

**RPG369Rb01 10µg**

**Recombinant ATPase, Na<sup>+</sup>/K<sup>+</sup> 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

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11th Edition (Revised in May, 2016)

## **[ PROPERTIES ]**

**Source:** Prokaryotic expression.

**Host:** *E. coli*

**Residues:** Thr339~Leu772

**Tags:** N-terminal His-Tag

**Tissue Specificity:** Endothelial Cells.

**Subcellular Location:** Cell membrane; Multi-pass membrane protein.

**Purity:** >90%

**Endotoxin Level:** <1.0EU per 1µg (determined by the LAL method).

**Traits:** Freeze-dried powder

**Buffer formulation:** 20mM Tris, 150mM NaCl, pH8.0, containing 1mM EDTA, 1mM DTT, 0.01% sarcosyl, 5%Trehalose and Proclin300.

**Original Concentration:** 200ug/mL

**Applications:** SDS-PAGE; WB; ELISA; IP; CoIP; Reporter Assays; Purification; Amine Reactive Labeling.

(May be suitable for use in other assays to be determined by the end user.)

**Predicted isoelectric point:** 5.3

**Predicted Molecular Mass:** 50.9kDa

**Accurate Molecular Mass:** 51kDa as determined by SDS-PAGE reducing conditions.

## [ USAGE ]

Reconstitute in 20mM Tris, 150mM NaCl (pH8.0) to a concentration of 0.1-1.0 mg/mL. Do not vortex.

## [ STORAGE AND STABILITY ]

**Storage:** Avoid repeated freeze/thaw cycles.

Store at 2-8°C for one month.

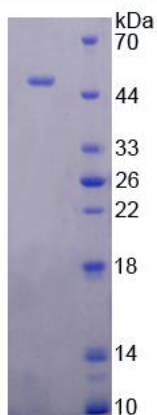
Aliquot and store at -80°C for 12 months.

**Stability Test:** The thermal stability is described by the loss rate. 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. The loss rate is less than 5% within the expiration date under appropriate storage condition.

## [ SEQUENCE ]

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TV TVCLTLTAKR
MARKNCLVKN LEAVETLGST STICSDKTGT LTQNRMTVAH MWFDNQIHEA
DTTENQSGVS FDKTSATWLA LSRIAGLCNR AVFQANQENL PILKRAVAGD
ASESALLKCI ELCCGSVKEM RERYTKIVEI PFNSTNKYQL SIHKNLNANE
PRHLLVMKGA PERILDRCSS ILLHGKEQPL DEELKDAFQN AYLELGGGLGE
RVLGFCHLLL PDEQFPEGFQ FDTDEVNFPV DNLCFIGLIS MIDPPRAAVP
DAVGKCRSAG IKVIMVTGDH PITAKAIAGK VGIISEGNET VEDIAARLNI
PVSQVNPRDA KACVVHGS DL KDMTSEQLDD ILKYHTEIVF ARTSPQQKLI
IVEGCQRQGA IVAVTGDGVN DSPALKKADI GVAMGIAGSD VSKQAADMIL
LDDNFASIVT GVEEGR LIFD NL
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**[ IDENTIFICATION ]**



**Figure 1. SDS-PAGE**