

**RPG374Hu01 100µg**

**Recombinant ATPase, Ca<sup>++</sup> 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).

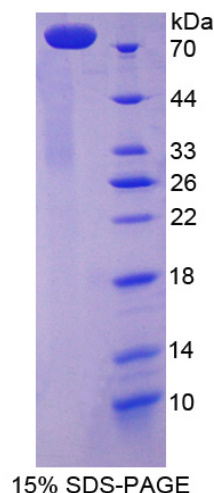
**Formulation:** Supplied as lyophilized form in PBS, pH7.4, containing 5% trehalose, 0.01% sarcosyl.

**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 MSVCRMFIELD  
RVEGDTCSLN EFTITGSTYA PIGEVHKDDK PVNCHQYDGL VELATICALC NDSALDYNEA  
KGVYEKVGEA TETALTCLVE KMNVFDTLTK GLSKIERANA CNSVIKQLMK KEFTLEFSRD  
RKSMSVYCTP NKPSRTSMK MFVKGAPEGV IDRCTHIRVG STKVPMMSGV KQKIMSVIRE  
WGSGSDLRC LALATHDNPL RREEMHLED ANFIKYETNL TFGCVGMLD PPRIEVASSV  
KLRCQAGIRV IMITGDNKGT AVAICRRIGI FGQDEDVTSK AFTGREFDEL NPSAQRDACL  
NARCFARVEP SHKSKIVEFL QSFDEITAMT GDGVNDAPAL KKAIEIGIAMG SGTAVAKTAS  
EMVLADDNFS TIVAAVEEGR AIYNNM