

Datasheet for ABIN5711343 LKB1 Protein (AA 1-429, partial) (His-SUMO Tag)





Overview

Quantity:	100 µg
Target:	LKB1 (STK11)
Protein Characteristics:	AA 1-429, partial
Origin:	Human
Source:	Escherichia coli (E. coli)
Protein Type:	Recombinant
Purification tag / Conjugate:	This LKB1 protein is labelled with His-SUMO Tag.
Application:	SDS-PAGE (SDS)
Product Details	
Sequence:	MEVVDPQQLG MFTEGELMSV GMDTFIHRID STEVIYQPRR KRAKLIGKYL MGDLLGEGSY
	MEVVDPQQLG MFTEGELMSV GMDTFIHRID STEVIYQPRR KRAKLIGKYL MGDLLGEGSY GKVKEVLDSE TLCRRAVKIL KKKKLRRIPN GEANVKKEIQ LLRRLRHKNV IQLVDVLYNE
	GKVKEVLDSE TLCRRAVKIL KKKKLRRIPN GEANVKKEIQ LLRRLRHKNV IQLVDVLYNE
	GKVKEVLDSE TLCRRAVKIL KKKKLRRIPN GEANVKKEIQ LLRRLRHKNV IQLVDVLYNE EKQKMYMVME YCVCGMQEML DSVPEKRFPV CQAHGYFCQL IDGLEYLHSQ GIVHKDIKPG
	GKVKEVLDSE TLCRRAVKIL KKKKLRRIPN GEANVKKEIQ LLRRLRHKNV IQLVDVLYNE EKQKMYMVME YCVCGMQEML DSVPEKRFPV CQAHGYFCQL IDGLEYLHSQ GIVHKDIKPG NLLLTTGGTL KISDLGVAEA LHPFAADDTC RTSQGSPAFQ PPEIANGLDT FSGFKVDIWS
	GKVKEVLDSE TLCRRAVKIL KKKKLRRIPN GEANVKKEIQ LLRRLRHKNV IQLVDVLYNE EKQKMYMVME YCVCGMQEML DSVPEKRFPV CQAHGYFCQL IDGLEYLHSQ GIVHKDIKPG NLLLTTGGTL KISDLGVAEA LHPFAADDTC RTSQGSPAFQ PPEIANGLDT FSGFKVDIWS AGVTLYNITT GLYPFEGDNI YKLFENIGKG SYAIPGDCGP PLSDLLKGML EYEPAKRFSI
	GKVKEVLDSE TLCRRAVKIL KKKKLRRIPN GEANVKKEIQ LLRRLRHKNV IQLVDVLYNE EKQKMYMVME YCVCGMQEML DSVPEKRFPV CQAHGYFCQL IDGLEYLHSQ GIVHKDIKPG NLLLTTGGTL KISDLGVAEA LHPFAADDTC RTSQGSPAFQ PPEIANGLDT FSGFKVDIWS AGVTLYNITT GLYPFEGDNI YKLFENIGKG SYAIPGDCGP PLSDLLKGML EYEPAKRFSI RQIRQHSWFR KKHPPAEAPV PIPPSPDTKD RWRSMTVVPY LEDLHGADED EDLFDIEDDI
	GKVKEVLDSE TLCRRAVKIL KKKKLRRIPN GEANVKKEIQ LLRRLRHKNV IQLVDVLYNE EKQKMYMVME YCVCGMQEML DSVPEKRFPV CQAHGYFCQL IDGLEYLHSQ GIVHKDIKPG NLLLTTGGTL KISDLGVAEA LHPFAADDTC RTSQGSPAFQ PPEIANGLDT FSGFKVDIWS AGVTLYNITT GLYPFEGDNI YKLFENIGKG SYAIPGDCGP PLSDLLKGML EYEPAKRFSI RQIRQHSWFR KKHPPAEAPV PIPPSPDTKD RWRSMTVVPY LEDLHGADED EDLFDIEDDI IYTQDFTVPG QVPEEEASHN GQRRGLPKAV CMNGTEAAQL STKSRAEGRA PNPARKACSA

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Target:	LKB1 (STK11)
Alternative Name:	STK11 (STK11 Products)
Background:	Tumor suppressor serine/threonine-protein kinase that controls the activity of AMP-activated
	protein kinase (AMPK) family mbers, thereby playing a role in various processes such as cell
	metabolism, cell polarity, apoptosis and DNA damage response. Acts by phosphorylating the T
	loop of AMPK family proteins, thus promoting their activity: phosphorylates PRKAA1, PRKAA2,
	BRSK1, BRSK2, MARK1, MARK2, MARK3, MARK4, NUAK1, NUAK2, SIK1, SIK2, SIK3 and SNRK
	but not MELK. Also phosphorylates non-AMPK family proteins such as STRADA, PTEN and
	possibly p53/TP53. Acts as a key upstream regulator of AMPK by mediating phosphorylation
	and activation of AMPK catalytic subunits PRKAA1 and PRKAA2 and thereby regulates
	processes including: inhibition of signaling pathways that promote cell growth and proliferation
	when energy levels are low, glucose homeostasis in liver, activation of autophagy when cells
	undergo nutrient deprivation, and B-cell differentiation in the germinal center in response to
	DNA damage. Also acts as a regulator of cellular polarity by rodeling the actin cytoskeleton.
	Required for cortical neuron polarization by mediating phosphorylation and activation of BRSK
	and BRSK2, leading to axon initiation and specification. Involved in DNA damage response:
	interacts with p53/TP53 and recruited to the CDKN1A/WAF1 promoter to participate in
	transcription activation. Able to phosphorylate p53/TP53, the relevance of such result in vivo is
	however unclear and phosphorylation may be indirect and mediated by downstream
	STK11/LKB1 kinase NUAK1. Also acts as a mediator of p53/TP53-dependent apoptosis via
	interaction with p53/TP53: translocates to the mitochondrion during apoptosis and regulates
	p53/TP53-dependent apoptosis pathways. In vein endothelial cells, inhibits PI3K/Akt signaling
	activity and thus induces apoptosis in response to the oxidant peroxynitrite (in vitro). Regulates
	UV radiation-induced DNA damage response mediated by CDKN1A. In association with NUAK1
	phosphorylates CDKN1A in response to UV radiation and contributes to its degradation which
	is necessary for optimal DNA repair
Molecular Weight:	64.1 kDa
UniProt:	Q15831
Pathways:	AMPK Signaling, Carbohydrate Homeostasis, Regulation of Carbohydrate Metabolic Process,
	Warburg Effect
Application Details	
Application Notes:	Optimal working dilution should be determined by the investigator.
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Application Details

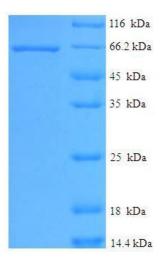
Restrictions:

For Research Use only

Handling

Format:	Liquid
Concentration:	0.1-2 mg/mL
Buffer:	20 mM Tris-HCl based buffer, pH 8.0
Storage:	-80 °C,4 °C,-20 °C
Storage Comment:	Store at -20°C, for extended storage, conserve at -20°C or -80°C. Repeated freezing and thawing is not recommended. Store working aliquots at 4°C for up to one week.

Images



SDS-PAGE		
Image 1.		
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