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YBA004Hu01 100 $\mu$ g

Angiotensin I Converting Enzyme (ACE)

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: Arg814~Gln1071 (Accession # P12821), with two  
N-terminal Tags, His-tag and GST-tag.

Host: *E. coli*

Subcellular Location: Secreted. Cell membrane;

Single-pass type I membrane protein.

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% sucrose, 0.01% sarcosyl.

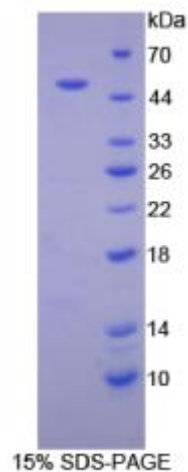
Predicted isoelectric point: 5.8

Predicted Molecular Mass: 61.5kDa

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 target protein is fused with two N-terminal Tags, His-tag and GST-tag, its sequence is listed below.

MSPILGYWKI KGLVQPTRLL LEYLEEKYEE HLYERDEGDK WRNKKFELGL EFPNLPYYID  
GDVKLTQ SMA IIRYIADKHN MLGGCPKERA EISMLEGAVL DIRYGVSRIA YSKDFETLKV  
DFLSKLP EML KMFEDRLCHK TYLNGDHVTH PDFMLYDALD VVLYMDPMCL DAFPKLVCFK  
KRIEAI PQID KYLKSSKYIA WPLQG WQATF GGGDHPPKSD GSTSGSGHHH HHSAGLVPR  
GSTAIGMKET AAKFERQHM DSPDLGTLEV LFQGPLGSEF-RLNGYVD AGDSWRSMYE  
TPSLEQDLER LFQELQPLYL NLHAYVRRAL HRHYGAQHIN LEGPIPAHLL GNMWAQTWSN  
IYDLVVPFPS APSMDTEAM LKQGWTPRRM FKEADFFTS LGLLPVPEF WNKSMLEKPT  
DGREVVCHAS AWFYNGKDF RIKQCTTVNL EDLVVAHHEM GHIQYFMQYK DLPVALREGA  
NPGFHEAIGDVLALSVSTPK HLHSLNLLSS EGGSEHDIN FLMKALDKI AFIPFSYLV D Q