

Human LDH-B Sandwich ELISA Kit Datasheet

Please read it entirely before use

Catalogue Number: KE00287

Size: 5*96T

Sensitivity: 0.04 ng/mL

Range: 0.625-40 ng/mL

Usage: For the quantitative detection of human LDH-B concentrations in serum, plasma and cell lysate.

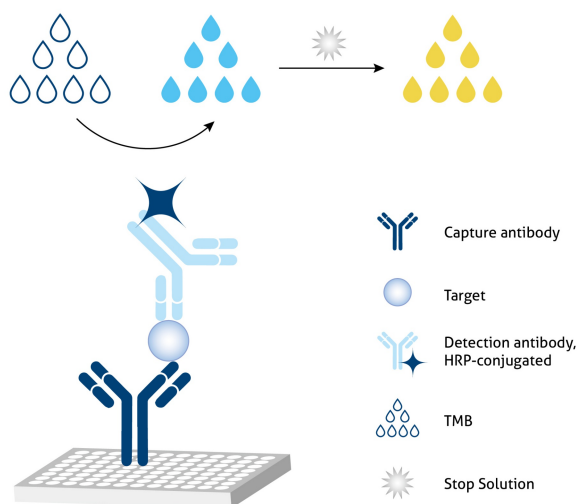
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1. Background

LDH is a tetrameric enzyme that is comprised by four subunits of two types: muscle/anaerobic (LDH-A) and heart/aerobic (LDH-B). Various combinations of LDH-A subunits and LDH-B subunits (which are produced from a different gene) can form homo- or heterotetramers, and assemble into five different combinations LDH-1 (4B), LDH-2 (3B,1A), LDH-3 (2B,2A), LDH-4 (1B,3A), and LDH-5 (4A). LDH-A converts pyruvate to lactate and is expressed in the skeletal muscle, while LDH-B converts lactate to pyruvate and is expressed more in the heart muscle.

2. Principle



Sandwich ELISA structure (Detection antibody labeled with HRP)

A capture antibody is pre-coated onto the bottom of wells which binds to analyte of interest. A detection antibody labeled with HRP also binds to the analyte. TMB acts as the HRP substrate and the solution color will change from colorless to blue. A stop solution containing sulfuric acid turns solution yellow. The color intensity is proportional to the quantity of bound protein which is measurable at 450 nm with the correction wavelength set at 630 nm.

3. Required Materials

- 3.1 A microplate reader capable of measuring absorbance at 450 nm with the correction wavelength set at 630 nm.
- 3.2 Calibrated, adjustable precision pipettes and disposable plastic tips. A manifold multi-channel pipette is recommended for large assays.
- 3.3 Plate washer: automated or manual.
- 3.4 Absorbent paper towels.
- 3.5 Glass or plastic tubes to prepare standard and sample dilutions.
- 3.6 Beakers and graduated cylinders.
- 3.7 Log-log or semi-log graph paper or computer and software for ELISA data analysis. A four-parameter logistic (4-PL) curve-fit is recommended.

4. Kit Components and Storage

Microplate - antibody coated 96-well microplate (8 well × 12 strips)	5 plates	Unopened Kit: Store at 2-8°C for 6 months or -20°C for 12 months. Opened Kit: All reagents stored at 2-8°C for 7 days. Please use a new standard for each assay.
Protein standard - 80 ng/bottle; lyophilized	10 bottles	
Detection antibody, HRP-conjugated (100×) - 600 µL/vial*	1 vial	
Additional Diluent AT-00287 - 30 mL/bottle. Only for human serum and plasma samples	1 bottle	
Sample Diluent PT 4B1 - 150 mL/bottle	2 bottles	
Detection Diluent - 150 mL/bottle	1 bottle	
Wash Buffer Concentrate (20×) - 150 mL/bottle	1 bottle	
Extraction Reagent - 150 mL/bottle	1 bottle	
Tetramethylbenzidine Substrate (TMB) - 60 mL/bottle	1 bottle	
Stop Solution - 60 mL/bottle	1 bottle	
Plate Cover Seals	15 pieces	

* Centrifugation immediately before use

5. Safety Notes

- 5.1 Avoid any skin and eye contact with Stop Solution and TMB. In case of contact, wash thoroughly with water.
- 5.2 Do not use the kit after the expiration date.
- 5.3 Do not mix or substitute reagents or materials from other kit lots or other sources.
- 5.4 Be sure to wear protective equipment such as gloves, masks and goggles during the experiment.
- 5.5 When using an automated plate washer, adding a 30 second soak period following the addition of Wash Buffer to improve assay precision

