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YBE802Hu01 100µg

Recombinant Glutamate Receptor, Ionotropic, AMPA 2 (GRIA2)

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: Asn25~Ala543 Tags: Two N-terminal Tags, His-tag and T7-tag Accession: P42262 Host: E. coli Subcellular Location: Cell membrane, Multi-pass membrane protein, Endoplasmic reticulum membrane, Cell junction, synapse, postsynaptic cell membrane. **Purity: >90%** 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: 6.2 Predicted Molecular Mass: 62.0kDa 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 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.

NSIQIG GLFPRGADQE YSAFRVGMVQ FSTSEFRLTP HIDNLEVANS FAVTNAFCSQ FSRGVYAIFG FYDKKSVNTI TSFCGTLHVS FITPSFPTDG THPFVIQMRP DLKGALLSLI EYYQWDKFAY LYDSDRGLST LQAVLDSAAE KKWQVTAINV GNINNDKKDE MYRSLFQDLE LKKERRVILD CERDKVNDIV DQVITIGKHV KGYHYIIANL GFTDGDLLKI QFGGANVSGF QIVDYDDSLV SKFIERWSTL EEKEYPGAHT TTIKYTSALT YDAVQVMTEA FRNLRKQRIE ISRRGNAGDC LANPAVPWGQ GVEIERALKQ VQVEGLSGNI KFDQNGKRIN YTINIMELKT NGPRKIGYWS EVDKMVVTLT ELPSGNDTSG LENKTVVVTT ILESPYVMMK KNHEMLEGNE RYEGYCVDLA AEIAKHCGFK YKLTIVGDGK YGARDADTKI WNGMVGELVY GKADIAIAPL TITLVREEVI DFSKPFMSLG ISIMIKKPQK SKPGVFSFLD PLA