

Nucleotide and Protein sequences

Nucleotide sequences.

NCBI

All Databases PubMed Nucleotide Protein Genome Structure OMIM PMC Journals Books

Search Nucleotide for plasmodium falciparum lactate dehydrogenase Go Clear Save Search

Limits Preview/Index History Clipboard Details

Display Summary Show 20 Sort By Send to

All: 46 Bacteria: 23 INSDC (GenBank): 29 RefSeq: 17 mRNA: 1

This search in Gene shows [3 results](#), including:

[PF13_0141](#) (*Plasmodium falciparum* 3D7): L-lactate dehydrogenase

[PF13_0144](#) (*Plasmodium falciparum* 3D7): oxidoreductase

[PFF0895w](#) (*Plasmodium falciparum* 3D7): malate dehydrogenase

Items 41 - 46 of 46 Previous Page 3 of 3

41. [A kit for diagnosing malaria comprising monoclonal antibodies to malarial lactate dehydrogenase and aldolase](#)
1,084 bp linear genomic
DI122512.1 GI:168421398

42. [Plasmodium falciparum isolate FCC1/HN lactate dehydrogenase \(LDH\) gene, complete cds](#)
951 bp linear genomic

Top Organisms [Tree]

Plasmodium falciparum (8)
Toxoplasma gondii (5)
Arabidopsis thaliana (4)
Toxoplasma gondii ME49 (4)
Mycobacterium bovis AF2122/97 (3)
All other taxa (26)

More...

Recent activity

Plasmodium falciparum L-lactate dehydrogenase (LDH-P) mRNA

Internet Protected Mode Off

A kit for diagnosing malaria comprising monoclonal antibodies to malarial lactate dehydrogenase and aldolase

41. 1,084 bp linear genomic

DI122512.1 GI:168421398

Plasmodium falciparum isolate FCC1/HN lactate dehydrogenase (LDH) gene, complete cds

42. 951 bp linear genomic

DQ825436.1 GI:111034850

Select this one

Plasmodium falciparum isolate FCBR L-lactate dehydrogenase (LDH) gene, complete cds

43. 951 bp linear genomic

DQ198262.1 GI:76563842

Plasmodium falciparum isolate K1 L-lactate dehydrogenase (LDH) gene, complete cds

44. 951 bp linear genomic

DQ198261.1 GI:76563840

Toxoplasma gondii malate dehydrogenase gene, complete cds

45. 951 bp linear genomic

AY972525.1 GI:62465590

Plasmodium reichenowi ldh gene for lactate dehydrogenase, complete cds

46. 951 bp linear genomic



Encore Nucleotide

My NCBI
[Sign In] [Register]

All Databases

PubMed

Nucleotide

Protein

Genome

Structure

PMC

Taxonomy

Books

Search Nucleotide for

Go Clear

Limits

Preview/Index

History

Clipboard

Details

Download

Save

Links

Format: GenBank FASTA Graphics More Formats

GenBank DQ198262.1

Plasmodium falciparum isolate FCBR L-lactate dehydrogenase (LDH) gene, complete cds

[Features](#) [Sequence](#)

LOCUS DQ198262 951 bp DNA linear INV 04-OCT-2005
 DEFINITION Plasmodium falciparum isolate FCBR L-lactate dehydrogenase (LDH) gene, complete cds.
 ACCESSION DQ198262
 VERSION DQ198262.1 GI:76563842
 KEYWORDS .
 SOURCE Plasmodium falciparum (malaria parasite P. falciparum)
 ORGANISM [Plasmodium falciparum](#)
 Eukaryota; Alveolata; Apicomplexa; Aconoidasida; Haemosporida; Plasmodium; Plasmodium (Laverania).
 REFERENCE 1 (bases 1 to 951)
 AUTHORS Turgut-Balik,D. and Holbrook,J.J.
 TITLE Determination of the DNA and aminoacid sequences of lactate dehydrogenase gene from Plasmodium falciparum strains K1 and RF FCBR: A route to the design of new antimalarials
 JOURNAL Turk. J. Biol. 25, 241-250 (2001)
 REFERENCE 2 (bases 1 to 951)
 AUTHORS Turgut-Balik,D., Shoemark,D.K., Moreton,K.M., Sessions,R.B. and Holbrook,J.J.
 TITLE Over-production of lactate dehydrogenase from Plasmodium falciparum opens a route to new antimalarials
 JOURNAL Biotechnol. Lett. 23, 917-921 (2001)
 REFERENCE 3 (bases 1 to 951)

Header

Change Region Shown

Customize View

Pick Primers

Design and test primers for this sequence using Primer-BLAST.

