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YBB806Hu01 100µg

Recombinant FK506 Binding Protein 12 Rapamycin Associated Protein (FRAP)

Organism Species: Homo sapiens (Human)

Instruction manual

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

9th Edition (Revised in Jul, 2013)

[PROPERTIES]

Residues: Ala2226~Val2488 (Accession # P42345), with

Host: E. coli

Subcellular Location: Endoplasmic reticulum

membrane; Cytoplasmic side. Golgi apparatus

membrane; Peripheral membrane protein;

two N-terminal Tags, His-tag and T7-tag.

Mitochondrion outer membrane; Lysosome.

Cytoplasm.

Purity: >95%

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

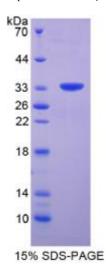
Formulation: Supplied as lyophilized form in PBS, pH7.4, containing 5% trehalose, 0.01% sarcosyl.

Predicted isoelectric point: 7.0

Predicted Molecular Mass: 33.7kDa

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

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





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[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 T7-tag, its sequence is listed below.

MGS SHHHH HH SSGLVP RGSH MASMTGG QQM GR GSEF- AVIPL STN SGLIGWV PHCDTLHALI RDYREKKKIL LNIEHRIMLR MAPDYDHLTL MQKVEVFEHA VNNTAGDDLA KLLWLKSPSS EVWFDRRTNY TRSLAVMSMV GYILGLGDRH PSNLMLDRLS GKILHIDFGD CFEVAMTREK FPEKIPFRLT RMLTNAMEVT GLDGNYRITC HTVMEVLREH KDSVMAVLEA FVYDPLLNWR LMDTNTKGNK RSRTRTDSYS AGQSVEILDG VELGEPAHKK TGTTVPESIH SFIGDGLV