Certificate of Analysis CSF1R, 100 µg

Colony Stimulating Factor 1 Receptor, Histidine-tagged



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Description:

Part Number: PR4598A

Lot Number: 2112355C

Immediate Storage: -80°C

Shipping Conditions: dry ice

Recombinant human, catalytic domain (amino acids 538-910), Histidinetagged, expressed in insect cells. Activated in-vitro via autophosphorylation.

Specific Activity:

480 nmoles of phosphate transferred to poly [Glu, Tyr] 4:1 substrate per minute per mg of total protein at 30°C. Activity determined at a final protein concentration of 2 μ g/mL.

Concentration:

0.15~mg/mL total protein as measured using the Bradford protein assay with BSA as a standard.

Calculated 3,350 nM.

Aliases:

FMS, CSFR, C-FMS, CD115

Storage and Handling:

For maximum recovery please spin prior to use. Unless noted below, aliquots of the 5 ug, 10ug and 20ug sizes of kinase are not recommended as materials can be used in original packaging until exhausted. For larger sizes, the number of freeze/thaws may be reduced by preparing aliquots, aliquots below 20 μ L are not recommended. **Please never store a kinase diluted.** If properly stored at -80°C, this product is guaranteed for 6 months from date of purchase.

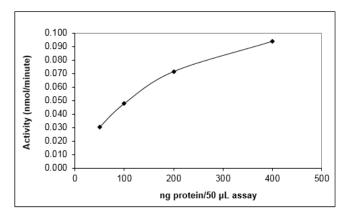
This product is not stable at room temperature (50% activity loss is observed within 6 hours at room temperature).

Storage Buffer:

50 mM Tris (pH 7.5), 100 mM NaCl, 0.05 mM EDTA, 0.05% Triton® X–100, 2 mM DTT and 50% Glycerol.

QUALITY ASSURANCE

CSF1R Activity Graph



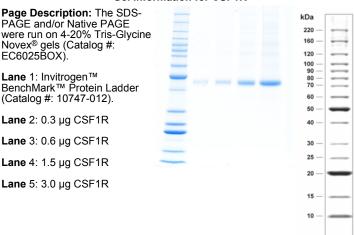
Dilution Buffer:

20 mM Tris (pH 7.5), 0.05% NP-40, 0.1 mg/mL BSA, 1 mM DTT and 10% Glycerol.

Assay Conditions:

CSF1R was pre-diluted in enzyme dilution buffer and assayed in 50 mM HEPES (pH 7.5), 10 mM MgCl₂, 10% Glycerol, 2.5 mM DTT, 0.01% Triton[®] X-100, 200 μM ATP, 200 μg/mL poly [Glu, Tyr] 4:1 substrate and trace [³²P]-γ-ATP for 10 minutes at 30°C.

Gel Information for CSF1R



Purity:

80% as determined by a Coomassie® blue stained SDS-PAGE gel.

Molecular Weight:

44.8 kDa. Calculated from the protein sequence(s).

Mass Spectrometry:

CSF1R was subjected to proteolytic digest followed by mass spec analysis. The resulting MS/MS data verified CSF1R identity by comparison against the amino acid sequence(s) of the recombinant protein.

Certificate of Analysis

Protein sequence alignment with reference sequence(s)

GenBank Accession Number: NP_005202

1 MYKYKQKPKY QVRWKIIESY EGNSYTFIDP TQLPYNEKWE FPRNNLQFGK TLGAGAFGKV VEATAFGLGK EDAVLKVAVK MLKSTAHADE KEALMSELKI IVGN CSFIR 537 LYKYKQKPKY QVRWKIIESY EGNSYTFIDP TQLPYNEKWE FPRNNLQFGK TLGAGAFGKV VEATAFGLGK EDAVLKVAVK MLKSTAHADE KEALMSELKI IVGN CSFIR 637 MSHLGQHENI VNLLGACTHG GPVLVITEYC CYGDLLNFLR RKAEAMLGPS LSPGQDPEGG VDYKNIHLEK KYVRRDSGFS SQGVDTYVEM RPVSTSSNDS 201 FSEQDLDKED GRPLELRDLL HFSSQVAQGM AFLASKNCIH RDVAARNVLL TNGHVAKIGD FGLARDIMND SNYIVKGNAR LPVKWMAPES IFDCVYTVQS 737 FSEQDLDKED GRPLELRDLL HFSSQVAQGM AFLASKNCIH RDVAARNVLL TNGHVAKIGD FGLARDIMND SNYIVKGNAR LPVKWMAPES IFDCVYTVQS 301 DVWSYGILLW EIFSLGLNPY PGILVNSKFY KLVKDGYQMA QPAFAPKNIY SIMQACWALE PTHRPTFQQI CSFLKGVEAC QLGTDDYDIP TTHHHHH. 837 DVWSYGILLW EIFSLGLNPY PGILVNSKFY KLVKDGYQMA QPAFAPKNIY SIMQACWALE PTHRPTFQQI CSFL

* highlighted residues denote differences from the reference protein sequence(s).

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Anita Targosz, Associate Director

Date: 24/Jul/2019

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