

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus
Q6WKZ4	RFIP1_HUMAN	Homo sapiens	Rab11 family-interacting protein 1	30.584871	NaN	S184;S202;S206;S234;S300;S315;S339;S341;S343;S345;S356;S357;S382;S435;S477;S529;S545;S758;S1135	28411811;23301498;40914422;37217939	MSLMVVSAGRGLGAVVSPTHVQVTVLQARGLRAKPGGTSDAYAVIQVGEKYATSVSERSLGAAPVWREATFELP SLLSSGPAAAAATLQTLVLRALLGLDKFLGRAEVDLRDLHRDQGRRTQWYKFKKSPGKDKKERGEIEVDIQFMR NNMTASMFDSLMDKSRNPFQGLKDKIKGKNKDSGSDTASAIIPSTTPSV DSDDES VVKDKKKKSKIKTLLSKSNLQKTPLSQSMSVLPSTKPEKVLRLPG DFQSQWDEDDNEDESSASDVMSHKRTASTDLKQLNQVNFPLPKKEGLS FLGGRLSKNDVLSRSNVCINGNHVYLEQPEAKGEIKDSSPSSSPSPKGFRRK KHLFSSSTENLAAGSWKEPAEGGGLSDRQLESSTKDSLKSMTLPSYRPAP LVSGDLRENMAPANSEATKEAKESK KPESRRSSLLSLMTGKDVAKGSEG ENPLTVPGREKEGMLMGVKGEDA SGPAEDLVRREKDTAAVVSROGSS LNLFDVQITEPEAEPEKSEPRPPIS SPRAPQTRAVKPRLEVPPEAQPTRL PSPTDSSSLPPLPSSSGQASVPEL GHGADTQSSSPSVFSSLSPIAAPIS TSTPIESWPLVDRGQAKSEGPPLLPK AELQTESLTPVPSGSSALGSLFKQP SFPANKGTEDSLMGRTRGTGTEKNT SSELEESLPEQPETGRQEEELPRFP CKKQDYSPSSGEAQEVFALSSSD GAVSPVGELAAGGDRDLESQAGSLV ESKARDAEEVAPPLPMGASVPSIDS MMRKLEEMGLNLRKDQKTKKRV FSEQLFTEEAVAGAALLVEGHSSCP QELNPAWSVAGNASDGEPPESPHA EDSERESVITPGPATCGAPASPADH LLPSQEEFSFVPMSEASSAKDTPL FRMEGEDALVTQYQSKASDHEGLLS DPLSDLQLVDFKSPIMADLNLSLPS IPEVASDDERIDQVEDDGDQVEDDG ETAKSSTLDIGALSLGLVVPCEPERGK GPSGEADRLVLGEGLCDFRLQAPQA SVTAPSEQTTEFGIHKPHLGSSSLD KQLPGPSGGEEKPMGNPSPSPPPG TSLDNPVPSPPSEIFPVTHSFPSSA HSDTHHTSTAESQKATAEGSAGRV ENFGKRKPLLQAWVSPSETHPVSAQ PGAGTGSAKHRLHPVKPMNAMATK VANCSLGTATIISENLNNEVMMKKY SPSDPAFAYAQLTHDELQVLKQKE TISKKEFQVRELEDYIDNLLVRVMEE TPNILRIPTQVGGKAGKM	True	False	4.774	1.605	1.027	1.17	1.663