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YBG374Hu01 100µg

Recombinant ATPase, Ca++ TransPorting, Cardiac Muscle, Slow Twitch 2 (ATP2A2) Organism Species: Homo sapiens (Human)

Instruction manual

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

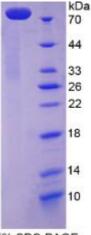
10th Edition (Revised in Jan, 2014)

[PROPERTIES]

Residues: Ile315~Met756 Tags: Two N-terminal Tags, His-tag and GST-tag Accession: P16615 Host: E. coli Subcellular Location: Endoplasmic reticulum membrane Multi-pass membrane protein. Sarcoplasmic reticulum membrane. **Purity: >90%** Endotoxin Level: <1.0EU per 1µg (determined by the LAL method). Formulation: Supplied as lyophilized form in PBS, pH7.4, 15% SDS-PAGE containing 5% trehalose, 0.01% sarcosyl. Predicted isoelectric point: 6.4 Predicted Molecular Mass: 78.2kDa 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.

ITTCLA LGTRRMAKKN AIVRSLPSVE TLGCTSVICS DKTGTLTTNQ MSVCRMFILD RVEGDTCSLN EFTITGSTYA PIGEVHKDDK PVNCHQYDGL VELATICALC NDSALDYNEA KGVYEKVGEA TETALTCLVE KMNVFDTELK GLSKIERANA CNSVIKQLMK KEFTLEFSRD RKSMSVYCTP NKPSRTSMSK MFVKGAPEGV IDRCTHIRVG STKVPMTSGV KQKIMSVIRE WGSGSDTLRC LALATHDNPL RREEMHLEDS ANFIKYETNL TFVGCVGMLD PPRIEVASSV KLCRQAGIRV IMITGDNKGT AVAICRRIGI FGQDEDVTSK AFTGREFDEL NPSAQRDACL NARCFARVEP SHKSKIVEFL QSFDEITAMT GDGVNDAPAL KKAEIGIAMG SGTAVAKTAS EMVLADDNFS TIVAAVEEGR AIYNNM