	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	gol app
0	Q14669	TRIPC_HUMAN	Homo sapiens	E3 ubiquitin-protein ligase TRIP12	22.849954	T387;T758;T764 (Q14669-2)	S12;S77;S85;S100;S310;S312;S942;S991;S997;S1016;S1030;S1317;S1322;S1329;S1376;T1377;S1427	34725712;36240223;28510447;38253038	MSNRPNNNPGGSLRRSQRNTAGAQ PQDDSIGGRSCSSSSAVIVPQPEDPD RANTSERQKTGQVPKKDNSRGVKRS ASPDYNRTNSPSSAKKPKALQHTES PSETNKPHSKSKKRHLDQEQQLKSA QSPSTSKAHTRKSGATGGSRSQKRK RTESSCVKSGSGSESTGAEERSAKPT KLASKSATSAKAGCSTITDSSSAAST SSSSSAVASASSTVPPGARVKQGKD QNKARRSRSASSPSPRRSSREKEQS KTGGSSKFDWAARFSPKVSLPKTKL SLPGSSKSETSKPGPSGLQAKLASLR KSTKKRSESPPAELPSLRRSTRQKTT GSCASTSRRGSGLGKRGAAEARRQE KMADPESNQEAVNSSAARTDEAPQ GAAGAVGMTTSGESESDDSEMGRL QALLEARGLPPHLFGPLGPRMSQLF HRTIGSGASSKAQQLLQGLQASDES QQLQAVIEMCQLLVMGNEETLGGF PVKSVVPALITLLQMEHNFDIMNHA CRALTYMMEALPRSSAVVVDAIPVFL EKLQVIQCIDVAEQALTALEMLSRR HSKAILQAGGLADCLLYLEFFSINAQ RNALAIAANCCQSITPDEFHFVADSL PLLTQRLTHQDKKSVESTCLCFARL VDNFQHEENLLQQVASKDLLTNVQ QLLVVTPPILSSGMFIMVVRMFSLM CSNCPTLAVQLMKQNIAETLHFLLC GASNGSCQEQIDLVPRSPQELYELTS LICELMPCLPKEGIFAVDTMLKKGN AQNTDGAIWQWRDDRGLWHPYNRI DSRIIEQINEDTGTARAIQRKPNPLA NSNTSGYSESKKDDARAQLMKEDP ELAKSFIKTLFGVLYEVYSSSAGPAV RHKCLRAILRIIYFADAELLKDVLKN HAVSSHIASMLSSQDLKIVVGALQM AEILMQKLPDIFSVYFRREGVMHQV KHLAESESLLTSPPKACTNGSGSMG STTSVSSGTATAATHAAADLGSPSLQ HSRDDSLDLSPQGRLSDVLKRKRLP KRGPRRPKYSPPRDDDKVDNQAKSP TTTQSPKSSFLASLNPKTWGRLSTQ SNSNNIEPARTAGGSGLARAASKDTI SNNREKIKGWIKEQAHKFVERYFSS ENMDGSNPALNVLQRLCAATEQLN LQVDGGAECLVEIRSIVSESDVSSFEI QHSGFVKQLLLYLTSKSEKDAVSREI RLKRFLHVFFSSPLPGEEPIGRVEPV GNAPLLALVHKMNNCLSQMEQFPV KVHDFPSGNGTGGSFSLNRGSQALK FFNTHQLKCQLQRHPDCANVKQWK GGPVKIDPLALVQAIERYLVVRGYGR VREDDEDSDDDGSDEEIDESLAAQF LNSGNVRHRLQFYIGEHLLPYNMTV YQAVRQFSIQAEDERESTDDESNPL GRAGIWTKTHTIWYKPVREDEESNK DCVGGKRGRAQTAPTKTSPRNAKKH DELWHDGVCPSVSNPLEVYLIPTPPE NITFEDPSLDVILLLRVLHAISRYWYY LYDNAMCKEIIPTSEFINSKLTAKAN RQLQDPLVIMTGNIPTWLTELGKTC PFFFPFDTRQMLFYVTAFDRDRAMQ RLLDTNPEINQSDSQDSRVAPRLDR KKRTVNREELLKQAESVMQDLGSSR AMLEIQYENEVGTGLGPTLEFYALVS QELQRADLGLWRGEEVTLSNPKGS QEGTKYIQNLQGLFALPFGRTAKPA HIAKVKMKFRFLGKLMAKAIMDFRL VDLPLGLPFYKWMLRQETSLTSHDL FDIDPVVARSVYHLEDIVRQKKRLEQ DKSQTKESLQYALETLTMNGCSVED LGLDFTLPGFPNIELKKGGKDIPVTI HNLEEYLRLVIFWALNEGVSRQFDS FRDGFESVFPLSHLQYFYPEELDQLL CGSKADTWDAKTLMECCRPDHGYT HDSRAVKFLFEILSSFDNEQQRLFL QFVTGSPRLPVGGFRSLNPPLTIVRK TFESTENPDDFLPSVMTCVNYLKLP DYSSIEIMREKLLIAAREGQQSFHLS	None	None	None	None	None	None	Nor