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YBD179Hu01 100ug Recombinant Nucleobindin 1 (NUCB1) Organism Species: Homo sapiens (Human) 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: Glu37~Leu461 Tags: Two N-terminal Tags, His-tag and T7-tag Accession: Q02818 Host: E. coli Subcellular Location: Golgi Apparatus. Cis-golgi Network Membrane; Peripheral Membrane Protein; Lumenal Side. **Purity: >90%** Endotoxin Level: <1.0EU per 1µg (determined by the LAL method). Formulation: Supplied as lyophilized form in 20mM Tris, 15% SDS-PAGE 150mM NaCl, pH8.0, containing 1mM EDTA, 1mM DTT, 0.01% sarcosyl, 5% trehalose, and preservative. Predicted isoelectric point: 5.1

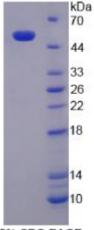
Predicted Molecular Mass: 53.8kDa

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 ddH₂O.





[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.

EETP ATE SPD TG LY YHRY LQE VID VLETD GH FRE KLQA AN AED I KSG KLSR ELD FVSHHVRTKL DELKRQEVSR LRMLLKAKMD AEQDPNVQVD HLNLLKQFEH LDPQNQHTFE ARDLELLIQT ATRDLAQYDA AHHEEFKRYE MLKEHERRRY LESLGEEQRK EAERKLEEQQ RRHREHPKVN VPGSQAQLKE VWEELDGLDP NRFNPKTFFI LHDINSDGVL DEQELEALFT K E L E K V Y D P K N E E D D M R E M E E E R L R M R E H V M K N V D T N Q D R LV T L E E F L A S T Q R K E F G D T G E G W E T V E M H P AY T E E E L R R F E E E L A A R E A E L N A K A Q R L S Q ET E A L G R S Q G RL E A Q K R E L Q QAV L H M E Q R K QQ Q Q Q G H K APA A H P E G Q L KFHPDTDDVP VPAPAGDQKE VDTSEKKLLE RLPEVEVPQH L