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#### YBB030Hu01 100µg

Recombinant Protein Tyrosine Phosphatase Receptor Type C (PTPRC) 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: Asp193~Lys575 (Accession # P08575),

with N-terminal His-Tag.

Host: E. coli

**Subcellular Location: Membrane; Single-pass** 

type I membrane protein. Membrane raft.

**Purity: >95%** 

Endotoxin Level: <1.0EU per 1µg

(determined by the LAL method).

Formulation: Supplied as lyophilized form in PBS,

pH7.4, containing 5% trehalose, 0.01% sarcosyl.

Predicted isoelectric point: 6.4

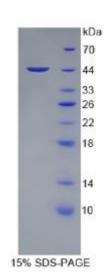
Predicted Molecular Mass: 45.1kDa

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 target protein is fused with N-terminal His-Tag, its sequence is listed below. MGHHHHHHSGSEF- DAYLNASE TTTLSPSGSA VISTTIATT PSKPTCDEKY ANITVDYLYN KETKLFTAKL NVNENVECGN NTCTNNEVHN LTECKNASVS ISHNSCTAPD KTLILDVPPG VEKFQLHDCT QVEKADTTIC LKWKNIETFT CDTQNITYRF QCGNMIFDNK EIKLENLEPE HEYKCDSEIL YNNHKFTNAS KIIKTDFGSP GEPQIIFCRS EAAHQGVITW NPPQRSFHNF TLCYIKETEK DCLNLDKNLI KYDLQNLKPY TKYVLSLHAY IIAKVQRNGS AAMCHFTTKS AP P S Q V W N M T VS M T S D N S M H VK C R P P R D R N G P H E RY H L E V EA G N T LV R N E SHKNCDFRVK DLQYSTDYTF KAYFHNGDYP GEPFILHHST SYNSK