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YBM011Hu01 100 $\mu$ g

Recombinant Peptidase D (PEPD)

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: Ala2~Lys493

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

Accession: P12955

Host: *E. coli*

Purity: >95%

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

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

Predicted isoelectric point: 5.6

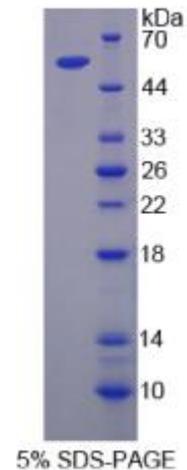
Predicted Molecular Mass: 58.1kDa

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.

AAATGPSFW LGNETLKVPL ALFALNRQRL CERLRKNPAV QAGSIVVLQG GEETQRYCTD  
TGVLFRQESF FHWAFGVTEP GCYGVIDVDT GKSTLFPRL PASHATWMGK IHSKEHFKEK  
YAVDDVQYVD EIASVLTSQK PSVLLTLRGV NTDSGSVCRE ASFDGISKFE VNNTILHPEI  
VECRVFKTDM ELEVLRNTK ISSEAHREVM KAVKVGKEY ELESLEHYC YSRGGMRHSS  
YTCICGSGEN SAVLHYGHAG APNDRTIQNG DMCLFDMGGE YYCFASDITC SFPANGKFTA  
D Q K A V Y E A V L R S S R A V M G A M K P G V W W P D M H R L A D R I H L E E L A H M G I L S G S  
VDAMVQAHLG AVFMPHGLGH FLGIDVHDVG GYPEGVERID EPGLRSLRTA RHLQPGMVL  
VEPGIYFIDH LLEALADPA RASFLNREVL QRFRGGGVR IEEDVVVTDS GIELLTCVPR  
TVEEIEACMA GCDKAFTPFS GPK