

**RPB491Mu01 100µg**  
**Recombinant Interferon Gamma Receptor 1 (IFNgR1)**  
**Organism Species: *Mus musculus* (Mouse)**  
***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:** Gly23~Asp253 (Accession # P15261),  
with N-terminal GST-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).

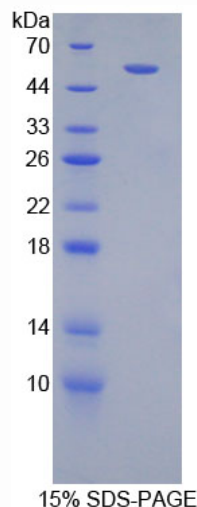
**Formulation:** Supplied as lyophilized form in PBS,  
pH7.4, containing 5% trehalose, 0.01% sarcosyl.

**Predicted isoelectric point:** 5.4

**Predicted Molecular Mass:** 52.6kDa

**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 GST-tag, its sequence is listed below.

MSPILGYWKI KGLVQPTRL L LEYLEEKYEE HLYERDEGDK WRNKKFELGL EFPNLPYYID  
GDVKLTQSMA IIRYIADKHN MLGGCPKERA EISMLEGAVL DIRYGVSR IAYSKDFETLKV  
DFLSKLPEML KMFEDRLCHK TYLNGDHVTH PDFMLYDALD VVLYMDPMCL DAFPKLVCFK  
KR IEAIPQID KYLKSSKYIA WPLQGWQATF GGGDHPPKSD GGPLGSPEF-GSGALTST  
EDPEPPSVPV PTNVLIKSYN LNPVVCWEYQ NMSQTPIFTV QVKVYSGSWT DSCTNISDHC  
CNIYEQIMYP DVSAWARVKA KVGQKESDYA RSKEFLMCLK GKVGP PGLEI RRKKEEQLSV  
LVFHPEVVVN GESQGT MFGD GSTCYTFDYT VYVEHNRSGE ILHTKHTVEK EECNETLCEL  
NISVSTLDSR YCISVDGISS FWQVRTEKSK DVCIPPFHDD RKD