Recent Activity

All links from this record

Date when record was made public

Gene index

Accession number



Internet

start

2 Windo...

2 Micros...

2 Novell ...

2 Micros...

2 Intern...

2 Nucleotid...

Type to search



14:06



JOURNAL Submitted (08-SEP-2005) Biology, Faculty of Arts and Sciences,
Elazig 23169, Turkey

FEATURES

source Location/Qualifiers
1..951
/organism="Plasmodium falciparum"
/mol_type="genomic DNA"
/isolate="FCBR"
/db_xref="taxon:[5833](#)"

gene <1..>951
/gene="LDH"

mRNA <1..>951
/gene="LDH"
/product="L-lactate dehydrogenase"

CDS 1..951
/gene="LDH"
/EC_number="[1.1.1.27](#)"
/codon_start=1
/product="L-lactate dehydrogenase"
/protein_id="[ABA46355.1](#)"
/db_xref="GI:76563843"
/translation="MAPKAKIVLVGSGMIGGVMATLIVQKNLGDVVLFDIVKNMPHGK
ALDTSHTNVMAYSNCVGSNTYDLAGDVVITAGFTKAPGKSDKEWNRDLLPLN
NKIMIEIGHHIKKNCPNAFIIVVTNPVDVMVQLLHQHSGVPKNIIGLGGVLDTSLRK
YYISQKLNVCPRDVNAHIVGAHGNKMVLKRYITVGGIPLQEFINNKLISDAELEAIF
DRTVNTALEIVNLHASPYVAPAAIIEMAESYKLKLVLCSTLLEGQYGHSDIFGG
TPVVLGANGVEQVIELQLNSEEKAKFDEAIAETKRMKALA"

Coding Sequence

Features

Sequence

ORIGIN

1 atggcaccaa aagaaaaat cgtttagtt ggctcaggta tgattggagg agtaatggct
61 acctaatttgc ttcagaaaaa tttaggagat gtatgggt tcgtatattgt aaagaacatg
121 ccacatggaa aagctttaga tacatctcat actaatgtta tgccatattc aaattgcaaa
181 gtaagtggtt caaacactta tgacgatttg gctggagcag atgttagata agtaacagct
241 ggatttacca aggccccagg aaagagtgc aaagaatggg atagagatga ttatttacca
301 ttaaacaaca agattatgtat tgaaattgggt ggtcatatttta agaagaatttgc tccaaatgt
361 ttatttatttgc ttgtacaaa cccagtagat gttatggta aatttattaca tcaacattca
421 ggtgttccat aaaaacaaat tattgggtta ggtgttgtat tagatacatc aagatttgc
481 tattacatat ctccaaaaattt aatgtatgc ccaagagatgc taaatgcaca cattgttaggt
541 gctcatggaa ataaaaatgtt tctttttaaa agatcatttgc ctgttaggtgg tatcccttta
601 caagaatttgc ttaaataacaa gttaaatttgc gatgtgtat tagaagctat atttgataga
661 actgttataa ctgcatttgc aattgttacac ttacatgttgc caccatgttgc accgttgc
721 gctgttattttgc tgcggatggc tgaatcttgc tttaaaaggatt tgaaaaaaatg attaatttgc
781 tcaacccgttgc tagaaggacac atatggacac tccgttatatttgc tccgttgc acctgttgc
841 ttaggtgttgc atgggtttgc acaagtttgc gaatttacaat taaatgttgc ggaaaaaaatg
901 aaatttgc aagccatagc tggaaactaag agaatttgc cattatgttgc a

//

DQ198262: ***Plasmodium falciparum*** LDH sequence.

- There are several IMPORTANT features given to you on the sequence page.
- HEADER
 - **DQ198262** is a UNIQUE number assigned to the Pf LDH gene sequence in a database.
 - Known as the: ACCESSION number (can have various versions: DQ198262.1)
 - **gi:76563842:** gene index identifies the current version
 - If sequence annotation is revised then the gi number changes but the accession number remains the same
 - LOCUS: Length of the sequence: **951bp**
 - DEFINITION: name of the sequence
 - SOURCE: Common name of *Plasmodium falciparum* (i.e. malaria parasite *P. falciparum*)
 - ORGANISM *Plasmodium falciparum*
 - More complete identification
 - Lineage: Eukaryota; Alveolata; Apicomplexa; Haemosporida; Plasmodium.
 - The hyperlink takes you to the...**TAXONOMY** database

- REFERENCE: gives publication citation related to the sequence entry
 - AUTHORS: Turgut-Balik,D., Shoemark,D.K., Moreton,K.M., Sessions,R.B. and Holbrook,J.J.
 - TITLE Over-production of lactate dehydrogenase from Plasmodium falciparum opens a route to new antimalarials. JOURNAL Biotechnol. Lett. 23, 917-921 (2001).

