RPH480Hu01 100µg

Recombinant 3-Oxoacid Coenzyme A Transferase 1 (OXCT1)

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)

[PROPERTIES]

Residues: Thr40~Leu489

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

Accession: P55809

Host: *E. coli*

Subcellular Location: Mitochondrion matrix.

Purity: >95%

Endotoxin Level: <1.0EU per 1µg
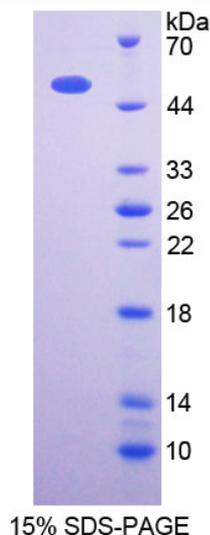
(determined by the LAL method).

Formulation: Supplied as lyophilized form in PBS, pH7.4, containing 5% trehalose, 0.01% sarcosyl.

Predicted isoelectric point: 6.3

Predicted Molecular Mass: 52.4kDa

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

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

kDa
70
44
33
26
22
18
14
10

15% SDS-PAGE

[USAGE]

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

[<u>STORAGE AND STABILITY</u>]

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.

[<u>SEQUENCES</u>]

The sequence of the target protein is listed below.

T KFYTDPVEAV KDIPDGATVL VGGFGLCGIP ENLIDALLKT GVKGLTAVSN NAGVDNFGLG
LLLRSKQIKR MVSSYVGENA EFERQYLSGE LEVELTPQGT LAERIRAGGA GVPAFYTPTG
YGTLVQEGGS PIKYNKDGSV AIASKPREVR EFNGQHFILE EAITGDFALV KAWKADRAGN
VIFRKSARNF NLPMCKAAET TVVEVEEIVD IGAFAPEDIH IPQIYVHRLI KGEKYEKRIE
RLSIRKEGDG EAKSAKPGDD VRERIIKRAA LEFEDGMYAN LGIGIPLLAS NFISPNITVH
LQSENGVLGL GPYPRQHEAD ADLINAGKET VTILPGASFF SSDESFAMIR GGHVDLTMLG
AMQVSKYGDL ANWMIPGKMV KGMGGAMDLV SSAKTKVVVT MEHSAKGNAH
KIMEKCTLPL TGKQCVNRII TEKAVFDVDK KKGLTLIEL