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## Rabbit anti-human tripartite motif-containing 21 polyclonal Antibody Catalog Number: CSB-PA024457GA01HU

Synonym Names	52 kDa Ro protein,52 kDa ribonucleoprotein autoantigen Ro/SS-A,RING finger protein
	81,Ro(SS-A),Sjoegren syndrome type A antigen,SS-A,Tripartite motif-containing protein 21,TRIM21,RNF81,
	RO52, SSA1
Product type	Primary antibodies
Description	Rabbit polyclonal to TRIM21
Clonality	Polyclonal
Isotype	IgG
Reacts with	Human, Mouse, Rat; Other species are not tested. Please decide the specificity by homology.
Conjugate	Non-conjugated
Purity	Antigen Affinity Purified
Storage buffer	PBS with 0.1% sodium azide and 50% glycerol pH 7.3.
Storage	Shipped at 4°C Upon delivery aliquot and store at -20°C or -80°C. Avoid repeated freeze.
Form	Liquid
Raised in	Rabbit
	ELISA; Use at an assay dependent dilution.
Tested applications	WB: 1:200-1:2000
	IHC: 1:20-1:200 (Recommender dilutions)
Positive WB detected in	A549 cells, HeLa cells, mouse pancreas tissue, HT-1080 cells
Positive IHC detected in	Human lung cancer
	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
Function	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.
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	stem cell differentiation."Rigbolt K.T., Prokhorova T.A., Akimov V., Henningsen J., Johansen P.T.,
	Kratchmarova I., Kassem M., Mann M., Olsen J.V., Blagoev B.Sci. Signal. 4:RS3-RS3(2011).
	[2]"Quantitative phosphoproteomics reveals widespread full phosphorylation site occupancy during
References	mitosis."Olsen J.V., Vermeulen M., Santamaria A., Kumar C., Miller M.L., Jensen L.J., Gnad F., Cox J.,
	Jensen T.S., Nigg E.A., Brunak S., Mann M.
	Sci. Signal. 3:RA3-RA3(2010).
	[3]"Dynamic movements of Ro52 cytoplasmic bodies along microtubules."
	Tanaka M., Tanji K., Niida M., Kamitani T.Histochem. Cell Biol. 133:273-284(2010).
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