	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellula region
0	Q8IWW6	RHG12_HUMAN	Homo sapiens	Rho GTPase-activating protein 12	26.75988	NaN	S165;S176;S201;S213;S215;T230;T231;S240;T243;S592	29485866	MKMADRSGKIIPGQVYIEVEYDYEYE AKDRKIVIKQGERYILVKKTNDDWW QVKPDENSKAFYVPAQYVKEVTRKA LMPPVKQVAGLPNNSTKIMQSLHLQ RSTENVNKLPELSSFGKPSSSVQGT GLIRDANQNFGPSYNQGQTVNLSLD LTHNNGKFNNDSHSPKVSSQNRTR SFGHFPGPEFLDVEKTSFSQEQSCD SAGEGSERIHQDSESGDELSSSSTE QIRATTPPNQGRPDSPVYANLQELKI SQSALPPLPGSPAIQINGEWETHKDS SGRCYYYNRGTQERTWKPPRWTRD ASISKGDFQNPGDQELLSSEENYYST SYSQSDSQCGSPPRGWSEELDERG HTLYTSDYTNEKWLKHVDDQGRQY YYSADGSRSEWELPKYNASSQQQRE IIKSRSLDRRLQEPIVLTKWRHSTIVL DTNDKESPTASKPCFPENESSPSSP KHQDTASSPKDQEKYGLLNVTKIAE NGKKVRKNWLSSWAVLQGSSLLFT KTQGSSTSWFGSNQSKPEFTVDLKG ATIEMASKDKSSKKNVFELKTRQGT ELLIQSDNDTVINDWFKVLSSTINNQ AVETDEGIEEEIPDSPGIEKHDKEKE QKDPKKLRSFKVSSIDSSEQKKTKK NLKKFLTRRPTLQAVREKGYIKDQVF GSNLANLCQRENGTVPKFVKLCIEH VEEHGLDIDGIYRVSGNLAVIQKLRF AVNHDEKLDLNDSKWEDIHVITGAL KMFFRELPEPLFTFNHFNDFVNAIK QEPRQRVAAVKDLIRQLPKPNQDTM QILFRHLRRVIENGEKNRMTYQSIAI VFGPTLLKPEKETGNIAVHTVYQNQI VELILLELSSIFGR	None	None	None	None	None	None	None	None	None