TEL:4006-871-227 Web:www.ybio.net Email:shybio@126.com

YBH838Hu01 100µg

Recombinant Isocitrate Dehydrogenase 2, mitochondrial (IDH2)

Organism Species: Homo sapiens (Human)

Instruction manual

FOR IN VITRO USE AND RESEARCH USE ONLY NOT FOR USE IN CLINICAL DIAGNOSTIC PROCEDURES

10th Edition (Revised in Jan, 2014)

kDa 70

33

26

22

18

14

10

15% SDS-PAGE

[PROPERTIES]

Residues: Ala40~Gln452

Tags: Two N-terminal Tags, His-tag and T7-tag

Accession: P48735

Host: E. coli

Subcellular Location: Mitochondrion.

Purity: >95%

Endotoxin Level: <1.0EU per 1 µ g (determined by the

LAL method).

Formulation: Supplied as lyophilized form in PBS,

pH7.4, containing 5% trehalose, 0.01% sarcosyl.

Predicted isoelectric point: 8.3

Predicted Molecular Mass:

50.3kDa

Applications: SDS-PAGE; WB; ELISA; IP.

(May be suitable for use in other assays to be determined by the end user.)



[USAGE]

Reconstitute in sterile PBS, pH7.2-pH7.4.

[STORAGE AND STABILITY]

Storage: Avoid repeated freeze/thaw cycles.

Store at 2-8°C for one month.

Aliquot and store at -80°C for 12 months.

Stability Test: The thermal stability is described by the loss rate of the target protein. The loss rate was determined by accelerated thermal degradation test, that is, incubate the protein at 37°C for 48h, and no obvious degradation and precipitation were observed. (Referring from China Biological Products Standard, which was calculated by the Arrhenius equation.) The loss of this protein is less than 5% within the expiration date under appropriate storage condition.

[SEQUENCES]

The sequence of the target protein is listed below.

A DKRIKVAKPV VEMDGDEMTR IIWQFIKEKL ILPHVDIQLK YFDLGLPNRD QTDDQVTIDS

ALATQKYSVA VKCATITPDE ARVEEFKLKK MWKSPNGTIR NILGGTVFRE PIICKNIPRL

V P G W T K P I T I G R H A H G D Q Y K AT D F VA D R A G T F K M V F T P K D G S G V K E W E V Y

NFPAGGVGMG MYNTDESISG FAHSCFQYAI QKKWPLYMST KNTILKAYDG RFKDIFQEIF

DKHYKTDFDK NKIWYEHRLI DDMVAQVLKS SGGFVWACKN YDGDVQSDIL AQGFGSLGLM

TSVLVCPDGK TIEAEAAHGT VTRHYREHQK GRPTSTNPIA SIFAWTRGLE HRGKLDGNQD

LIRFAQMLEK VCVETVESGA MTKDLAGCIH GLSNVKLNEH FLNTTDFLDT IKSNLDRALG RQ