



RPE766Mu01 10 μ g
Recombinant Interleukin 22 Receptor (IL22R)
Organism Species: *Mus musculus* (Mouse)
Instruction manual

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

12th Edition (Revised in Aug, 2016)

[**PROPERTIES**]

Source: Prokaryotic expression.

Host: *E. coli*

Residues: Pro258~Cys552

Tags: N-terminal His-Tag

Subcellular Location: Membrane; Single-pass type I membrane protein.

Purity: >95%

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: 200µg/mL

Applications: Positive Control; Immunogen; SDS-PAGE; WB.

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

Predicted isoelectric point: 5.0

Predicted Molecular Mass: 35.4kDa

Accurate Molecular Mass: 40kDa as determined by SDS-PAGE reducing conditions.

Note: The possible reasons that the actual band size differs from the predicted are as follows:

1. Splice variants: Alternative splicing may create different sized proteins from the same gene.
2. Relative charge: The composition of amino acids may affects the charge of the protein.
3. Post-translational modification: Phosphorylation, glycosylation, methylation etc.
4. Post-translation cleavage: Many proteins are synthesized as pro-proteins, and then cleaved to give the active form.
5. Polymerization of the target protein: Dimerization, multimerization etc.

[**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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PVP PNSLNVQRVL TFQPLRFIQE HVLIPVLDLS GPSSLPQPIQ
YSQVVVSGPR EPPGAVWRQS LSDLTYVGQS DVSILQPTNV PAQQTLSPPS
YAPKAVPEVQ PPSYAPQVAS DAKALFYSPQ QGMKTRPATY DPQDILDSCP
ASYAVCVEDS GKDSTPGILS TPKYLKTKGQ LQEDTLVRSC LPGDLSLQKV
TSLGEGETQR PKSLPSPLGF CTDRGPDLHT LRSEEPETPR YLKGALSLLS
SVQIEGHPVS LPLHVHSVSC SPSDEGPPSW GLLDSLVCCK DEGPAVETEA
MC
```

[IDENTIFICATION]

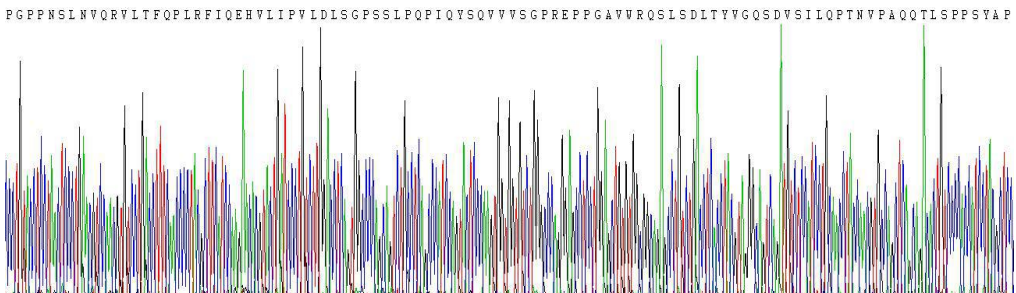


Figure 1. Gene Sequencing (Extract)

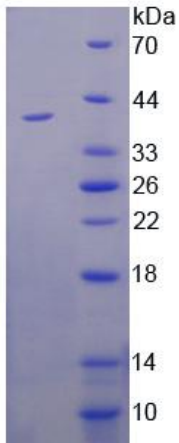


Figure 2. SDS-PAGE