Cloud-Clone Corp.

#### RPB250Bo01 100µg Recombinant 5'-Nucleotidase, Ecto (NT5E) 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)

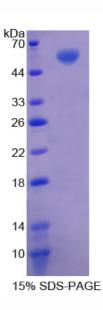
#### [PROPERTIES]

Residues: Ser48~Val527 Tags: Two N-terminal Tags, His-tag and T7-tag Accession: Q05927 Host: *E. coli* Subcellular Location: Cell membrane; Lipid-anchor, GPI-anchor. Purity: >95% Endotoxin Level: <1.0EU per 1µg (determined by the LAL method). Formulation: Supplied as Iyophilized form in 20mM Tris, 150mM NaCl, pH8.0, containing 1mM EDTA, 1mM DTT, 0.01% sarcosyl, 5% trehalose, and preservative. Predicted isoelectric point: 6.7 Predicted Molecular Mass: 56.2kDa 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 ddH<sub>2</sub>O.



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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.

## [<u>SEQUENCES</u>]

The sequence of the target protein is listed below.

SSK CVNASRCVGG VARLATKVHQ IRRAEPHVLL LDAGDQYQGT IWFTVYKGTE VAHFMNALGY DAMALGNHEF DNGVEGLIDP LLKEVNFPIL SANIKAKGPL ASKISGLYSP YKILTVGDEV VGIVGYTSKE TPFLSNPGTN LVFEDEITAL QPEVDKLKTL NVNKIIALGH SGFEVDKLIA QKVKGVDVVV GGHSNTFLYT GNPPSKEVPA GQYPFIVTSD DGRKVPVVQA YAFGKYLGYL KVEFDEKGNV VTSHGNPILL NSSIPEDPNI KADINKWRVK LDNYSTQELG KTIVYLDGTA QSCRFRECNM GNLICDAMIN NNLRHPDEMS WNHVSMCILN GGGIRSPIDE RNNGTITWEN LAAVLPFGGT FDLVQLKGST LKKAFEHSVH RYGQATGEFL QVGGIHVVYD ISRNPGDRVV KLEVLCTQCR VPSYEPLRMD KVYKVILPSF LVSGGDGFQM IKDEKIKHDS GDQDINV