

## Product datasheet for **SC216316**

### ALDH7A1 (NM\_001182) Human 3' UTR Clone

#### Product data:

Product Type:	3' UTR Clones
Product Name:	ALDH7A1 (NM_001182) Human 3' UTR Clone
Vector:	pMirTarget (PS100062)
Symbol:	ALDH7A1
Synonyms:	ATQ1; EPD; PDE
ACCN:	NM_001182
Insert Size:	2000 bp



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**Insert Sequence:**

>SC216316 3'UTR clone of NM\_001182

The sequence shown below is from the reference sequence of NM\_001182. The complete sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

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GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAGCGATCGCC
CCTCTGGCCCAAGGAATCAAGTTTCAGTAAGGTGTTTTAGATGAACATCCCTTAATTTGAGGTGTTC
AGCAGCTGTTTTGGAGAAGACAAAGAAAATTAAGTTTTCCCTGAATAAATGCATTATTATGACTGTG
ACAGTGACTAATCCCCTATGACCCCAAAGCCCTGATTAATCAAGAGATTCTTTTTAAAAATCAAA
ATAAAATTGTTACAACATAGCCATAGTTACTAAAAGATGAGTTAGGTGGATTTTTATTATGTCAACTAG
TTGTACATGGCTTTTTAAAAGTTAATGATTATTTGTAATTAGAAAAAATAGTACGTACCAGGGTAGG
AATTTGGGAAAATACAGAACCGAGGAGAAAACAGAAGTCTTTCAGTAGATACAGGGTGTCTCTGACC
AAGTGAAGGATTCAGGGCGGGGGTGAATATTGCTTGACATTACCCACCTATGGCATGTGTTGGATGT
CGTGTGATTGAAAGGGGATGCATTCCTGTATGCTTGACCACAGTTCAGTATTCATGTGAATAAC
TTGCAGTGCTTAAATGACAACAGGCATTCATCACAGTTATGGCTCTTGCTTATGAAGGCTGTGTCAAC
TTGGAAGAGATTTCTGGGTAGACAGATTTTGTCTGTGCTGGAGTCTGCCACCTGCTGGAGGACTCT
GAGGCCAAAATTGCCCTGCAGATTGTATAGAAAATGCGTCTGGAGGCTGGGCAAGGTGGCTTACGCC
TATAATCCAGCACTTTGAGAGGCCGAGGCGGGTGGATCAGTAGAGGCTAGGAGTTTGGAGCCAGCCTG
GCCGATATCGTGAACCCCGTCTTTACCAAAAAATACAAAAATAGCCAGTGTGGTGGCAGGCACCTG
TAATCCAGCTATTAGGGAGGCTGAGGCAGGAGAATTGTGTGAACCCAGGAGGCAGAGTTGCAGTGAG
CCTAGATTGTGCCACTGCCTGGGCAACAGTGAATCCTGTCTACAAAAAAGTGA
TCTGGATTTTTTATACAACCTTAGACCCTTTAGCTTTAGCGTCTGCGGTTGCCCTGGATCT
GTTCTCAATCCTCAGTGTGTGGCAGCATGTGGTCATAGAGAGCTGGGCAAGTTCACCTTCTCTTTG
CTGACAGTCTCACCTTTCTCACTGGGAAGCTGCACAGGAGCCTTTGGGCTGGTTCAGCCCAGAGGCC
CTGTTCTCCTGCCTTCTGTCATTCTCTGCTCCCTTCTGCATGGCACCCCTCTACTCTGCCAGGTTA
GAATGGAGCAGAGGCCTTTGAACAAGATGACATTTGTAGACAACCTGATCCATCAGGATGTGGAGAGAT
CTGGACAAATGGCAAATAATTTGGGTTGAATTATGGCCAGAGAAGTAAGCAAATGAAAACTAAACA
ATTATAATGTATCTGGGAAGCAAAGTATTATAATATGTCCCATGGCCAGCTGCGAGCAGTATTTAA
CTAGACATGCTGTGCAATATTGAACATGAATCAAGCCAAATACTGATATAGCTATTGAGAGAAGTT
GGCAGGAAGAAATCATGGGTGAGGGTATGTGGGAGTGATAATCTTAGCTATAGATTTTAGTAAAAGT
ATTCTGAAAGCTTTCTTTGAAACAAATTTATAATTTTCTTCTACATGTTTCTAGACAAGGACTCAAT
TTTTTTGGTTAATGTTCAATAAAACATATGCCTTCAATTTGTTCTCTTTTGTCTTTTTTGGCTGAT
TTAGCAGATATAAATCACTTGTTAGTATTTTAACTATCCAAGTAAAATGTACATTTGTAAGTGTAT
TACTGGTCTTATATTTGAAGGTGCAAGGGTCTTGTGCTGCTCATTACTGGTTATCTTGAGCATTTCAT
AACTGCTGCCTCCTCAATCCTGTGTTCCAACCGCTACTGCTGGCAATGCCAGCCTCCATTGTGG
ACGCGTAAGCGGCCGCGGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCAACCTGCCATCA
CGAGATTCGATTCACCGCCGCTTCTATGAAAGG
    
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**Restriction Sites:**

Sgfl-Mlul

**OTI Disclaimer:**

Our molecular clone sequence data has been matched to the sequence identifier above as a point of reference. Note that the complete sequence of this clone is largely the same as the reference sequence but may contain minor differences, e.g., single nucleotide polymorphisms (SNPs).

**Components:**

The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The package also includes 100 pmols of both the corresponding 5' and 3' vector primers in separate vials.

**RefSeq:**

[NM\\_001182.5](#)

**Summary:** The protein encoded by this gene is a member of subfamily 7 in the aldehyde dehydrogenase gene family. These enzymes are thought to play a major role in the detoxification of aldehydes generated by alcohol metabolism and lipid peroxidation. This particular member has homology to a previously described protein from the green garden pea, the 26g pea turgor protein. It is also involved in lysine catabolism that is known to occur in the mitochondrial matrix. Recent reports show that this protein is found both in the cytosol and the mitochondria, and the two forms likely arise from the use of alternative translation initiation sites. An additional variant encoding a different isoform has also been found for this gene. Mutations in this gene are associated with pyridoxine-dependent epilepsy. Several related pseudogenes have also been identified. [provided by RefSeq, Jan 2011]

**Locus ID:** 501

**MW:** 76.5