

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
0 O94972	TRIM37_HUMAN	Homo sapiens	E3 ubiquitin-protein ligase TRIM37	26.21016	T195	S454	29351928	MDEQSVESIAEVFRFCFICMEKLRDA RLCPHCSKLCFCFSCIRRWLTEQRAQ CPHCRAPLQLRELVNCRWAEVETQ QLDTLQLCSLTKHEENEKDKCENH HEKLSVFCWTKCKCICHQCALWGG MHGGHTFKPLAEIYEQHVTKVNEEV AKLRRRLMELISLVQEVERNVEAVR NAKDERVREIRNAVEMMIARLDTQL KNKLITLMGQKTSLSLQTELESLLO EVEHQLRSCSKSELISKSSEILMMF QQVHRKPMASFTVTPVPPDFTSELV PSYDSATFVLENFSTLRORADPVYSP PLQVSGLCWRLKVYPDGNVVRGYY LSVFLELSAGLPETSKYEYRVMVH QSCNDPTKNIREFASDFEVGECWG YNRFFRLDLLANEGYLNPNQNDTVIL RFQVRSPTFFQKSRDQHWYITQLEA AQTSYIQQINNLERLTIELSRTQKS RDLSPDNHLSQPNDDALETRAKKS ACSDMLLEGGPTTASVREAKEDEED EELIQNEDYHHELSDGDLDLVLVE DEVNQLDGSSSSASSTATSNTTEEND IDEETMSGENDVEYNNMELEGEL MEDAAAAGPAGSSHGYVSSSRISR RTHLCSAATSLLDIDPLILIHLLDLK DRSSIENLWGLQPRPPASLLQPTASY SRKDKDQRKQAMWRVPSDLKMLK RLKTQMAEVRMKTQVKNLSEIKS SSAASGDMQTSLSADQAALAACGT ENSGRLQDLGMELLAKSSVANCYIR NSTNKKSNPKPARSSVAGSLSLRR AVDPGENSRSKGDCQTLSEKSPGSS QSGSRHSSPRALIHGSIGDILPKTED RQCKALDSDAVVAVVFGSLPAVEKR RKMVTLGANAKGGHLEGLQMTDLE NNSETGELQPVLPEGASAAPEEGMS SDSDIECDTENEQEHTSVGGFHD SFMVMTQPPDEDTHSSFPDGEQIGP EDLSFNTDENSGR	True	False	5.0	3.437	1.227	1.004	0.555	1.067	1.184