

RPA090Hu01 100µg

Recombinant Colony Stimulating Factor 1, Macrophage (MCSF)

Organism Species: Homo sapiens (Human)

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: Glu33~Ser496

Tags: Two N-terminal Tags, His-tag and GST-tag

Accession: P09603

Host: *E. coli*

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

Purity: >90%

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

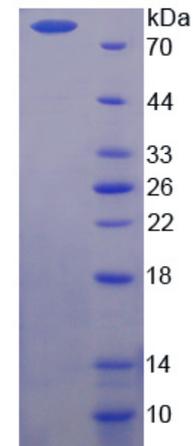
Formulation: Supplied as lyophilized form in PBS, pH7.4, containing 5% trehalose, 0.01% sarcosyl.

Predicted isoelectric point: 5.0

Predicted Molecular Mass: 80.2kDa

Applications: SDS-PAGE; WB; ELISA; IP.

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



15% SDS-PAGE

[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.

EEVSEYCS HMIGSGHLQS LQRLIDSQME TSCQITFEFV DQEQLKDPVC YLKKAFLLVQ
DIMEDTMRFR DNTPNIAIV QLQELSLRLK SCFTKDYEEL DKACVRTFYE TPLQLLEKVK
NVFNETHKLL DKDWNIFSKN CNNSFAECSS QDVVTKPCDN CLYPKAIPSS DPASVSPHQF
LAPSMAPVAG LTWEDSEGTE GSSLLPGEQP LHTVDPGSAK QRPPRSTCQS FEPPETPVVK
DSTIGGSPQP RPSVGAFNPG MEDILDSAMG TNWVPEEASG EASEIPVPQG TELSPSRPGG
GSMQTEPARP SNFLSASSPL PASAKGQQA DVTGTALPRV GPVRPTGQDW
NHTPQKTDHP SALLRDPPEP GSPRISSLRP QGLSNPSTLS AQPQLSRSHS SGSVLPLGEL
EGRRSTRDRR SPAEPEGGA SEGAARPLPR FNSVPLTDTG HERQSEGSFS PQLQES