

YBC386Hu01 100µg

Recombinant Creatine Kinase, Mitochondrial 2, Sarcomeric (CKMT2) 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)

[<u>PROPERTIES</u>]

Residues: Glu40~Lys419

Tags: Two N-terminal Tags, His-tag and T7-tag Accession: P17540

Host: *E. coli*

Subcellular Location: Mitochondrion inner

membrane; Peripheral membrane protein;

Intermembrane side.

Purity: >95%

Endotoxin Level: <1.0EU per $1\,\mu\,g$ (determined by the

LAL method).

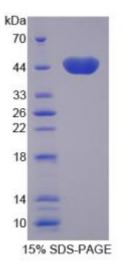
Formulation: Supplied as lyophilized form in PBS,

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

Predicted isoelectric point: 7.3

Predicted Molecular Mass:

47.0kDa





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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. E V R E Q P R L F P P S A D Y P D L R K H N N C M A E C LT P A I YA K L R N K V TP N G Y T L D Q C IQTGVDNPGH PFIKTVGMVA GDEESYEVFA DLFDPVIKLR HNGYDPRVMK HTTDLDASKI TOGOFDEHYV LSSRVRTGRS IRGLSLPPAC TRAERREVEN VAITALEGLK GDLAGRYYKL SEMTEQDQQR LIDDHFLFDK PVSPLLTCAG MARDWPDARG IWHNYDKTFL IWINEEDHTR VISMEKGGNM KRVFERFCRG LKEVERLIQE RGWEFMWNER LGYILTCPSN LGTGLRAGVH VRIPKLSKDP RFSKILENLR LQKRGTGGVD TAAVADVYDI SNIDRIGRSE VELVQIVIDG VNYLVDCEKK LERGQDIKVP PPLPQFGKK