

Datasheet for ABIN3134259 CDK5 Protein (AA 1-292) (Strep Tag)



Overview

| Quantity: | 1 mg |
|-------------------------------|---|
| Target: | CDK5 |
| Protein Characteristics: | AA 1-292 |
| Origin: | Mouse |
| Source: | Cell-free protein synthesis (CFPS) |
| Protein Type: | Recombinant |
| Purification tag / Conjugate: | This CDK5 protein is labelled with Strep Tag. |
| Application: | SDS-PAGE (SDS), Western Blotting (WB), ELISA |

Product Details

| Brand: | AliCE® |
|------------------|---|
| Sequence: | MQKYEKLEKI GEGTYGTVFK AKNRETHEIV ALKRVRLDDD DEGVPSSALR EICLLKELKH |
| | KNIVRLHDVL HSDKKLTLVF EFCDQDLKKY FDSCNGDLDP EIVKSFLFQL LKGLGFCHSR |
| | NVLHRDLKPQ NLLINRNGEL KLADFGLARA FGIPVRCYSA EVVTLWYRPP DVLFGAKLYS |
| | TSIDMWSAGC IFAELANAGR PLFPGNDVDD QLKRIFRLLG TPTEEQWPAM TKLPDYKPYP |
| | MYPATTSLVN VVPKLNATGR DLLQNLLKCN PVQRISAEEA LQHPYFSDFC PP |
| | Sequence without tag. The proposed Strep-Tag is based on experience s with the expression |
| | system, a different complexity of the protein could make another tag necessary. In case you |
| | have a special request, please contact us. |
| Characteristics: | Key Benefits: |
| | Made in Germany - from design to production - by highly experienced protein experts. |

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- Protein expressed with ALiCE® and purified in one-step affinity chromatography
- These proteins are normally active (enzymatically functional) as our customers have reported (not tested by us and not guaranteed).
- State-of-the-art algorithm used for plasmid design (Gene synthesis).

This protein is a **made-to-order protein** and will be made for the first time for your order. Our experts in the lab try to ensure that you receive soluble protein.

The big advantage of ordering our **made-to-order proteins** in comparison to ordering custom made proteins from other companies is that there is no financial obligation in case the protein cannot be expressed or purified.

Expression System:

- ALICE®, our Almost Living Cell-Free Expression System is based on a lysate obtained from Nicotiana tabacum c.v.. This contains all the protein expression machinery needed to produce even the most difficult-to-express proteins, including those that require post-translational modifications.
- During lysate production, the cell wall and other cellular components that are not required for
 protein production are removed, leaving only the protein production machinery and the
 mitochondria to drive the reaction. During our lysate completion steps, the additional
 components needed for protein production (amino acids, cofactors, etc.) are added to
 produce something that functions like a cell, but without the constraints of a living system all that's needed is the DNA that codes for the desired protein!

Concentration:

- The concentration of our recombinant proteins is measured using the absorbance at 280nm.
- The protein's absorbance will be measured against its specific reference buffer.
- We use the Expasy's ProtParam tool to determine the absorption coefficient of each protein.

| Purification: | One-step Strep-tag purification of proteins expressed in Almost Living Cell-Free Expression System (AliCE®). |
|---------------|--|
| Purity: | > 70-80 % as determined by SDS PAGE, Western Blot and analytical SEC (HPLC). |
| Grade: | custom-made |

Target Details

| Target: | CDK5 |
|-------------------|----------------------|
| Alternative Name: | Cdk5 (CDK5 Products) |

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Cyclin-dependent kinase 5 (EC 2.7.11.1) (Cell division protein kinase 5) (Cyclin-dependent-like kinase 5) (Serine/threonine-protein kinase PSSALRE) (Tau protein kinase II catalytic subunit) (TPKII catalytic subunit),FUNCTION: Proline-directed serine/threonine-protein kinase essential for neuronal cell cycle arrest and differentiation and may be involved in apoptotic cell death in neuronal diseases by triggering abortive cell cycle re-entry. Interacts with D1 and D3-type G1 cyclins. Phosphorylates SRC, NOS3, VIM/vimentin, p35/CDK5R1, MEF2A, SIPA1L1, SH3GLB1, PXN, PAK1, MCAM/MUC18, SEPT5, SYN1, DNM1, AMPH, SYNJ1, CDK16, RAC1, RHOA, CDC42, TONEBP/NFAT5, MAPT/TAU, MAP1B, histone H1, p53/TP53, HDAC1, APEX1, PTK2/FAK1, huntingtin/HTT, ATM, MAP2, NEFH and NEFM. Regulates several neuronal development and physiological processes including neuronal survival, migration and differentiation, axonal and neurite growth, synaptogenesis, oligodendrocyte differentiation, synaptic plasticity and neurotransmission, by phosphorylating key proteins. Negatively regulates the CACNA1B/CAV2.2 -mediated Ca(2+) release probability at hippocampal neuronal soma and synaptic terminals (By similarity). Activated by interaction with CDK5R1 (p35) and CDK5R2 (p39), especially in postmitotic neurons, and promotes CDK5R1 (p35) expression in an autostimulation loop. Phosphorylates many downstream substrates such as Rho and Ras family small GTPases (e.g. PAK1, RAC1, RHOA, CDC42) or microtubule-binding proteins (e.g. MAPT/TAU, MAP2, MAP1B), and modulates actin dynamics to regulate neurite growth and/or spine morphogenesis. Phosphorylates also exocytosis associated proteins such as MCAM/MUC18, SEPT5, SYN1, and CDK16/PCTAIRE1 as well as endocytosis associated proteins such as DNM1, AMPH and SYNJ1 at synaptic terminals. In the mature central nervous system (CNS), regulates neurotransmitter movements by phosphorylating substrates associated with neurotransmitter release and synapse plasticity, synaptic vesicle exocytosis, vesicles fusion with the presynaptic membrane, and endocytosis. Promotes cell survival by activating anti-apoptotic proteins BCL2 and STAT3, and negatively regulating of JNK3/MAPK10 activity. Phosphorylation of p53/TP53 in response to genotoxic and oxidative stresses enhances its stabilization by preventing ubiquitin ligase-mediated proteasomal degradation, and induces transactivation of p53/TP53 target genes, thus regulating apoptosis. Phosphorylation of p35/CDK5R1 enhances its stabilization by preventing calpain-mediated proteolysis producing p25/CDK5R1 and avoiding ubiquitin ligase-mediated proteasomal degradation. During aberrant cell-cycle activity and DNA damage, p25/CDK5 activity elicits cellcycle activity and double-strand DNA breaks that precedes neuronal death by deregulating HDAC1. DNA damage triggered phosphorylation of huntingtin/HTT in nuclei of neurons protects neurons against polyglutamine expansion as well as DNA damage mediated toxicity. Phosphorylation of PXN reduces its interaction with PTK2/FAK1 in matrix-cell focal adhesions (MCFA) during oligodendrocytes (OLs) differentiation. Negative regulator of Wnt/beta-catenin