- **FEATURES**

- Information about gene, gene product and regions of biological significance
 - Coding sequence (CDS), mRNA, introns, exons
 - Enzyme Committee number. **E.C.**
 - Unique international identifier (eg: [1.1.1.27](#))
 - Click on link to go to EXPASY website
- Amino acid sequence number : [ABA46355.1](#)
 - Links to **Entrez Protein** database
 - The amino-acid sequence is given
- **SEQUENCE**
 - The DNA or base sequence is given. (Note: what is the base sequence of the first codon?)
 - **CDS (coding sequence)** starts with ATG (DNA) or AUG if mRNA

NiceZyme View of ENZYME: EC 1.1.1.27

Official Name

L-lactate dehydrogenase.

Alternative Name(s)

L-lactic acid dehydrogenase.

L-lactic dehydrogenase.

Reaction catalysed**Comment(s)**

- Also oxidizes other (S)-2-hydroxymonocarboxylic acids.
- NADP(+) acts, more slowly, with the animal, but not the bacterial, enzyme.

Cross-references

Biochemical Pathways; map number(s)	E6 ; H4
PROSITE	PDOC00062
BRENDA	1.1.1.27
PUMA2	1.1.1.27
PRIAM enzyme-specific profiles	1.1.1.27
KEGG Ligand Database for Enzyme Nomenclature	1.1.1.27
IUBMB Enzyme Nomenclature	1.1.1.27
IntEnz	1.1.1.27
MEDLINE	Find literature relating to 1.1.1.27
MetaCyc	1.1.1.27

Q81RW4, LDH1_BACAN; P62047, LDH1_BACC1; Q81EP4, LDH1_BACCR;
Q63CN1, LDH1_BACCZ; Q6HK31, LDH1_BACHK; P59050, LDH1_BIFLO;
Q97MD1, LDH1_CLOAB; Q839C1, LDH1_ENTFA; Q5FMBO, LDH1_LACAC;

Features (Nucleotide database)

- Coding sequence (CDS)
- Ribosomal binding sites (RBS)
- Start/stop
- Promotor sequences
- Introns/exons
- Repeat regions etc.
- Remember:
 - Promotor – shows the position of the promoter box e.g. -35 for transcription
 - RBS is the ribosomal binding site
 - CDS stands for coding segment/sequence (i.e. open reading frame: ORF).

[promoter](#)

```
/mol_type="genomic DNA"
/db_xref="taxon:562"
286..291
/note="-35 region"
310..316
/note="-10 region"
322..324
/note="put. transcription start region"
```

Transcription[promoter](#)

310..316

(Note: -10 region)[misc_feature](#)

322..324

(Note: put. transcription start region)

330..333

(Note: put. rRNA binding site)[RBS](#)

343..798

(Note: unnamed protein product; dUTP-ase (aa 1-151))[CDS](#)*/codon_start=1**/transl_table=11**/protein_id="CAA25859.1"**/db_xref="GI:41297"**/db_xref="GOA:P06968"**/db_xref="InterPro:IPR008180"**/db_xref="InterPro:IPR008181"**/db_xref="PDB:1DUD"**/db_xref="PDB:1 DUP"**/db_xref="PDB:1EUS"**/db_xref="PDB:1EUW"**/db_xref="PDB:1RN8"**/db_xref="PDB:1RNJ"**/db_xref="PDB:1SEH"**/db_xref="PDB:1SYL"**/db_xref="UniProtKB/Swiss-Prot:P06968"*

translation="MKKIDVKILDPRVGKEFPLPTYATSGSAGLDLRACLNDAVELAP
 GTTTLVPTGLAIHIADPSLAAMMLPRSGLGHKGIVLGNLVGLIDSYQGQLMISVWN
 RGQDSFTIOPGERIAQMIFVPVVQAEFNLVEDFDATDRGEGGFHSGRQ"
 831..851

[misc_feature](#)*/note="put. stem-loop structure"*[repeat_unit](#)

831..838

/note="inverted repeat A"[repeat_unit](#)

844..851

/note="inverted repeat A'"[misc_feature](#)

