TEL:4006-871-227 Web:www.ybio.net Email:shybio@126.com

### YBC019Hu01 100µg

### Recombinant Platelet Activating Factor Acetylhydrolase 2 (PAFAH2)

Organism Species: Homo sapiens (Human)

Instruction manua1

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

10th Edition (Revised in Jan, 2014)

# [ PROPERTIES ]

Residues: Met1~Leu392

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

Accession: Q99487

Host: E. coli

Subcellular Location: Cytoplasm.

Purity: >90%

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

method).

Formulation: Supplied as lyophilized form in PBS, pH7.4,

containing 5% trehalose, 0.01% sarcosyl.

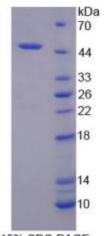
Predicted isoelectric point: 6.4

Predicted Molecular Mass:

47. 7kDa

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

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



15% SDS-PAGE



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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 sequence of the target protein is listed below.

MG V N Q S V G F P PV T G P H LV G C G D V M E G Q N L Q G S F F R L F Y P C Q K A E E T M E Q P

LWIPRYEYCT GLAEYLQFNK RCGGLLFNLA VGSCRLPVSW NGPFKTKDSG YPLIIFSHGL

GAFRTLYSAF CMELASRGFV VAVPEHRDRS AATTYFCKQA PEENQPTNES LQEEWIPFRR

VEEGEKEFHV RNPQVHQRVS ECLRVLKILQ EVTAGQTVFN ILPGGLDLMT LKGNIDMSRV

AVMGHSFGGA TAILALAKET QFRCAVALDA WMFPLERDFY PKARGPVFFI NTEKFQTMES

VNLMKKICAQ HEQSRIITVL GSVHRSQTDF AFVTGNLIGK FFSTETRGSL DPYEGQEVMV

RAMLAFLQKH LDLKEDYNQW NNLIEGIGPS LTPGAPHHLS SL