# antibodies

## Datasheet for ABIN5711419 TRIM21 Protein (AA 1-470, full length) (His-SUMO Tag)



Overview

Image

Quantity:	100 µg
Target:	TRIM21
Protein Characteristics:	full length, AA 1-470
Origin:	Mouse
Source:	Escherichia coli (E. coli)
Protein Type:	Recombinant
Purification tag / Conjugate:	This TRIM21 protein is labelled with His-SUMO Tag.
Application:	SDS-PAGE (SDS)

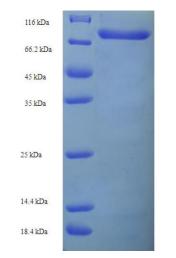
#### Product Details

Sequence:	MSPSTTSKMS LEKMWEEVTC SICLDPMVEP MSIECGHCFC KECIFEVGKN GGSSCPECRQ
	QFLLRNLRPN RHIANMVENL KQIAQNTKKS TQETHCMKHG EKLHLFCEED GQALCWVCAQ
	SGKHRDHTRV PIEEAAKVYQ EKIHVVLEKL RKGKELAEKM EMDLTMQRTD WKRNIDTQKS
	RIHAEFALQN SLLAQEEQRQ LQRLEKDQRE YLRLLGKKEA ELAEKNQALQ ELISELERRI
	RGSELELLQE VRIILERSGS WNLDTLDIDA PDLTSTCPVP GRKKMLRTCW VHITLDRNTA
	NSWLIISKDR RQVRMGDTHQ NVSDNKERFS NYPMVLGAQR FSSGKMYWEV DVTQKEAWDL
	GVCRDSVQRK GQFSLSPENG FWTIWLWQDS YEAGTSPQTT LHIQVPPCQI GIFVDYEAGV
	VSFYNITDHG SLIYTFSECV FAGPLRPFFN VGFNYSGGNA APLKLCPLKM
Purification:	SDS-PAGE
Purity:	> 90 %

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Target:	TRIM21
Alternative Name:	R052 (TRIM21 Products)
Background:	E3 ubiquitin-protein ligase whose activity is dependent on E2 enzymes, UBE2D1, UBE2D2,
	UBE2E1 and UBE2E2. Forms a ubiquitin ligase complex in cooperation with the E2 UBE2D2 that
	is used not only for the ubiquitination of USP4 and IKBKB but also for its self-ubiquitination.
	Component of cullin-RING-based SCF (SKP1-CUL1-F-box protein) E3 ubiquitin-protein ligase
	complexes such as SCF(SKP2)-like complexes. A TRIM21-containing SCF(SKP2)-like complex
	is shown to mediate ubiquitination of CDKN1B ('Thr-187' phosphorylated-form), thereby
	promoting its degradation by the proteasome. Monoubiquitinates IKBKB that will negatively
	regulates Tax-induced NF-kappa-B signaling. Negatively regulates IFN-beta production post-
	pathogen recognition by polyubiquitin-mediated degradation of IRF3. Mediates the ubiquitin-
	mediated proteasomal degradation of IgG1 heavy chain, which is linked to the VCP-mediated
	ER-associated degradation (ERAD) pathway. Promotes IRF8 ubiquitination, which enhanced the
	ability of IRF8 to stimulate cytokine genes transcription in macrophages. Plays a role in the
	regulation of the cell cycle progression. Enhances the decapping activity of DCP2. Exists as a
	ribonucleoprotein particle present in all mammalian cells studied and composed of a single
	polypeptide and one of four small RNA molecules. At least two isoforms are present in
	nucleated and red blood cells, and tissue specific differences in RO/SSA proteins have been
	identified. The common feature of these proteins is their ability to bind HY RNAs.2.
Molecular Weight:	70.14 kDa
UniProt:	Q62191
Application Details	
Application Notes:	Optimal working dilution should be determined by the investigator.
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

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### SDS-PAGE

Image 1.

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#### Images