Cloud-Clone Corp.

RPB853Mu01 100µg

Recombinant Matrix Metalloproteinase 19 (MMP19)

Organism Species: Mus musculus (Mouse)

#### 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: Tyr99~Asp512 Tags: Two N-terminal Tags, His-tag and T7-tag Accession: Q9JHI0 Host: *E. coli* Subcellular Location: Cell membrane. Lipid-anchor. GPI-anchor. Extracellular side. Secreted, extracellular space, extracellular matrix. Purity: >95% Endotoxin Level: <1.0EU per 1µg (determined by the LAL method). Formulation: Supplied as Iyophilized form in PBS, pH7.4, containing 5% trehalose, 0.01% sarcosyl. Predicted isoelectric point: 8.9 Predicted Molecular Mass: 50.3kDa

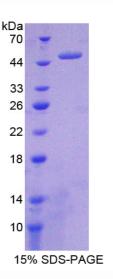
Applications: SDS-PAGE; WB; ELISA; IP.

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

# [ <u>USAGE</u> ]

Reconstitute in sterile PBS, pH7.2-pH7.4.

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## [ 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.

YL LLGHWRKKNL TFRIFNVPST LSLPRVRAAL HQAFKYWSSV APLTFREVKA GWADIRLSFH GRQSLYCSNT FDGPGKVLAH ADIPELGSIH FDKDELWTEG TYQGVNLRII AAHEVGHALG LGHSRYTQAL MAPVYAGYQP FFKLHPDDVA GIQALYGKRS PETRDEEEET EMLTVSPVTA KPGPMPNPCS GEVDAMVLGP RGKTYAFKGD YVWTVTDSGP GPLFQISALW EGLPGNLDAA VYSPRTRRTH FFKGNKVWRY VDFKMSPGFP MKFNRVEPNL DAALYWPVNQ KVFLFKGSGY WQWDELARTD LSRYPKPIKE LFTGVPDRPS AAMSWQDGQV YFFKGKEYWR LNQQLRVAKG YPRNTTHWMH CGSQTPDTNS STGDVTPSTT DTVLGTTPST MGSTLDIPSA TD