866..893

/note="put. stem-loop structure"[repeat_unit](#)

866..872

/note="imp. inverted repeat B"[repeat_unit](#)

888..893

*.....***Shine Dalgarno (Translation)****dUTPase**

File Edit View Favorites Tools Help

Search Nucleotide for Go Clear

Limits Preview/Index History Clipboard Details

Format: GenBank [FASTA](#) [Graphics](#) More Formats

Download Save Links

GenBank DQ198262.1

Plasmodium falciparum isolate FCBR L-lactate dehydrogenase (LDH) gene, complete cds

[Features](#) [Sequence](#)

LOCUS	DQ198262	951 bp	DNA	linear	INV 04-OCT-2005
DEFINITION	Plasmodium falciparum isolate FCBR L-lactate dehydrogenase (LDH) gene, complete cds.				
ACCESSION	DQ198262				
VERSION	DQ198262.1 GI:76563842				
KEYWORDS	.				
SOURCE	Plasmodium falciparum (malaria parasite P. falciparum)				
ORGANISM	Plasmodium falciparum Eukaryota; Alveolata; Apicomplexa; Aconoidasida; Haemosporida; Plasmodium; Plasmodium (Laverania).				
REFERENCE	1 (bases 1 to 951)				
AUTHORS	Turgut-Balik,D. and Holbrook,J.J.				
TITLE	Determination of the DNA and aminoacid sequences of lactate dehydrogenase gene from Plasmodium falciparum strains K1 and PF FCBR: A route to the design of new antimalarials				
JOURNAL	Turk. J. Biol. 25, 241-250 (2001)				
REFERENCE	2 (bases 1 to 951)				
AUTHORS	Turgut-Balik,D., Shoemark,D.K., Moreton,K.M., Sessions,R.B. and Holbrook,J.J.				
TITLE	Over-production of lactate dehydrogenase from Plasmodium falciparum opens a route to new antimalarials				
JOURNAL	Biotechnol. Lett. 23, 917-921 (2001)				
REFERENCE	3 (bases 1 to 951)				
AUTHORS	Turgut-Balik,D., Moreton,K. and Holbrook,J.J.				
TITLE	Direct Submission				
JOURNAL	Submitted (08-SEP-2005) Biology, Faculty of Arts and Sciences, Elazig 23169, Turkey				
FEATURES	Location/Qualifiers				
source	1 951				

[Change Region Shown](#)

[Customize View](#)

Pick Primers
Design and test primers for this sequence using Primer-BLAST.

Recent Activity

[Turn Off](#) [Clear](#)

- Plasmodium falciparum isolate FCBR L-lactate dehydrogenase (LDH) gene, complete cds
- DQ198262 (1)
- dDQ198262 (0)
- Mus musculus SPARC-like 1 (Sparc1), mRNA
- decorin (421)

[Nucleotide](#)

All links from this record

- ▶ Protein
- ▶ PubMed (Weighted)
- ▶ Taxonomy
- ▶ Related Sequences

Internet

FASTA format

Default input format for sequence analysis software (nucleotide and protein) such as BLAST etc.

- For downloading the sequence click on **FASTA**
 - **COPY and PASTE** the information into an empty MS WORD document.
NOTE: **FASTA** starts with the Gene index.
 - Save as **text file** (no formatting)
- Case-sensitive: always use **CAPITAL** letters
- Use Courier font for easy alignment
- Format:
>My_Sequence_Name → Definition line
ARFDSHJDKDMFSMNDN → Sequence

File Edit View Favorites Tools Help

Search Nucleotide for Go Clear

Limits Preview/Index History Clipboard Details

Format: GenBank [FASTA](#) [Graphics](#) [More Formats](#)

Download Save Links

GenBank DQ198262.1

Plasmodium falciparum isolate FCBR L-lactate dehydrogenase (LDH) gene, complete cds

[Features](#) [Sequence](#)

