

Product datasheet for **SC111756**

ALDH1L1 (NM_012190) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	ALDH1L1 (NM_012190) Human Untagged Clone
Tag:	Tag Free
Symbol:	ALDH1L1
Synonyms:	10-fTHF; 10-FTHFDH; FDH; FTHFD
Mammalian Cell Selection:	None
Vector:	<u>pCMV6-XL4</u>
E. coli Selection:	Ampicillin (100 ug/mL)



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Fully Sequenced ORF: >OriGene ORF within SC111756 sequence for NM_012190 edited (data generated by NextGen Sequencing)

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ATGAAGATTGCAGTGATTGGACAGAGCCTGTTTGGCCAGGAAGTTTACTGCCACCTGAGG
AAGGAGGGCCACGAAGTGGTGGGTGTGTTCACTGTTCCAGACAAGGATGGAAAGGCCGAC
CCCCTGGGTCTGGAAGCTGAGAAGGATGGAGTGCCGGTATTCAAGTACTCCCGTGGCGT
GCAAAAAGGACAGGCTTTGCCTGATGTGGTGGCAAATAACCAGGCTTTGGGGCCGAGCTC
AACGTCCTGCCCTTCTGCAGCCAATTCATCCCCATGGAGATAATCAGTGCCCCCGGCAT
GGCTCCATCATCTATCACCCGTCCTGCTCCCTAGGCACCGAGGGCCTCGGCCATCAAC
TGGACCCTCATTACGGAGATAAGAAAAGGGGGTTTTCCATCTTCTGGGCGGATGATGGT
CTGGACACCGGAGACCTGCTGCTGCAGAAGGAGTGTGAGGTGCTCCCGGACGACACCGTG
AGCACGCTGTACAACCGCTTCTCTTCCCTGAAGGCATCAAAGGGATGGTGCAGGCCGTG
AGGCTGATCGCTGAGGGCAAAGCCCCAGACTCCCTCAGCCTGAGGAAGGAGCCACCTAT
GAGGGGATTCAGAAGAAGGAGACAGCCAAGATCAACTGGGACCAGCCGGCAGAGGCCATT
CACAACTGGATCCGCGGGAACGACAAGGTGCCGGGAGCCTGGACAGAGGCCCTGTGAACAG
AAACTGACATTTTCAACTCAACGCTGAACACTTCAGGCCTGGTCCCGAGGGAGACGCT
TTGCCATCCCAGGAGCCATCGGCCAGGGGTGGTCAACAAAGCAGGACTCATCCTCTTT
GGGAATGATGACAAAATGCTGCTGGTGAAGAATATTCAGCTGGAGGATGGCAAAATGATC
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CTGGTTACTGCGGAGGCTGTGCGGAGTTTTTGGCAGCGGATCCTCCCAAAGTCTGGAG
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ACCATCAATCCCACCGATGGAAGTGTCTGTCAGGTATCCCTGGCCAAAGTCAACCGAC
GTCGACAAGGCAGTGGCCGACGCAAGGATGCCTTTGAGAATGGACGGTGGGGGAAGATC
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AGGGACATCAACAAGGCCCTGTATGTGAGTGAAGCTCCAGGCAGGCACTGTGTTTGTG
AACACGTACAACAAGGCCGCTGATGTGAGTGAAGCTCCAGGCAGGCACTGTGTTTGTG
GGCAAAGATCTAGGAGAGCGGCTCTGAACGAGTACCTGCGGGTCAAGACAGTGACCTTC
GAATACTGA

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Clone variation with respect to NM_012190.2
 988 g=>t;1185 g=>t;1192 g=>c;1441 a=>g

5' Read Nucleotide Sequence:

