

EPA049Ra61 100ug

Eukaryotic Interferon Gamma (IFNg)

Organism Species: Rattus norvegicus (Rat)

Instruction manual

FOR IN VITRO USE AND RESEARCH USE ONLY

NOT FOR USE IN CLINICAL DIAGNOSTIC PROCEDURES

11th Edition (Revised in May, 2016)

[PROPERTIES]

Source: Eukaryotic expression.

Host: 293F cell

Residues: Gln23~Cys156

Tags: N-terminal His Tag

Homology: Human 39%, mouse 83%

Tissue Specificity: Spleen.

Subcellular Location: Secreted.

Purity: >98%

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

Traits: Freeze-dried powder

Buffer Formulation: 20mM Tris, 150mM NaCl, pH8.0, containing 1mM EDTA, 1mM DTT, 5%Trehalose and Proclin300.

Original Concentration: 200ug/mL

Predicted isoelectric point: 9.5

Predicted Molecular Mass: 17.1kDa

Accurate Molecular Mass: 19kDa as determined by SDS-PAGE reducing conditions.

Applications: SDS-PAGE; WB; ELISA; IP; CoIP; EMSA; Reporter Assays; Purification; Amine Reactive Labeling.

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

[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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QGT L I E S L E S L K N Y F N S S S M D A M E G K S L  
L L D I W R N W Q K D G N T K I L E S Q I I S F Y L R L F E V L K D N Q A I S N N I S V I E S H L I  
T N F F S N S K A K K D A F M S I A K F E V N N P Q I Q H K A V N E L I R V I H Q L S P E S S L R K  
R K R S R C
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[IDENTIFICATION]

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TGAGGGCAGCTATTGAAAGCTGAGAGCTGAGAGCTATTTTACTCGAGTGGATGGATGCTGGGAGGAAAGAGGCTCTCTTGGATATCTGGAGGAGCTGGCAAGAGGCGGTAACTGCAAAATCTTGAGAGCCAGATTATCTCTTCTACCTGAGACTCTTGGAGCTTGAAAGCAAGCAGGCCATC  
Q G T L I E S L E S L K N Y F N S S W D A T E G K S L L L D I V R N V Q K D G N T K I E S Q I I S F Y L R L F E V L K D N Q A I
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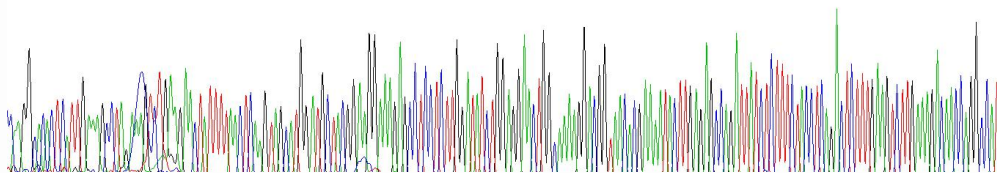


Figure 1. Gene Sequencing (extract)



Figure 2. SDS-PAGE