LOCUS	DQ198262	951 bp	DNA	linear	INV 04-OCT-2005
DEFINITION	Plasmodium falciparum isolate FCBR L-lactate dehydrogenase (LDH) gene, complete cds.				
ACCESSION	DQ198262				
VERSION	DQ198262.1 GI:76563842				
KEYWORDS	.				
SOURCE	Plasmodium falciparum (malaria parasite P. falciparum)				
ORGANISM	Plasmodium falciparum Eukaryota; Alveolata; Apicomplexa; Aconoidasida; Haemosporida; Plasmodium; Plasmodium (Laverania).				
REFERENCE	1 (bases 1 to 951)				
AUTHORS	Turgut-Balik,D. and Holbrook,J.J.				
TITLE	Determination of the DNA and aminoacid sequences of lactate dehydrogenase gene from Plasmodium falciparum strains K1 and PF FCBR: A route to the design of new antimalarials				
JOURNAL	Turk. J. Biol. 25, 241-250 (2001)				
REFERENCE	2 (bases 1 to 951)				
AUTHORS	Turgut-Balik,D., Shoemark,D.K., Moreton,K.M., Sessions,R.B. and Holbrook,J.J.				
TITLE	Over-production of lactate dehydrogenase from Plasmodium falciparum opens a route to new antimalarials				
JOURNAL	Biotechnol. Lett. 23, 917-921 (2001)				
REFERENCE	3 (bases 1 to 951)				
AUTHORS	Turgut-Balik,D., Moreton,K. and Holbrook,J.J.				
TITLE	Direct Submission				
JOURNAL	Submitted (08-SEP-2005) Biology, Faculty of Arts and Sciences, Elazig 23169, Turkey				
FEATURES	Location/Qualifiers				
source	1 951				

[Change Region Shown](#)

[Customize View](#)

Pick Primers
Design and test primers for this sequence using Primer-BLAST.

Recent Activity

[Turn Off](#) [Clear](#)

- [Plasmodium falciparum isolate FCBR L-lactate dehydrogenase \(LDH\) gene, complete cds](#)
- [DQ198262 \(1\)](#)
- [dDQ198262 \(0\)](#)
- [Mus musculus SPARC-like 1 \(Sparc1\), mRNA](#)
- [decorin \(421\)](#)

[Nucleotide](#)

All links from this record

- ▶ Protein
- ▶ PubMed (Weighted)
- ▶ Taxonomy
- ▶ Related Sequences

Internet



CGCTCAGGATAAGACTTCGGCCGCTAGAGATCGGATCCCCGGCGATTATATAGCTCGATCGATC
 TTCTCTATATGGTATATACACACAGCGCGGATAGCACTGACTGATCTA
 CACAGACGTTCTACGTTCTACACTTAACTACCAATTCGGAGAGGGGGGAATGAGK
Nucleotide

My NCBI [?](#)
[\[Sign In\]](#) [\[Register\]](#)

All Databases

PubMed

Nucleotide

Protein

Genome

Structure

PMC

Taxonomy

Books

Search **Nucleotide** [▼](#) for [Go](#) [Clear](#)

Limits [Preview/Index](#) [History](#) [Clipboard](#) [Details](#)

Format: [GenBank](#) [FASTA](#) [Graphics](#) [More Formats▼](#)

[Download▼](#) [Save▼](#) [Links▼](#)

GenBank DQ198262.1

Plasmodium falciparum isolate FCBR L-lactate dehydrogenase (LDH) gene, complete cds

>gi|76563842|gb|DQ198262.1| Plasmodium falciparum isolate FCBR L-lactate dehydrogenase (LDH) gene, complete cds
 ATGGCACCAAAAGCAAAAATCGTTTACTGGCTCAGGTATGATTGGAGGAGTAATGGCTACCTTAATTG
 TTCAGAAAAATTAGGAGATGTAGTTTGTTCGATATTGTAAGAACATGCCACATGGAAAAGCTTTAGA
 TACATCTCATACTAATGTTATGGCATATTCAAATTGCAAAGTAAGTGGTCAACACTTATGACGATTG
 GCTGGAGCAGATGTAGTAATAGTAAACAGCTGGATTTACCAAGGCCCCCAGGGAAAGACTGACAAAGAATGGA
 ATAGAGATGATTATTACCATTAACAAACAAGATTATGATTGAAATTGGTGGTCATATTAAAGAACATTG
 TCCAATGCTTTATTATTGTTGAAACAAACCCAGTAGATGTTATGGTACAATTATACATCAACATTCA
 GGTGTTCTAAAAACAAAGATTATGGTTAGGTGGTGATTAGATAACATCAAGATTGAAGTTACATAT
 CTCAGAAAATTAAATGTATGCCAAGAGATGTAATGCACACATTGTAAGGTGCTCATGGAAATAAAATGGT
 TCTTTAAAAAGATACTTACATTGTAGGTGGTATCCCTTACAAGAATTATTAAATAACAAGTTAATTCT
 GATGCTGAATTAGAAGCTATATTGATAGAAGCTGTTAATACTGCAATTAGAAATTGTAACACTACATGCAT
 CACCATATGTTGCACCAAGCTGCTGCTATTATGAAATGGCTGAATCTACTTAAAGATTGAAAAAGT
 ATTAAATTGCTCAACCTGTTAGAAGGACAATATGGACACTCCGATATATTGGTGGTACACCTGTTGTT
 TTAGGTGCTAATGGTGGTGAACAAGTTATCGAATTACAATTAAAGTGAGGAAAAAGCTAAATTGATG
 AAGCCATAGCTGAAACTAAGAGAATGAAGGCATTAGCTTAA

[Change Region Shown](#)

[Customize View](#)

Pick Primers

Design and test primers for this sequence using Primer-BLAST.

