

**RPE837Ra01 100µg**  
**Aldehyde Dehydrogenase 7 Family, Member A1 (ALDH7A1)**  
**Organism Species: Rattus norvegicus (Rat)**  
***Instruction manual***

FOR IN VITRO USE AND RESEARCH USE ONLY  
NOT FOR USE IN CLINICAL DIAGNOSTIC PROCEDURES

9th Edition (Revised in Jul, 2013)

**[ PROPERTIES ]**

**Residues:** Leu28~Gln539 (Accession # Q64057),  
with N-terminal His-Tag.

**Host:** *E. coli*

**Subcellular Location:** Nucleus. Cytoplasm,  
cytosol. Mitochondrion.

**Purity:** >95%

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

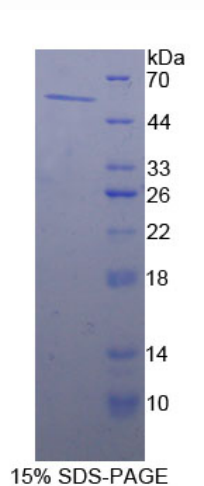
**Formulation:** Supplied as lyophilized form in PBS,  
pH7.4, containing 5% sucrose, 0.01% sarcosyl.

**Predicted isoelectric point:** 6.5

**Predicted Molecular Mass:** 57.0kDa

**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 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 target protein is fused with N-terminal His-Tag, its sequence is listed below.

MGHHHHHSGS-LMS TLLIHHPQYA WLQDLGLRED NEGVFNGSWG GRGEVITTYC  
PANNEPIARV RQASMKYEE TIGKAKKAWN IWADIPAPKR GEIVRKIGDA LREKIQLLGR  
LVSLEMKGIL VEGIGEVQEY VDVCYAAGL SRMIGGPTLP SERPGHALME QWNPLGLVGI  
ITAFNFPVAV FGWNNIALI TGNVCLWKGA PTTSLVSIIV TKIIAKVLED NLLPGAICSL  
TCGGADMGT A MARDERNLL SFTGSTQVVK QVALMVQERF GKSLLELGGN NAIIFEDAD  
LSLVLP S AAVGTAGQRC TTVRRLFLHE SIHDEVVDRL KNAYSQIRVG NPWDPNILYG  
PLHTKQAVSM FVQAVEEAKK EGGTVVYGGK VMDHPGNYVE PTIVTGLVHD APIVHKETFA  
PILYVFKFKN EEEVFEWNE VKQGLSSIF TKDLGRIFRW LGPKGSDCGI VNVNIPTSGA  
EIGGAFGG EK HTGGGREGS DAWKQYMRRS TCTINYSTAL PLAQGKIFQ