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| signaling pathway. Activator of the GAIT (IFN-gar | nma-activated inhibitor of translation) |
|--|--|
| pathway, which suppresses expression of a post- | -transcriptional regulon of proinflammatory |
| genes in myeloid cells, phosphorylates the linker | domain of glutamyl-prolyl tRNA synthetase |
| (EPRS) in a IFN-gamma-dependent manner, the i | nitial event in assembly of the GAIT complex. |
| Phosphorylation of SH3GLB1 is required for auto | phagy induction in starved neurons. |
| Phosphorylation of TONEBP/NFAT5 in response | to osmotic stress mediates its rapid nuclear |
| localization. MEF2 is inactivated by phosphorylat | ion in nucleus in response to neurotoxin, thus |
| leading to neuronal apoptosis. APEX1 AP-endode | oxyribonuclease is repressed by |
| phosphorylation, resulting in accumulation of DN | A damage and contributing to neuronal death. |
| NOS3 phosphorylation down regulates NOS3-der | ived nitrite (NO) levels. SRC phosphorylation |
| mediates its ubiquitin-dependent degradation and | d thus leads to cytoskeletal reorganization. |
| May regulate endothelial cell migration and angio | genesis via the modulation of lamellipodia |
| formation. Involved in dendritic spine morphogen | esis by mediating the EFNA1-EPHA4 |
| signaling. The complex p35/CDK5 participates in | the regulation of the circadian clock by |
| modulating the function of CLOCK protein: phosp | horylates CLOCK at 'Thr-451' and 'Thr-461' |
| and regulates the transcriptional activity of the C | LOCK-BMAL1 heterodimer in association with |
| altered stability and subcellular distribution. {ECC | :0000250 UniProtKB:Q03114, |
| ECO:0000269 PubMed:12941275, ECO:0000269 | PubMed:17143272, |
| ECO:0000269 PubMed:20473298, ECO:0000269 | PubMed:21210220, |
| ECO:0000269 PubMed:24235147}. | |

| Molecular Weight: | 33.3 kDa |
|-------------------|--|
| UniProt: | P49615 |
| Pathways: | Cell Division Cycle, Regulation of Muscle Cell Differentiation, Synaptic Membrane, Regulation of |
| | Cell Size, Skeletal Muscle Fiber Development, Synaptic Vesicle Exocytosis |

Application Details

| Application Notes: | In addition to the applications listed above we expect the protein to work for functional studies as well. As the protein has not been tested for functional studies yet we cannot offer a guarantee though. |
|--------------------|---|
| Comment: | ALICE®, our Almost Living Cell-Free Expression System is based on a lysate obtained from Nicotiana tabacum c.v This contains all the protein expression machinery needed to produce even the most difficult-to-express proteins, including those that require post-translational modifications. During lysate production, the cell wall and other cellular components that are not required for |

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| Application Details | |
|---|---|
| | protein production are removed, leaving only the protein production machinery and the mitochondria to drive the reaction. During our lysate completion steps, the additional components needed for protein production (amino acids, cofactors, etc.) are added to produce something that functions like a cell, but without the constraints of a living system - all that's needed is the DNA that codes for the desired protein! |
| Restrictions: | For Research Use only |
| Handling | |
| Format: | Liquid |
| | |
| Buffer: | The buffer composition is at the discretion of the manufacturer. Standard Storage Buffer: PBS pH 7.4, 10 % Glycerol Might differ depending on protein. |
| Buffer: Handling Advice: | The buffer composition is at the discretion of the manufacturer. Standard Storage Buffer: PBS pH 7.4, 10 % Glycerol Might differ depending on protein. Avoid repeated freeze-thaw cycles. |
| Buffer: Handling Advice: Storage: | The buffer composition is at the discretion of the manufacturer. Standard Storage Buffer: PBS pH 7.4, 10 % Glycerol Might differ depending on protein. Avoid repeated freeze-thaw cycles. -80 °C |
| Buffer: Handling Advice: Storage: Storage Comment: | The buffer composition is at the discretion of the manufacturer. Standard Storage Buffer: PBS pH 7.4, 10 % Glycerol Might differ depending on protein. Avoid repeated freeze-thaw cycles. -80 °C Store at -80°C. |