

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q6PCX9	TRI37_MOUSE	Mus musculus	E3 ubiquitin-protein ligase TRIM37	25.019509	S785	S454	34418053	MDEQSVESIAEVFRFCFICMEKLRDA RLCPHCCKLCCFSCIRRWLTEQRAQ CPHCRAPLQLRELVNCRWAEVETQ QLDTLQLCSLTKHEENEKDKCENH HEKLSVFCWTCKKCIHQCALWGG MHGGHTFKPLAEIYEQHVTKVNEEV AKLRRRLMELISLVQEVERNVEAVR NAKDERVREIRNAVEMMIARLDTQL KNKLITLMGQKTSLTQETELLESLLQ EVEHQLRSCSKSELISKSSSEILMMF QQVHRKPMASFVTTVPVPPDFTSELV PSYDSATFVLENFSTLRQRADPVYSP PLQVSGLCWRLKVYPDGNVVRGYY LSVFLELSAGLPETSKYEYRVMVH QSCNDPTKNHREFASDFEVGECWG YNRFFRLDLLANEGYLNRRQNDTVIL RFQVRSPTFFQKCRDQHWYITQLEA AQTYIQQINNLERLTIELSRTQKS RDLSPPDNHLSPQNDSPETRKKKA GSCSDMLLEGGPTCASVRETKEDD EEEKIQNEDYHHELSDGDLDLVLV EDEVNHLDGSSSSASSTATSNTEEN DIDEETMSGENDVEYNSMELEEGE LMEDAAAAGPPGSSHSYVGASSRMS RRTHLCSAATSSLLDIDPLILHLLDL KDRSSMENLWGLQPRPSASLLQPTA SYSRKDKDQRKQAMWRVPSDLKM LKRLKTQMAEVRCKMTDVKTTLSDI KGSSVASTDMQTNLFCADQAALTTC GPENSGRLQDLGMELLAKSSVAGCY IRNPTNKKNSPKSARAIAGSLSLRRA VDSGENSRKGDQVLAEGSSGSSQ SGSRHSSPRALTHGIIGDLLPKSEDR QCKALDSAVVVAVFNGLPTVEKRR KMVTLGTNAKGGRLGEMQMADLES HSEAGEVQPTLPEGASAAPEEGMSS DSDIECDTENEQEHEHTSMGAFND PFLAQPDEDSHSSFPDGEQIDPEN LHFNPEDEGGGR	None	None	None	None	None	None	None		

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