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YB97050Mu01 100 μ g

Transcription Factor A, Mitochondrial (TFAM)

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)

[PROPERTIES]

Residues: Ser43-His243 (Accession # P40630), with two N-terminal Tags, His-tag and GST-tag.

Host: *E. coli*

Subcellular Location: Mitochondrion. Mitochondrion matrix > mitochondrion nucleoid.

Purity: >95%

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

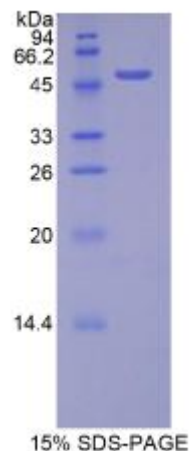
Formulation: Supplied as lyophilized form in PBS, pH7.4, containing 5% sucrose, 0.01% sarcosyl.

Predicted isoelectric point: 9.1

Predicted Molecular Mass: 50.6kDa

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

MRNKKFELGL EFPNLPYYID GDVCLTQSMA IIRYIADKHN MLGGCPKERA EISMLEGAVL
DIRYGVSRIA YSKDFETLKV DFLSKLPEML KMFEDRLCHK TYLNGDHVTH PDFMLYDALD
VVLYMDPMCL DAFPKLVCFK KRIEAIQID KYLKSSKYIA WPLQGWQATF GGGDHPKSD
G S T S G S G H H H H H S A G L V P R G S T A 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
LFQGPLGSEF- SSMGSYPK KPMSSYLRF S TEQLPKFKAK HPDAKLSELV RKIAALWREL
PEAEKKVYEA DFKAEWKAYK EAVSKYKEQL TPSQLMGMEK EARQRLKKK ALVKRRELIL
LGKPKRPRSA YNIYVSESFQ EAKDDSAQGK LKLVNEAWKN LSPEEKQAYI QLAKDDRIRY
DNEMKSWEEQ MAEVGRSDLI RRSVKRSGDI SEH