

RPD046Ra01 100µg
Recombinant Natriuretic Peptide Receptor 3 (NPR3)
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)

kDa 70

## [PROPERTIES]

**Residues:** Ala45~Pro467 (Accession # P41740) linked with SNHKSSVDKLAAALEHHHHHHH, with two N-terminal Tags, His-tag and T7-tag.

Host: E. coli

Subcellular Location: Membrane; Single-pass

type I membrane protein.

**Purity: >95%** 

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

Formulation: Supplied as lyophilized form in PBS,

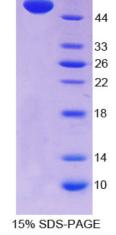
pH7.4, containing 5% trehalose, 0.01% sarcosyl.

Predicted isoelectric point: 6.4

Predicted Molecular Mass: 53.4kDa

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 two N-terminal Tags, His-tag and T7-tag, its sequence is listed below.

MGSSHHHHHH SSGLVPRGSH MASMTGGQQM GRGS- AQKIEV LVLLPRDDSY LFSLARVRPA IEYALRSVEG NGTGRKLLPP GTRFQVAYED SDCGNRALFS LVDRVAAARG AKPDLILGPV CEYAAAPVAR LASHWDLPML SAGALAAGFQ HKDTEYSHLT RVAPAYAKMG EMMLALFRHH HWSRAALLYS DDKLERNCYF TLEGVHEVFQ EEGLHTSAYN FDETKDLDLD DIVRYIQGSE RVVIMCASGD TIRRIMLAVH RHGMTSGDYA FFNIELFNSS SYGDGSWKRG DKHDFEAKQA YSSLQTVTLL RTAKPEFEKF SMEVKSSVEK QGLNEEDYVN MFVEGFHDAI LLYVLALHEV LRAGYSKKDG GKIIQQTWNR TFEGIAGQVS IDANGDRYGD FSVVAMTDTE AGTQEVIGDY FGKEGRFKMR SNVKYPWGSL KLRIDETRIV EHTNSSP SNHKSSVDKLAAALEHHHHHH