RPA097Bo01 100 mg
Recombinant Matrix Metalloproteinase 1 (MMP1)
Organism Species: Bos taurus; Bovine (Cattle)
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: Phe19~Asn469
Tags: Two N-terminal Tags, His-tag and T7-tag
Accession: P28053
Host: E. coli
Subcellular Location: Secreted, extracellular
space, extracellular matrix.
Purity: >90\%
Endotoxin Level: <1.0EU per $1 \mu \mathrm{~g}$
(determined by the LAL method).
Formulation: Supplied as lyophilized form in PBS,

pH7.4, containing 5\% trehalose, $0.01 \%$ sarcosyl.
Predicted isoelectric point: 5.8
Predicted Molecular Mass: 55.1 kDa
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 $2-8^{\circ} \mathrm{C}$ for one month.
Aliquot and store at $-80^{\circ} \mathrm{C}$ 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 $37^{\circ} \mathrm{C}$ for 48 h , 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.
FP AATSETQEQD VETVKKYLEN YYNLNSNGKK VERQRNGGLI TEKLKQMQKF FGLRVTGKPD AETLNVMKQP RCGVPDVAPF VLTPGKSCWE NTNLTYRIEN YTPDLSRADV DQAIEKAFQL WSNVTPLTFT KVSEGQADIM ISFVRGDHRD NSPFDGPGGN LAHAFQPGAG IGGDAHFDDD EWWTSNFQDY NLYRVAAHEF GHSLGLAHST DIGALMYPSY TFSGDVQLSQ DDIDGIQAIY GPSQNPTQPV GPQTPEVCDS KLTFDAITTI RGEVMFFKDR FYMRTNPLYP EVELNFISVF WPQLPNGLQA AYEVADRDEV RFFKGNKYWA VKGQDVLRGY PRDIYRSFGF PRTVKSIDAA VSEEDTGKTY FFVANKCWRY DEYKQSMDAG YPKMIAEDFP GIGNKVDAVF QKGGFFYFFH GRRQYKFDPQ TKRILTLLKA NSWFNCRKN

