

## Product datasheet for SC326928

### LDHA (NM\_001165414) Human Untagged Clone

#### Product data:

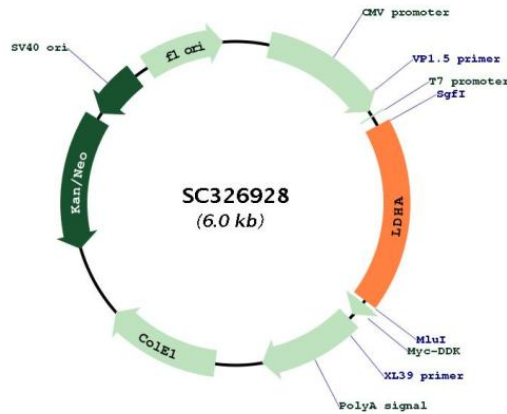
Product Type:	Expression Plasmids
Product Name:	LDHA (NM_001165414) Human Untagged Clone
Tag:	Tag Free
Symbol:	LDHA
Synonyms:	GSD11; HEL-S-133P; LDHM; PIG19
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Fully Sequenced ORF:	>SC326928 representing NM_001165414. Blue=Insert sequence Red=Cloning site Green=Tag(s)

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GCTCGTTTAGTGAACCGTCAGAATTTGTAAACGACTACTATAGGGCGCCGGGAATTCGTCGACTG
GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC
ATGGGTGAACCTCAGGAGGCTATACTTACACCCAAACGTCGATATTCCTTTCCACGCTAAGATTCCT
TTTGGTCCAAGTCCAATATGGCAACTCTAAAGGATCAGCTGATTTATAATCTTCTAAAGGAAGAACAG
ACCCCCAGAATAAGATTACAGTTGTTGGGTTGGTGCTGTTGGCATGGCCTGTGCCATCAGTATCTTA
ATGAAGGACTTGGCAGATGAACCTGCTCTTGTGATGTCATCGAAGACAAATTGAAGGGAGAGATGATG
GATCTCCAACATGGCAGCCTTTCTTAGAACACCAAAGATTGTCTCTGGCAAAGACTATAATGTAAC
GCAAACCTCAAGCTGGTCATTATCACGGCTGGGGCACGTCAGCAAGAGGGAGAAAGCCGCTTAATTTG
GTCCAGCGTAACGTGAACATCTTTAAATTCATCATTCTAATGTTGTAATAACAGCCCGAAGTCAAG
TTGCTTATTGTTTCAAATCCAGTGGATATCTTGACCTACGTGGCTTGAAGATAAGTGGTTTTCCAAA
AACCGTGTATTGGAAGCGGTTGCAATCTGGATTACGCCGATTCCGTTACCTAATGGGGGAAAGGCTG
GGAGTTCACCCATTAAGCTGTCTGGTGGTCTTGGGGAACATGGAGATTCCAGTGTGCCGTGATGG
AGTGAATGAATGTTGCTGGTGTCTCTGAAGACTCTGCACCCAGATTTAGGGACTGATAAAGATAAG
GAACAGTGGAAAGAGGTTCAAGCAGGTGGTTGAGAGTGCTTATGAGGTGATCAAACCTCAAAGGCTAC
ACATCTGGGCTATTGGACTCTGTAGCAGATTTGGCAGAGAGTATAATGAAGAATCTTAGGCGGGTG
CACCCGTTTCCACCATGATTAAGGGTCTTACGGAATAAAGGATGATGTCTTCTTAGTGTTCCTTGC
ATTTTGGGACAGAATGGAATCTCAGACCTTGTGAAGGTGACTCTGACTTCTGAGGAAGAGGCCGTTTG
AAGAAGAGTGCAGATACACTTTGGGGGATCCAAAAGGAGCTGCAATTTTAA
ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAATGATATCCTGGAT
TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC
```

Restriction Sites: SgfI-MluI



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**Plasmid Map:**


**ACCN:** NM\_001165414

**Insert Size:** 1086 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).

**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:** [NM\\_001165414.1](#)

**RefSeq Size:** 2323 bp

**RefSeq ORF:** 1086 bp

**Locus ID:** 3939

UniProt ID:	<a href="#">P00338</a>
Cytogenetics:	11p15.1
Protein Families:	Druggable Genome
Protein Pathways:	Cysteine and methionine metabolism, Glycolysis / Gluconeogenesis, Metabolic pathways, Propanoate metabolism, Pyruvate metabolism
MW:	39.8 kDa
Gene Summary:	<p>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]</p> <p>Transcript Variant: This variant (3) uses an alternate 5' exon that results in a distinct 5' UTR and causes translation initiation at an alternate start codon, compared to variant 1. The resulting isoform (3) has a longer 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>