

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

YBS092Ra01 100µg

Recombinant Hyaluronoglucosaminidase 1 (HYAL1)

Organism Species: Rattus norvegicus (Rat)

Instruction manual

kDa 70

44

33

26

22 **18**

14

10

15% SDS-PAGE

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

10th Edition (Revised in Jan, 2014)

[PROPERTIES]

Residues: Ser40 Met449

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

Accession: Q76HN1

Host: E. coli

Subcellular Location: Secreted. Lysosome.

Purity: >95%

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: 5.3

Predicted Molecular Mass: 50.1kDa

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

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

[USAGE]

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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.

S MV S N R P F I T V WN A D T H W C L K DH G V D V D V S V FD V VA N K E Q N FQ G P N M T I F Y

REELGTYPYY TPTGEPVFGG LPONASLVTH LAHAFODIKA AMPEPDFSGL AVIDWEAWRP

RWA F N W D S K D IY Q Q R S M E LV RA E H P D W P E T LV E A E A Q G Q F QE A A E AW M A G

TL Q L G Q V L R P R G LW G Y Y G F P D C Y N Y D F L S P N Y T G Q C S L S I H D Q N D Q L G W L

WNQSYALYPS IYLPAALMGT GKSQMYVRYR VQEAFRLALV SRDPHVPIMP YVQIFYEKTD

YLLPLEELEH SLGESAAQGA AGAVLWISSE KTSTKESCQA IKAYMDSTLG PFILNVTSAA

LLCSEALCSG RGRCVRHPSY PEALLTLSPA SFSIEPTHDG RPLSLKGTLS LKDRAQMAMK

FKCRCYRGWS GEWCKKQDM