
| RefSeq Size: | 3445 bp |
| RefSeq ORF: | 1983 bp |
| Locus ID: | 51 |
| UniProt ID: | Q15067 |
| Cytogenetics: | 17q25.1 |
| Protein Families: | Druggable Genome |

Protein Pathways:

Gene Summary:
alpha-Linolenic acid metabolism, Biosynthesis of unsaturated fatty acids, Fatty acid metabolism, Metabolic pathways, PPAR signaling pathway
The protein encoded by this gene is the first enzyme of the fatty acid beta-oxidation pathway, which catalyzes the desaturation of acyl-CoAs to 2-trans-enoyl-CoAs. It donates electrons directly to molecular oxygen, thereby producing hydrogen peroxide. Defects in this gene result in pseudoneonatal adrenoleukodystrophy, a disease that is characterized by accumulation of very long chain fatty acids. Alternatively spliced transcript variants encoding different isoforms have been identified. [provided by RefSeq, Jul 2008] Transcript Variant: This variant (2) has multiple differences in the coding region but maintains the reading frame, compared to variant 1. The encoded isoform (b, alternately known as ACOX1a per PMID: 17603022), is the same length as isoform a. 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.