6. Sample Collection and Storage

- 6.1 Serum: Allow blood samples to clot for 30 minutes, followed by centrifugation for 15 minutes at 1000×g. Clear serum can be assayed immediately or aliquoted and stored at -20°C. Avoid repeated freeze-thaw cycles.
- 6.2 Plasma: Use EDTA, heparin, or citrate as an anticoagulant for plasma collection. Centrifuge for 15 minutes at 1000×g within 30 minutes of collection. The plasma can be assayed immediately or aliquoted and stored at -20°C. Avoid repeated freeze-thaw cycles.
- 6.3 Cell Lysate:
 - 1) Collect cells and wash by centrifuging at 500 x g for 5 minutes before resuspension in pre-cooled PBS buffer. Perform this step three times.
 - 2) Count cells and then discard the supernatant.
 - 3) Add protease inhibitor cocktail to the Extraction Reagent to a final concentration immediately prior to performing cell lysis.
 - 4) Add 1 mL of Extraction reagent (containing protease inhibitor cocktail) Per 1 x 10⁷ cells, Incubate cell suspension on ice for 30 minutes, use ultrasound to treat the samples.
 - 5) Centrifuge cell lysate at 10,000 x g for 5 minutes at 4°C.
 - 6) Measure the concentration of total protein in cell lysate using BCA assay. Where possible, keep samples on ice to avoid protein degradation.

7. Regent Preparation

7.1 Wash Buffer (1X): If crystals have formed in the concentrate, warm to room temperature and mix gently until the crystals have completely dissolved. Add 30 mL of Wash Buffer Concentrate(20X) to 570 mL deionized or distilled water to prepare 1X Wash Buffer.

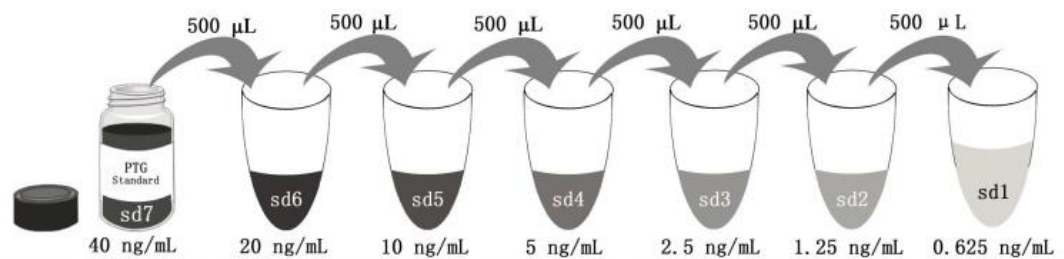
7.2 Detection Antibody, HRP-conjugated(1X): Dilute 100X Detection Antibody, HRP-conjugated 1:100 using Detection Diluent prior to assay. Suggested 1:100 dilution: 10 μ L 100X Detection Antibody, HRP-conjugated + 990 μ L Detection Diluent (Centrifuge the 100 X Detection Antibody solution, HRP-conjugated for a few seconds prior to use)

7.3 Sample Dilution: Different samples should be diluted with corresponding Sample Diluent, samples may require further dilution if the readout values are higher than the highest standard OD reading. Variations in sample collection, processing and storage may affect the results of the measurement.

Recommended Dilution for different sample types: 1:4 or 1:8 is recommended for human serum and plasma; 1:2 to 1:1,000 is recommended for cell lysate.

7.4 Standard Serial Dilution:

Add 2 mL Sample Diluent PT 4B1 in the protein standard.



Add # μ L of Standard diluted in the previous step	—	500 μ L	500 μ L	500 μ L	500 μ L	500 μ L	500 μ L
# μ L of Sample Diluent PT 4B1	2000 μL	500 μ L	500 μ L	500 μ L	500 μ L	500 μ L	500 μ L
	"sd7"	"sd6"	"sd5"	"sd4"	"sd3"	"sd2"	"sd1"

8. Assay Procedure Summary

Bring all reagents to room temperature before use (Detection antibody, HRP-conjugated can be used immediately). To avoid cross-contamination, change pipette tips between additions of each standard level, between sample additions, and between reagent additions. Also, use separate reservoirs for each reagent.

8.1 Take out the required number of microplate strips and return excess strips to the foil pouch containing the drying reagent pack and reseal; store at 4°C immediately. Microplate strips should be used in one week.