Recent Activity

[Turn Off](#) [Clear](#)

[gb|DQ198262.1| \(951 lett...\)](#)

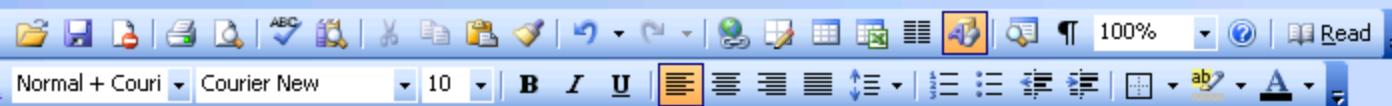
Plasmodium falciparum isolate FCBR L-lactate dehydrogenase (LDH) gene, complete cds

[DQ198262 \(1\)](#)

[dDQ198262 \(0\)](#) Nucleotide

Mus musculus SPARC-like 1 (Sparc1), mRNA

All links from this record



Normal + Courier

10

B

I

U

100%

?

Read

Save As

Save in:

- My Recent Documents
 - Desktop
 - My Documents
 - My Computer
 - My Network Places
- Desktop
- ROGU Membership form 2009.doc
 - Unused Desktop Shortcuts and setup files
 - SecureVPN
 - My Network Places
 - My Computer
 - My Documents

4 15 17 18

actate
ATTG
TAGA
TTTG
TGGA
ATTG
TTCA
ATAT
TGGT
TTCT
GCAT
AAGT
TGTT
GATG

File name:

gi.doc

Save

Cancel

Save as type:

Word Document (*.doc)

- Word Page (*.htm; *.html)
- Web Page, Filtered (*.htm; *.html)
- Document Template (*.dot)
- Rich Text Format (*.rtf)
- Plain Text (*.txt) **selected**
- Word 2007 Document (*.docx)

FASTA format: dUTPase

```
>gi|2443575|gb|AF018429.1|HSDUT1 Homo sapiens dUTPase (DUT) gene, exons 1 and 2
TCGGAAAAATGGGGGCCAGAGCAAACAAGAACAGAGCGAAAGCAAGAGGGCTAGGCAGCCAGAGGCGGCAGC
AAGACTCAAGACGCCAACGGCGCTTCTGGGCCAGGGCTTGCGCCATCCCTGGCTGCCGGGC
ACCGCCTCTCCACGCCCTCGTCCGGCGGCTGCGACTGCTCCGAGGTATGTTCCCAGGACGGCG
CGTCTTCAGGGTGAAGCCTGGCGCACGTCCGGAGGTGCCGAGGACCCAACCAGCCCCAAACTCTGGGGGA
AATGACTCCCCTCTGCCCTGCCCGCGCTGCTACCATTCTACGTCCTGCTCGCTCAGCGATG
CAAAACGCGCAGGCACGGCAGAGGGCGAAGCCGCGGTACTCTCCGGGCCAGGCCCGCCCTCGGCCGCG
CGGCCGCGCAGCACGGATTCCCCGGCGCTGTCCAGCGCTGGCCGCTGAGCCAAGGCTGCCGCCGGAGG
CAGTACAGTCGGGCCGCTGGCTGGAAGGGCGAGCTCTAAGGCGGGGGAGGCCCGGCCGGGCCG
GGTAGGAAAGCGGGGGAGGGGCTCCGGCGCTGGAAGGAATCCAACGCGGCTGAGGCTGTGGGGAGG
TAGGGTGGCGAGCGGCCCTCTGCGCGGGGGGGGGGGGGGGGGGGGTGGTCCATTAGGGTCCCCTGGC
GAGGGGGCGGCTTCTAGTGTGTGAAGGCGACGCCCTAGAACGCTCCCCTCAAAGTTGGCCCCACGCGCT
GAATGTGGAAAGTTGACTGGGACCCAGTAGTTCCCATCCCAAACCTGCTTCCGAGAAGGGCTTCAAAC
CCAAAATGTGAATCCCGCCTCCCCTCTCAGCCAGAACTGTGGACTCGTCCGGGAGGGCGGTGGTGG
GGCGGGGCTGGCGGGAAATTGGGTTGGCGCGCTCCCTGCGCGACGCTCATCGTGCCTCCTCTT
CCCCCGGTGGCTCCTCGCTCGCCCTGGCTCTGCCATGCCCTGCTGAAGAGACACCCGCCATTCA
CCCAGTAAGCGGGCCCGGCCTCGGGAGGTGGCGGCATGCAGCTCCGCTTGGCCGGCTCTCGAGCAG
CCACGGCCCCCACCGGGCTCCGCGCGCCGGCTACGACCTGTACAGGTGAGCGGGGACCTGCC
GCGAGGAGGCTGGGAAGGGCGGCCGTCCGCTGCCACAGCTAGAACAGTCACCGGAGAGATCACAGGAA
CACACTAGCTATAAATAGGATTCTGCCCTTTCGTGTTAAAATTAGCTTCATCTTGGCATAAAT
TAAATAGAGATTGGCAAAGACTGCAGAATAAGTAAAATAGCTATACGGTGTCTAGCAAGGCCTACTT
TGCAACGTTATTGTGCCCTTCTAAATAGAAGATAGAGAGGAAGGCCATGGTGGCTTCGAAGTGGCC
CGAGGGTGATGCTGTGCTCAATAGAAAAACCAAGGTGAGAGCCTAGATGTGAGCGTGAAAATACCTAAGA
AGGATGAACGAAGATGCATCTGCCTTAAAAGTTATTCTATACATTGACCGGCCAGGGCGGAATTG
AGAACATCTGAAAACGAAGGCAGACTGCCTGTATCTACCACACTTCATCTACAGCACGTTACTG
TACTAAAATCCGTATGCTGTTAGTCCTCACACATCCCTAACTAGATA
```

