YBC888BBo01 $100 \mu \mathrm{~g}$<br>Chromogranin B (CHGB) Organism Species: Bos taurus; Bovine (Cattle)<br>Instruction manual

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

## [ PROPERTIES ]

Residues: Leu309²Glu484 (Accession \# P23389), with two N-terminal Tags, His-tag and GST-tag.

Host: E. coli
Subcellular Location: Cytoplasmic vesicle, secretory vesicle membrane; Peripheral membrane protein; Lumenal side.

Purity: >95\%
Endotoxin Level: <1. OEU per $1 \mu \mathrm{~g}$ (determined by the
 LAL method).

Formulation: Supplied as lyophilized form in PBS, pH7.4, containing $5 \%$ sucrose, $0.01 \%$ sarcosyl.

Predicted isoelectric point: 5.9
Predicted Molecular Mass: 51.8kDa
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 target protein is fused with two N -terminal Tags, His-tag and GSTtag, its sequence is listed below.

MSPILGYWKI KGLVQPTRLL LEYLEEKYEE HLYERDEGDK WRNKKFELGL EFPNLPYYID gdvkltasma iiryiadkhn mlggcpkera eismlegavl dirygvsria yskdfetlkv DFLSKLPEML KMFEDRLCHK TYLNGDHVTH PDFMLYDALD VVLYMDPMCL DAFPKLVCFK KRIEAIPQID KYLKSSKYIA WPLQGWQATF GGGDHPPKSD GSTSGSGHHH HHHSAGLVPR
 RHSAAQAPGDLQGARFGGRGRGEHQALRRPSEESLEQENKRHGLSPDLNM A Q GYSEESEEERGPAPGPSYRARGGEAAAYSTLGQTDEKRFLGETHHRVQ ES Q R D K A R R R LP GELRNYLD YGEEKGEEAARGKWQPQGDPRDADENREEA RLRGKQYAPH HITE