8.2 Preset the layout of the microplate, including control group, standard group and sample group;

For human serum or plasma, add 50 µL of Additional Diluent to the appropriate wells (No need incubation and wash);

For cell lysate, no need to add Additional Diluent, directly follow the next step.

8.3. Add 100 µL of each standard and sample to the appropriate wells. (Make sure sample addition is uninterrupted and completed within 5 to 10 minutes).

8.4 Seal plate with cover seal, pressing it firmly onto top of microwells. Incubate the plate for 2 hours at 37°C.

8.5 Wash

1) Gently remove the cover seal. Discard the liquid from wells by aspirating or decanting. Remove any residual solution by tapping the plate a few times on fresh paper towels.

2) Wash 4 times with 1X Wash Buffer, using at least 350-400 µL per well. Following the last wash, firmly tap plates on fresh towels 10 times to remove residual Wash Buffer. Avoid getting any towel fibers in the wells or wells drying out completely.

8.6 Add 100 µL of 1X Detection antibody, HRP-conjugated solution (refer to Reagent Preparation 7.2) to each well. Seal plate with cover seal and incubate for 60 minutes at 37°C.

8.7 Repeat wash step in 8.5.

8.8 Signal development: Add 100 µL of TMB substrate solution to each well, protected from light. Incubate for 15 to 20 minutes. Substrate Solution should remain colorless until added to the plate.

8.9 Quenching color development: Add 100 µL of Stop Solution to each well in the same order as addition of the TMB substrate. Mix by tapping the side of the plate gently. NB: Avoid skin and eye contact with the Stop solution.

8.10 Read results: Immediately after adding Stop solution read the absorbance on a microplate reader at a wavelength of 450 nm. If possible, perform a double wavelength readout (450 nm and 630 nm).

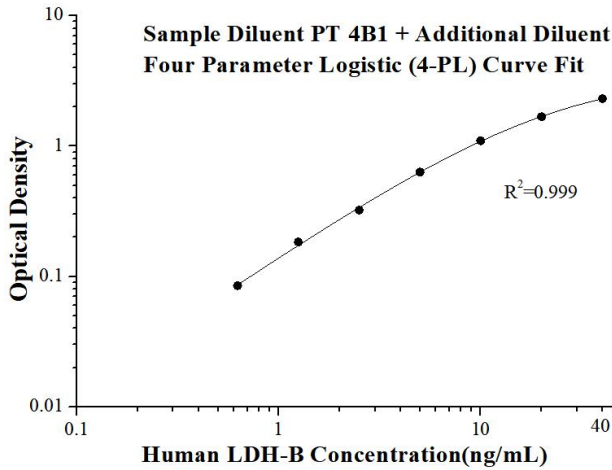
8.11 Data analysis: Calculate the average of the duplicate readings (OD value) for each standard and sample, and subtract the average of the zero standard absorbance. Construct a standard curve by plotting the mean absorbance for each standard on the y-axis against the concentration on the x-axis, use four-parameter logistic curve-fit (4-PL) analysis to do this. If the samples have been diluted, the OD readout from the standard curve must be multiplied by the dilution factor used.

Step	Reagent	Volume	Incubation	Wash	Notes
1	Additional diluent (Only for humna serum and plasma sample test)	50 µL	0 min	Do not wash	Add additional diluent 50 µL per well then add standard and samples immediately
2	Standard and Samples	100 µL	120 min	4 times	Cover Wells incubate at 37°C
3	Diluent Detection antibody, HRP-conjugated Solution	100 µL	60 min	4 times	Cover Wells incubate at 37°C
4	TMB Substrate	100 µL	15-20 min	Do not wash	Incubate in the dark at 37°C
5	Stop Solution	100 µL	0 min	Do not wash	-
6	Read plate at 450 nm and 630 nm immediately after adding Stop solution. DO NOT exceed 5 minutes.				

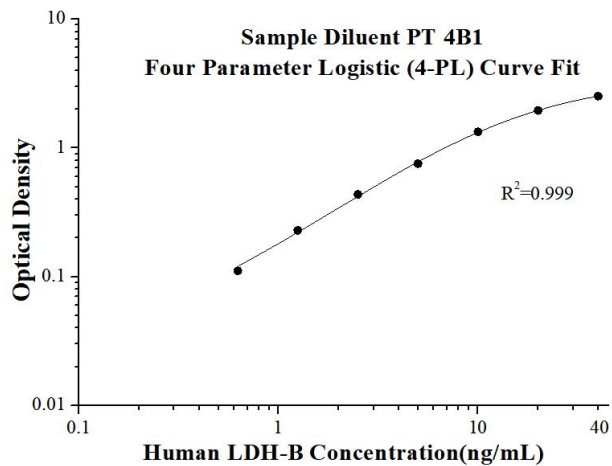
9. Validation Data

9.1 Standard curve

These standard curves are provided for demonstration only. A standard curve should be generated for each set of samples assayed.



