C Cloud-Clone Corp.

RPG369Rb01 100uq

Recombinant ATPase, Na+/K+ Transporting Alpha 1 Polypeptide (ATP1a1) Organism Species: Oryctolagus cuniculus (Rabbit)

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: Thr339~Leu772 **Tags:** Two N-terminal Tags, His-tag and T7-tag Accession: Q9N0Z6 Host: E. coli Subcellular Location: Cell membrane; Multi-pass membrane protein. **Purity:** >90% Endotoxin Level: <1.0EU per 1µg (determined by the LAL method). **Formulation:** Supplied as lyophilized form in 20mM Tris, 150mM NaCl, pH8.0, containing 1mM EDTA, 1mM DTT, 0.01% sarcosyl, 5% trehalose, and preservative. Predicted isoelectric point: 5.3 Predicted Molecular Mass: 50.9kDa 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.



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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.

TV TVCLTLTAKR MARKNCLVKN LEAVETLGST STICSDKTGT LTQNRMTVAH MWFDNQIHEA DTTENQSGVS FDKTSATWLA LSRIAGLCNR AVFQANQENL PILKRAVAGD ASESALLKCI ELCCGSVKEM RERYTKIVEI PFNSTNKYQL SIHKNLNANE PRHLLVMKGA PERILDRCSS ILLHGKEQPL DEELKDAFQN AYLELGGLGE RVLGFCHLLL PDEQFPEGFQ FDTDEVNFPV DNLCFIGLIS MIDPPRAAVP DAVGKCRSAG IKVIMVTGDH PITAKAIAKG VGIISEGNET VEDIAARLNI PVSQVNPRDA KACVVHGSDL KDMTSEQLDD ILKYHTEIVF ARTSPQQKLI IVEGCQRQGA IVAVTGDGVN DSPALKKADI GVAMGIAGSD VSKQAADMIL LDDNFASIVT GVEEGRLIFD NL