

Product datasheet for SC326810

LDHA (NM 001165416) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: LDHA (NM 001165416) Human Untagged Clone

Tag: Tag Free Symbol: LDHA

Synonyms: GSD11; HEL-S-133P; LDHM; PIG19

Mammalian Cell

Selection:

None

Vector: pCMV6-XL5

E. coli Selection: Ampicillin (100 ug/mL)

Fully Sequenced ORF: >NCBI ORF sequence for NM_001165416, the custom clone sequence may differ by one or

more nucleotides

ATGGCAACTCTAAAGGATCAGCTGATTTATAATCTTCTAAAGGAAGAACAGACCCCCCAG
AATAAGATTACAGTTGTTGGGGTTGGTGCTGTTGGCATGGCCTGTGCCATCAGTATCTTA
ATGAAGGACTTGGCAGATGAACTTGCTCTTGTTGATGTCATCGAAGACAAATTGAAGGGA
GAGATGATGGATCTCCAACATGGCAGCCTTTTCCTTAGAACACCCAAAGATTGTCTCTGGC
AAAGACTATAATGTAACTGCAAACTCCAAGCTGGTCATTATCACGGCTGGGGCACGTCAG
CAAGAGGGAGAAAGCCGTCTTAATTTGGTCCAGCGTAACGTGAACATCTTTAAATTCATC
ATTCCTAATGTTGTAAAATACAGCCCGAACTGCAAGTTGCTTATTGTTTCAAATCCAGTG
GATATCTTGACCTACGTGGCTTGGAAGATAAGTGGTTTTCCCAAAAACCGTGTTATTGGA
AGCGGTTGCAATCTGGATTCAGCCCGATTCCGTTACCTAATGGGGGAAAGGCTGGGAGTT
CACCCATTAAGCTGTCATGGGTGGTCCTTGGGGAACATCTGCACCCAGATTTAGGGACT
GGAGTGGAATGAATGTTGCTGGTGTCTCTCTGAAGACTCTGCACCCAGATTTAGGGACT
GATAAAGATAAGGAACAGTGGAAAGAGGTTCACAAGCAGGTGTTGAGAGGGTCTTTACG

 GAA

Restriction Sites: Please inquire **ACCN:** NM 001165416

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).



OriGene Technologies, Inc. 9620 Medical Center Drive, Ste 200

CN: techsupport@origene.cn

Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com

LDHA (NM_001165416) Human Untagged Clone - SC326810

OTI Annotation: This TrueClone is provided through our Custom Cloning Process that includes sub-cloning

into OriGene's pCMV6 vector and full sequencing to provide a non-variant match to the expected reference without frameshifts, and is delivered as lyophilized plasmid DNA.

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 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>NM 001165416.1, NP 001158888.1</u>

 RefSeq Size:
 2102 bp

 RefSeq ORF:
 726 bp

 Locus ID:
 3939

 UniProt ID:
 P00338

 Cytogenetics:
 11p15.1

Protein Families: Druggable Genome

Protein Pathways: Cysteine and methionine metabolism, Glycolysis / Gluconeogenesis, Metabolic pathways,

Propanoate metabolism, Pyruvate metabolism

Gene Summary: The protein encoded by this gene catalyzes the conversion of L-lactate and NAD to pyruvate

and NADH in the final step of anaerobic glycolysis. The protein is found predominantly in muscle tissue and belongs to the lactate dehydrogenase family. Mutations in this gene have been linked to exertional myoglobinuria. Multiple transcript variants encoding different isoforms have been found for this gene. The human genome contains several non-

transcribed pseudogenes of this gene. [provided by RefSeq, Sep 2008]

Transcript Variant: This variant (5) lacks an alternate exon in the 3' coding region, compared to variant 1, which results in a frameshift. The resulting isoform (5) lacks a segment of the LDH domain and has a shorter and distinct C-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.