FASTA format: L-Lactate dehydrogenase protein

```
>gi|76563843|gb|ABA46355.1| L-lactate dehydrogenase [Plasmodium falciparum]
MAPKAKIVLVGSGMIGGMATLIVQKNLGDVVLFDIVKNMPHGKALDTSHTNVM
AYSNCVKSGSNTYDDLADVVIVTAGFTKAPGKSDKEWRDPLLPLNNKIMI
EIGGHIKKNCPNAFIIVVTNPVDVMVQLLHQHSGVPKNKIIGLGGVLDTSRLKYYI
SQKLNVCPRDVNAHIVGAHGNKMVLLKRYITVGGIPLQEFINNKLISDAELEAIF
DRTVNTALEIVNLHASPYVAPAAIIEMAESYLKDLKKVLICSTLLEGQYGHSDIF
GGTPVVLGANGVEQVIELQLNSEEKAKFDEAIAETKRMKALA
```

Protein sequences

- NCBI Protein
 - Plasmodium falciparum lactate dehydrogenase
 - [ABA46355.1](#)
 - [DQ198262.](#)
- Retrieve the protein sequence from the EXPASY website (Expert Protein Analysis System)
 - [WWW.EXPASY.CH](#)



AUTHORS Turgut-Balik,D., Shoemark,D.K., Moreton,K.M., Sessions,R.B. and Holbrook,J.J.

TITLE Over-production of lactate dehydrogenase from Plasmodium falciparum opens a route to new antimalarials

JOURNAL Biotechnol. Lett. 23, 917-921 (2001)

REFERENCE 3 (bases 1 to 951)

AUTHORS Turgut-Balik,D., Moreton,K. and Holbrook,J.J.

TITLE Direct Submission

JOURNAL Submitted (08-SEP-2005) Biology, Faculty of Arts and Sciences, Elazig 23169, Turkey

FEATURES Location/Qualifiers

source 1..951
 /organism="Plasmodium falciparum"
 /mol_type="genomic DNA"
 /isolate="FCBR"
 /db_xref="taxon:[5833](#)"

gene <1..>951
 /gene="LDH"

mRNA <1..>951
 /gene="LDH"
 /product="L-lactate dehydrogenase"