(ng/mL)	O.D	Average	Corrected
0	0.077 0.072	0.074	-
0.625	0.160 0.159	0.160	0.085
1.25	0.257 0.261	0.259	0.184
2.5	0.383 0.413	0.398	0.323
5	0.674 0.736	0.705	0.631
10	1.144 1.200	1.172	1.098
20	1.681 1.821	1.751	1.677
40	2.393 2.370	2.382	2.307



(ng/mL)	O.D	Average	Corrected
0	0.076 0.077	0.076	-
0.625	0.191 0.183	0.187	0.111
1.25	0.302 0.309	0.305	0.229
2.5	0.520 0.503	0.511	0.435
5	0.819 0.844	0.831	0.755
10	1.411 1.408	1.410	1.333
20	2.018 2.043	2.030	1.954
40	2.642 2.556	2.599	2.522

9.2 Precision

Intra-assay Precision (Precision within an assay) Three samples of known concentration were tested 20 times on one plate to assess intra-assay precision.

Inter-assay Precision (Precision between assays) Three samples of known concentration were tested in 24 separate assays to assess inter-assay precision.

Intra-assay Precision					Inter-assay Precision				
Sample	n	Mean (ng/mL)	SD	CV%	Sample	n	Mean (ng/mL)	SD	CV%
1	20	18.65	0.72	3.84	1	24	17.81	0.79	4.46
2	20	4.56	0.20	4.47	2	24	4.49	0.14	3.16
3	20	0.99	0.08	8.25	3	24	1.13	0.09	8.15

9.3 Recovery

The recovery of human LDH-B spiked to three different levels throughout the range of the assay in various matrices was evaluated.

Sample Type		Average% of Expected	Range (%)
Human serum	1:2	114	86-130
	1:4	113	103-130
Cell lysate	1:4	102	92-117
	1:8	103	92-122

9.4 Sample values

Serum - Human serum samples were evaluated for the presence of human LDH-B in this assay.

Sample Type	Mean (ng/mL)	Range (ng/mL)
Human serum (n=16)	24.1	9.4-47.3

Cell lysate

	LDH-B (ng/mL)	Total protein (mg/mL)
Hela	11,920.0	4.4
HepG2	20.0	6.7
HEK-293	7,360.0	5.1
MCF-7	720.0	4.7

9.5 Sensitivity

The minimum detectable dose of human LDH-B is 0.04 ng/mL. This was determined by adding two standard deviations to the concentration corresponding to the mean O.D. of 20 zero standard replicates.

9.6 Linearity

To assess the linearity of the assay, samples were diluted with the appropriate **Sample Diluent** to produce samples with values within the dynamic range of the assay.

		Human serum	Cell lysate
1:2	Average% of Expected	100	100
	Range (%)	-	-
1:4	Average% of Expected	100	98
	Range (%)	76-115	92-102
1:8	Average% of Expected	102	91
	Range (%)	88-117	79-103
1:16	Average% of Expected	112	97
	Range (%)	103-121	88-107

9.7 Specificity

This assay recognizes natural and recombinant human LDH-B.

A sample containing 1.6 ug/mL of the recombinant human LDHA reads as 0.70 ng/mL (1.4% cross-reactivity).

10. References

- 1 Wu J, Liu L, Hu H, Gao Z, Lu S. Bioinformatic analysis and experimental identification of blood biomarkers for chronic nonunion. *J Orthop Surg Res.* 2020;15(1):208. Published 2020 Jun 5. doi:10.1186/s13018-020-01735-1
- 2 Li H, Li J, Wang Y, Yang T. Proteomic analysis of effluents from perfused human heart for transplantation: identification of potential biomarkers for ischemic heart damage. *Proteome Sci.* 2012;10(1):21. Published 2012 Mar 23. doi:10.1186/1477-5956-10-21
- 3 Duka T, Collins Z, Anderson SM, et al. Divergent lactate dehydrogenase isoenzyme profile in cellular compartments of primate forebrain structures. *Mol Cell Neurosci.* 2017;82:137-142. doi:10.1016/j.mcn.2017.04.007