

Datasheet for ABIN3137554 Parkin Protein (AA 1-464) (Strep Tag)



Overview

Quantity:	250 µg
Target:	Parkin (PARK2)
Protein Characteristics:	AA 1-464
Origin:	Mouse
Source:	Cell-free protein synthesis (CFPS)
Protein Type:	Recombinant
Purification tag / Conjugate:	This Parkin protein is labelled with Strep Tag.
Application:	SDS-PAGE (SDS), Western Blotting (WB), ELISA

Product Details

Brand:	AliCE®
Sequence:	MIVFVRFNSS YGFPVEVDSD TSILQLKEVV AKRQGVPADQ LRVIFAGKEL PNHLTVQNCD
	LEQQSIVHIV QRPRRRSHET NASGGDEPQS TSEGSIWESR SLTRVDLSSH TLPVDSVGLA
	VILDTDSKRD SEAARGPVKP TYNSFFIYCK GPCHKVQPGK LRVQCGTCKQ ATLTLAQGPS
	CWDDVLIPNR MSGECQSPDC PGTRAEFFFK CGAHPTSDKD TSVALNLITS NRRSIPCIAC
	TDVRSPVLVF QCNHRHVICL DCFHLYCVTR LNDRQFVHDA QLGYSLPCVA GCPNSLIKEL
	HHFRILGEEQ YTRYQQYGAE ECVLQMGGVL CPRPGCGAGL LPEQGQRKVT CEGGNGLGCG
	FVFCRDCKEA YHEGDCDSLL EPSGATSQAY RVDKRAAEQA RWEEASKETI KKTTKPCPRC
	NVPIEKNGGC MHMKCPQPQC KLEWCWNCGC EWNRACMGDH WFDV
	Sequence without tag. The proposed Strep-Tag is based on experience \ensuremath{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.

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Product Details

Characteristics:

Key Benefits:

- Made in Germany from design to production by highly experienced protein experts.
- 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 posttranslational 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

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Target Details	
Target:	Parkin (PARK2)
Alternative Name:	Prkn (PARK2 Products)
Background:	E3 ubiquitin-protein ligase parkin (EC 2.3.2.31) (Parkin RBR E3 ubiquitin-protein
	ligase),FUNCTION: Functions within a multiprotein E3 ubiquitin ligase complex, catalyzing the
	covalent attachment of ubiquitin moieties onto substrate proteins (PubMed:32047033,
	PubMed:29311685). Substrates include SYT11 and VDAC1 (PubMed:32047033,
	PubMed:29311685). Other substrates are BCL2, CCNE1, GPR37, RHOT1/MIRO1, MFN1, MFN2,
	STUB1, SNCAIP, SEPTIN5, TOMM20, USP30, ZNF746, MIRO1 and AIMP2 (By similarity).
	Mediates monoubiquitination as well as 'Lys-6', 'Lys-11', 'Lys-48'-linked and 'Lys-63'-linked
	polyubiquitination of substrates depending on the context (PubMed:32047033,
	PubMed:25474007). Participates in the removal and/or detoxification of abnormally folded or
	damaged protein by mediating 'Lys-63'-linked polyubiquitination of misfolded proteins such as
	PARK7: 'Lys-63'-linked polyubiquitinated misfolded proteins are then recognized by HDAC6,
	leading to their recruitment to aggresomes, followed by degradation (By similarity). Mediates
	'Lys-63'-linked polyubiquitination of a 22 kDa O-linked glycosylated isoform of SNCAIP, possibly
	playing a role in Lewy-body formation (By similarity). Mediates monoubiquitination of BCL2,
	thereby acting as a positive regulator of autophagy (By similarity). Protects against
	mitochondrial dysfunction during cellular stress, by acting downstream of PINK1 to coordinate
	mitochondrial quality control mechanisms that remove and replace dysfunctional
	mitochondrial components (PubMed:32047033, PubMed:25474007, PubMed:22082830,
	PubMed:24898855). Depending on the severity of mitochondrial damage and/or dysfunction,
	activity ranges from preventing apoptosis and stimulating mitochondrial biogenesis to
	regulating mitochondrial dynamics and eliminating severely damaged mitochondria via
	mitophagy (PubMed:32047033, PubMed:22082830, PubMed:24898855). Activation and
	recruitment onto the outer membrane of damaged/dysfunctional mitochondria (OMM) requires
	PINK1-mediated phosphorylation of both PRKN and ubiquitin (PubMed:25474007). After
	mitochondrial damage, functions with PINK1 to mediate the decision between mitophagy or
	preventing apoptosis by inducing either the poly- or monoubiquitination of VDAC1, respectively,
	polyubiquitination of VDAC1 promotes mitophagy, while monoubiquitination of VDAC1
	decreases mitochondrial calcium influx which ultimately inhibits apoptosis
	(PubMed:32047033). When cellular stress results in irreversible mitochondrial damage,
	promotes the autophagic degradation of dysfunctional depolarized mitochondria (mitophagy)
	by promoting the ubiquitination of mitochondrial proteins such as TOMM20, RHOT1/MIRO1,
	MFN1 and USP30 (PubMed:21753002). Preferentially assembles 'Lys-6'-, 'Lys-11'- and 'Lys-63'-
	linked polyubiquitin chains, leading to mitophagy (By similarity). The PINK1-PRKN pathway also

promotes fission of damaged mitochondria by PINK1-mediated phosphorylation which
promotes the PRKN-dependent degradation of mitochondrial proteins involved in fission such
as MFN2 (PubMed:24192653). This prevents the refusion of unhealthy mitochondria with the
mitochondrial network or initiates mitochondrial fragmentation facilitating their later
engulfment by autophagosomes (By similarity). Regulates motility of damaged mitochondria
via the ubiquitination and subsequent degradation of MIRO1 and MIRO2, in motor neurons, this
likely inhibits mitochondrial intracellular anterograde transport along the axons which probably
increases the chance of the mitochondria undergoing mitophagy in the soma (By similarity).
Involved in mitochondrial biogenesis via the 'Lys-48'-linked polyubiquitination of transcriptional
repressor ZNF746/PARIS which leads to its subsequent proteasomal degradation and allows
activation of the transcription factor PPARGC1A (By similarity). Limits the production of
reactive oxygen species (ROS) (By similarity). Regulates cyclin-E during neuronal apoptosis (By
similarity). In collaboration with CHPF isoform 2, may enhance cell viability and protect cells
from oxidative stress (PubMed:22082830). Independently of its ubiquitin ligase activity,
protects from apoptosis by the transcriptional repression of p53/TP53 (PubMed:19801972).
May protect neurons against alpha synuclein toxicity, proteasomal dysfunction, GPR37
accumulation, and kainate-induced excitotoxicity (By similarity). May play a role in controlling
neurotransmitter trafficking at the presynaptic terminal and in calcium-dependent exocytosis.
May represent a tumor suppressor gene (By similarity). {ECO:0000250 UniProtKB:060260,
ECO:0000269 PubMed:19801972, ECO:0000269 PubMed:21753002,
EC0:0000269 PubMed:22082830, EC0:0000269 PubMed:24192653,
EC0:0000269 PubMed:24898855, EC0:0000269 PubMed:25474007,
ECO:0000269 PubMed:29311685, ECO:0000269 PubMed:32047033}.

Molecular Weight:	51.6 kDa
UniProt:	Q9WVS6
Pathways:	Autophagy, Ubiquitin Proteasome Pathway
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

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	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!
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.
Handling Advice:	Avoid repeated freeze-thaw cycles.
Storage:	-80 °C
Storage Comment:	Store at -80°C.

Expiry Date:

12 months