Certificate of Analysis CSNK1G2 (CK1 gamma 2), 10 µg

Casein Kinase 1 gamma 2, Histidine-tagged

Part Number: PV3499 Lot Number: 2711134B Immediate Storage: -80°C Shipping Conditions: dry ice



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Description:

Recombinant human full length protein, Histidine-tagged, expressed in insect cells. No special measures were taken to activate this kinase.

Specific Activity:

230 nmoles of phosphate transferred to dephosphorylated casein substrate per minute per mg of total protein at 30°C. Activity determined at a final protein concentration of 2 μ g/mL.

Concentration:

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

Calculated 4,570 nM.

Aliases:

CK1g2

Storage and Handling:

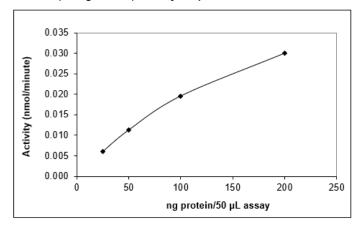
For maximum recovery please spin prior to use. Unless noted below, aliquots of the 5 ug, 10ug and 20ug sizes of kinase are not recommended as materials can be used in original packaging until exhausted. For larger sizes, the number of freeze/thaws may be reduced by preparing aliquots, aliquots below 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.

Storage Buffer:

50 mM Tris (pH 7.5), 150 mM NaCl, 0.5 mM EDTA, 0.04% Triton® X–100, 4 mM DTT and 50% Glycerol.

QUALITY ASSURANCE

CSNK1G2 (CK1 gamma 2) Activity Graph



Dilution Buffer:

20 mM Tris (pH 7.5), 0.05% Triton® X–100, 0.1 mg/mL BSA, 2 mM DTT, 0.5 mM Na $_3\rm VO_4$ and 10% Glycerol.

Assay Conditions:

CSNK1G2 (CK1 gamma 2) was pre-diluted in enzyme dilution buffer and assayed in 12.5 mM Tris (pH 7.5), 10 mM MgCl₂, 1 mM EGTA, 0.5 mM Na₃VO₄, 5 mM β –glycerophosphate, 2 mM DTT, 0.01% Triton® X-100, 200 μM ATP, 420 μg/mL dephosphorylated casein substrate and trace [32 P]-γ-ATP for 10 minutes at 30°C.

Gel Information for CSNK1G2 (CK1 gamma 2)



Purity:

95% as determined by a SDS-PAGE gel stained with SimplyBlue™ SafeStain.

Molecular Weight:

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

Mass Spectrometry:

CSNK1G2 (CK1 gamma 2) was subjected to proteolytic digest followed by mass spec analysis. The resulting MS/MS data verified CSNK1G2 (CK1 gamma 2) identity by comparison against the amino acid sequence(s) of the recombinant protein.

Part Number: PV3499, Lot Number: 2711134B

Protein sequence alignment with reference sequence(s)

GenBank Accession Number: NP_001310

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1 MDFDKKGGKG ETEEGRRMSK AGGGRSSHGI RSSGTSSGVL MVGPNFRVGK KIGCGNFGEL RLGKNLYTNE YVAIKLEPIK SRAPQLHLEY RFYKQLSATE IVGN CSNK1G2 1 MDFDKKGGKG ETEEGRRMSK AGGGRSSHGI RSSGTSSGVL MVGPNFRVGK KIGCGNFGEL RLGKNLYTNE YVAIKLEPIK SRAPQLHLEY RFYKQLSATE NP_001310

101 GVPQVYYFGP CGNYNAMVLE LLGPSLEDLF DLCDRTFTLK TVLMIAIQLI TRMEYVHTKS LIYRDVKPEN FLVGRPGTKR QHAIHIIDFG LAKEYIDPET 101 GVPQVYYFGP CGNYNAMVLE LLGPSLEDLF DLCDRTFTLK TVLMIAIQLI TRMEYVHTKS LIYRDVKPEN FLVGRPGTKR QHAIHIIDFG LAKEYIDPET 102 KKHIPYREHK SLTGTARYMS INTHLGKEQS RRDDLEALGH MFMYFLRGSL PWQGLKADTL KERYQKIGDT KRATPIEVLC ENFPEEMATY LRYVRRLDFF 102 KKHIPYREHK SLTGTARYMS INTHLGKEQS RRDDLEALGH MFMYFLRGSL PWQGLKADTL KERYQKIGDT KRATPIEVLC ENFPEEMATY LRYVRRLDFF 103 KKHIPYREHK SLTGTARYMS INTHLGKEQS RRDDLEALGH MFMYFLRGSL PWQGLKADTL KERYQKIGDT KRATPIEVLC ENFPEEMATY LRYVRRLDFF 103 KKHIPYREHK SLTGTARYMS INTHLGKEQS RRDDLEALGH MFMYFLRGSL PWQGLKADTL KERYQKIGDT KRATPIEVLC ENFPEEMATY LRYVRRLDFF 103 KKHIPYREHK SLTGTARYMS INTHLGKEQS RRDDLEALGH MFMYFLRGSL PWQGLKADTL KERYQKIGDT KRATPIEVLC ENFPEEMATY LRYVRRLDFF 103 KKHIPYREHK SLTGTARYMS INTHLGKEQS RRDDLEALGH MFMYFLRGSL PWQGLKADTL KERYQKIGDT KRATPIEVLC ENFPEEMATY LRYVRRLDFF 103 KKHIPYREHK SLTGTARYMS INTHLGKEQS RRDDLEALGH MFMYFLRGSL PWQGLKADTL KERYQKIGDT KRATPIEVLC ENFPEEMATY LRYVRRLDFF 104 KKHIPYREHK SLTGTARYMS INTHLGKEQS RRDDLEALGH MFMYFLRGSL PWQGLKADTL KERYQKIGDT KRATPIEVLC ENFPEEMATY LRYVRRLDFF 104 KKHIPYRLDFF 105 KKHIPYREHK SLTGTARYMS INTHLGKEQS RRDDLEALGH MFMYFLRGSL PWQGLKADTL KERYQKIGDT KRATPIEVLC ENFPEEMATY LRYVRRLDFF 104 KKHIPYRLDFF 105 KKHIPYRLDFF
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Chevohn Joseph, Director, Quality Date: 10/May/2023

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For Research Use Only. Not for use in diagnostic procedures.

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