

**RPC972Mu01 100µg**

**Recombinant Transforming Growth Factor Beta Receptor II (TGFβR2)**

**Organism Species: *Mus musculus* (Mouse)**

***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:** Ile209~Thr590

**Tags:** Two N-terminal Tags, His-tag and T7-tag

**Accession:** Q62312

**Host:** *E. coli*

**Subcellular Location:** Cell 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:** 5.6

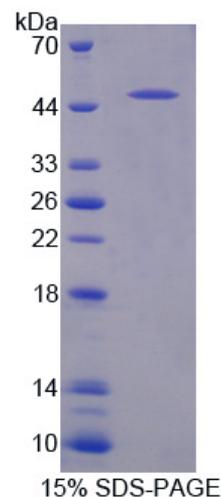
**Predicted Molecular Mass:** 47.5kDa

**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 sequence of the target protein is listed below.

II FYCYRVHRQQ KLSPSWESSK PRKLMDFSDN CAILEDERS DISSTCANNI NHNTELLPIE  
LDTLVGKGRF AEVYKAKLKQ NTSEQFETVA VKIFPYEEYS SWKTEKDIFS DINLKHENIL  
QFLTAEERKT ELGKQYWLIT AFHAKGNLQE YLTRHVISWE DLRKLGSSLA RGHHLHSDH  
TPCGRPKMPI VHRDLKSSNI LVKNDLTCCL CDFGLSLRLD PTLVDDLAN SGQVGTARYM  
APEVLESRMN LENVESFKQT DVYSMALVLW EMTSRCNAVG EVKDYEPFPG SKVREHPCVE  
SMKDSVLRDR GRPEIPSWL NHQGIQVCE TLTECWHDHP EARLTAQCVA ERFSELEHPE  
RLSGRSCSQE KIPEDGSLNT