>OriGene 5' read for NM_012190 unedited
 TTCGGNCCTCAGCTGCGGCTCATTTAAGTCTAACCTGCCCGGACGCCCGACCAACTCC
 TGCGCGGGTCTTCCAACCTCCTGCTACCATGAAGATGCAGTGATTGGACAACCTGTT
 TGGCCAGAATTTACTGCCACCTGAGAAGAGGCCACGAATGTNNGGTAGNTCACTGTTCCA
 TACAAGGATTGAAAGGCCGACCCCTGGGTCTGGAAGCTGAGAAGGATGGAGTGCCGGTA
 TTCAAGTACTCCCTGTGGCGTGCAAAAGGACAGGCTTTGCCTGATGTGGTGGCAAATAC
 CAGGCTTTGGGGCCGAGCTCAACGTCCCTGCCCTTCTGCAGCCAATTCATCCCCATGGAG
 ATAATCAGTGCCCCCGGCATGGCTCCATCATCTATCACCCGTCAGTCTCCCTAGGCAC
 CGAGGGGCTCGGCCATCAACTGGACCCTCATTACGGAGATAAGAAAGGGGGTTTTTC
 ATCTTCTGGGCGGATGATGGTCTGGACACCGGAGACCTGCTGCTGCAGAAAGAGTGTGAG
 GTGCTCCCGGACGACACCGTGAGCAGCTGTACAACCGTTCTCTTCCCTGAGGCATCA
 AAGGGATGGTGCAGGCCGTGAGGCTGATCCCTGANGCAAAACCCCAAACCTCCCTCACCC
 TGAGGAAGGAACCTATGAGGGGATTCAGAAAAGGAGAAGCCAGAATCACTTGGGAC
 CAGCCCGCAGAGCCATTTCAACTTGGATCCGGGGAAACAAACAAGTCCCGGGAGCCT
 GGGCAAAAGGCTGTTGAACAGAACTGGAATTNTTTCAACTTAACGCCTTGAACATTCA
 AGGCCTGGGCCCGAGGAAACCTTTGCCATCCCAGGAACCCCTCCGCCCCAGGGTG
 GTC

3' Read Nucleotide Sequence:

>OriGene 3' read for NM_012190 unedited
 ACTCTGGACCGCGCNCGAATCTANGATNCGTTTTTTTTTTTTTTTTTTTTTTTTTTTCT
 CTTATAAGCTTTATTCTCCCTGGGAGGGGCACACCTCACCCAGCAAGGGCTGCTTCTG
 ACTGGACAGAGGGCGTCCAAAATGCAAACATGGTGTGCAGGCAGGAGGGCTTCCACTAGC
 CCCCCAGGTGGGAGGTGCTGTGCACCCAGGCTCAAGAGGGAGGGGCCCCAGCCACGAGG
 GAGGGGCAGGGACTTTCTTCTCACAAGACCTTTCTTCAAGTATTGAAAGGTCAGTGTCTT
 GACCCGACAGTACTCGTTCAAAGCCGCCTCTCCTAAATCTTTGCCAAATCCAGACTGTTT
 GAATCCTCCGAAGGGAGCGGCCACGTCCGGTCTTGTGTACGTGTTGACAAACACAGTGCC
 TGCCTGGAGCTTGTCACTGACATACAGGGCCTTGTGATGTCCTGGTGAAGACACCAGA
 AGCCAGGCCAAATTCGTGGCATTGGCCCGAGACAGCACGGCATCCAAGTCCCCATCAGC
 AAACCGAGAGATGATCATGACAGGCCCGAAGGACTCCTCCTTGGCTATGAACATGTGGTC
 TTCCACGTCTGTGAAAACAGTTGGCTCAAAGAAGAACCCTGGCCGAGGGACCTGATTCCT
 GCCGACAGACAGTGTGGCCCTTCTTACGCCATGCTGGCAGTACTCCATCAGCTTCAC
 AAGGTGGGCATGGTATTCTGCGGCCCGTGGTGGTGTCCCTGTCCAGCGGGGTGCCCA
 CCTTATCTTCCGCACCTTNTACCACTCTNCGCACGAACNATCATGAATGGAGTCTNCG
 ACAAGAGTGCCTGCTGCATGCAATCTCTCTTTGTTGAAGAAACAGACTCATCCCATC
 TGCACANCTTGGTGAAGTACAGTACAGCAAAATGATGAGGGTGACTTCCGCCCCAGTCAGG
 GACTTCTNACGACTT

Restriction Sites:

NotI-NotI

ACCN:

NM_012190

Insert Size:

3100 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_012190.2 , NP_036322.2
RefSeq Size:	3125 bp
RefSeq ORF:	2709 bp
Locus ID:	10840
UniProt ID:	O75891
Cytogenetics:	3q21.3
Domains:	aldehyd, formyl_transf, formyl_trans_C
Protein Families:	Druggable Genome
Protein Pathways:	One carbon pool by folate
Gene Summary:	<p>The protein encoded by this gene catalyzes the conversion of 10-formyltetrahydrofolate, nicotinamide adenine dinucleotide phosphate (NADP+), and water to tetrahydrofolate, NADPH, and carbon dioxide. The encoded protein belongs to the aldehyde dehydrogenase family. Loss of function or expression of this gene is associated with decreased apoptosis, increased cell motility, and cancer progression. There is an antisense transcript that overlaps on the opposite strand with this gene locus. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Jun 2012]</p> <p>Transcript Variant: This variant (2) contains a distinct 5' UTR, lacks an in-frame portion of the 5' coding region, and initiates translation at an alternate in-frame downstream start codon, compared to variant 1. The resulting isoform (2) has a shorter N-terminus, compared to 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>