CDS 1..951
 /gene="LDH"
 /EC_number="[1.1.1.27](#)"
 /codon_start=1
 /product="L-lactate dehydrogenase"
 /protein_id="[ABA46355.1](#)"
 /db_xref="GI:[76563843](#)"
 /translation="MAPKAKIVLVGSGMIGGMATLIVQKNLGDVVLFDIVKNMPHKGK
 ALDTSHTNVMAYSNCKVGSNTYDDLADVVIVTAGFTKAPGKSDKEWRDDLLPLN
 NKIMIEIGGHIIKNCNPNAFIIVVTPVDVMWQLHQHSVGPKNKIIGLGGVLDTSLRK
 YYISQKLNVCPRDVNAHIVGAHGNKMVLKRYITVGGIPLQEFINNKLISDAELEAIF
 DRTVNTALEIVNLHASPYVAPAAAIIMEASYLKDLKKVLICSTLLEGQYGHSDIFGG
 TPVVVLGANGVEQVIELQLNSEEKAKFDEAIAETKRMKALA"

ORIGIN

```

1 atggcaccaa aagcaaaaat cgtttagtt ggctcaggta tgattggagg agtaatggct
61 acctaattt ttcagaaaaa tttaggagat gtatggat tcgatattgt aaagaacatg
121 ccatggaa aagctttaga tacatctcat actaatgtta tggcatattc aaattgcaaa

```

Click on here takes you to
Entrez Protein

Entrez Protein

Favorites L-lactate dehydrogenas... X difference between 3008-...

NCBI Resources How To

Protein Translations of Life

Search: Protein Limits Advanced search Help

Display Settings: GenPept Send to:

Search Clear

[Display Settings:](#) GenPept

[Send to:](#)

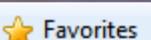
L-lactate dehydrogenase [Plasmodium falciparum]

GenBank: ABA46355.1

[FASTA](#) [Graphics](#)

[Go to:](#)

LOCUS ABA46355 316 aa linear INV 04-OCT-2005
DEFINITION L-lactate dehydrogenase [Plasmodium falciparum].
ACCESSION ABA46355
VERSION ABA46355.1 GI:76563843
DBSOURCE accession [DQ198262.1](#)
KEYWORDS .
SOURCE Plasmodium falciparum (malaria parasite P. falciparum)
ORGANISM [Plasmodium falciparum](#)
Eukaryota; Alveolata; Apicomplexa; Aconoidasida; Haemosporida;
Plasmodium; Plasmodium (Laverania).
REFERENCE 1 (residues 1 to 316)
AUTHORS Turgut-Balik,D. and Holbrook,J.J.
TITLE Determination of the DNA and aminoacid sequences of lactate
dehydrogenase gene from Plasmodium falciparum strains K1 and PF
FCBR: A route to the design of new antimalarials
JOURNAL Turk. J. Biol. 25, 241-250 (2001)
REFERENCE 2 (residues 1 to 316)
AUTHORS Turqut-Balik,D., Shoemark,D.K., Moreton,K.M., Sessions,R.B. and



L-lactate dehydrogenase



difference between 3008...

Region

```
/EC_number="1.1.1.21"
1..316
/region_name="PTZ00082"
/note="L-lactate dehydrogenase; Provisional"
/db_xref="CDD:173376"
```

Region

```
7..314
/region_name="LDH-like_MDH"
/note="L-lactate dehydrogenase-like malate dehydrogenase
proteins; cd01339"
/db_xref="CDD:133424"
```

Site

```
order(13..15,35..36,79..82,105,125,127,150,154,182)
/site_type="other"
/note="NAD(P) binding site"
/db_xref="CDD:133424"
```

Site

```
order(17,22,25,39..40,43,46..47,49..50,54,56..58,157..158,
160,162,231,236..238,241..242,245)
/site_type="other"
/note="dimer interface"
/db_xref="CDD:133424"
```

Site

```
order(56,58,168,173,175,177,194,196..198,252,254..257,283,
292)
/site_type="other"
/note="tetramer (dimer of dimers) interface"
/db_xref="CDD:133424"
```

Site

```
order(95,127,158,182,225,235)
/site_type="other"
/note="substrate binding site"
/db_xref="CDD:133424"
```

CDS

```
1..316
/gene="LDH"
/coded_by="DQ198262.1:1..951"
```

ORIGIN

```
1 mapkakivlv gsgmiggvma tlivqknlgd vvlfddivknm phgkaldtsh tnvmayasnck
61 vsgsntyddl agadvvivta gftkapgksd kewnrddllp lnnkimieig ghikkncpna
121 fiivvtnpvd vmvqllhqhs gpvknkiigl ggvltdtsrlk yyisqklncv prdnahivg
181 shankmullk rvitvaggml gefinnklis daeleasifdr tntaleivn lhasnwana
```