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

YB95368Mu01 100µg Carnitine Palmitoyltransferase 1A, Liver (CPT1A)

> Organism: Mus musculus (Mouse) Instruction manual

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

7th Edition (Revised in May, 2013)

kDa

94 66.2

45

33

26

20

### [PROPERTIES] Residues: Ala568~Lys773 (Accession # P97742), with two Nterminal Tags, His-tag and GST-tag. Host: F coli Subcellular Location: Mitochondrion outer membrane; Multipass membrane protein. **Purity: >95%** 14.4 **Endotoxin Level:** <1.0EU per 1µg (determined by the LAL 15% SDS-PAGE method). Formulation: Supplied as lyophilized form in PBS, pH7.4, containing 5% sucrose, 0.01% sarcosyl. Predicted isoelectric point: 6.7 Predicted Molecular Mass: 50.5kDa 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.

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## [ 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 target protein is fused with two N-terminal Tags, His-tag and GST-tag, its sequence is listed below.

MRNKKFELGL EFPNLPYYID GDVKLTQSMA IIRYIADKHN MLGGCPKERA EISMLEGAVL DIRYGVSRIA YSKDFETLKV DFLSKLPEML KMFEDRLCHK TYLNGDHVTH PDFMLYDALD VVLYMDPMCL DAFPKLVCFK KRIEAIPQID KYLKSSKYIA WPLQGWQATF GGGDHPPKSD G S T S G S G H H H H H H S A G LV P R G S TA I G M K E T A A A K F E R Q H M D S P D L G T L E V LFQ G P LG S E F- A FI QLA LQ L A HY K DMG K FC LT YE AS MTR LF R EG RTE TV R SC T T ESCNFVLAMM DPTTTAEQRF KLFKIACEKH QHLYRLAMTG AGIDRHLFCL YVVSKYLAVD SPFLKEVLSE PWRLSTSQTP QQQVELFDFE KYPDYVSCGG GFGPVADDGY GVSYIIVGEN FIHFHISSKF SSPETDSHRF GKHLRQAMMD IITLFGLTAN SKK