	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membr
0	O14559	RHG33_HUMAN	Homo sapiens	Rho GTPase-activating protein 33	20.111456	S67;S74;T593;S955;S978	S8;S570;S636;S727;T1169	30379171;29351928	MVARSTDSLDGPGEGSVQPLPTAGG PSVKGKPGKRLSAPRGPFPRLADCA HFHYENVDFGHIQLLLSPDREGPSL SGENELVFGVQVTCQGRSWPVLRSY DDFRSLDAHLHRCIFDRRFSCLPEL PPPPEGARAAQMLVPLLLQYLETLSG LVDSNLNCGPVLTWMELDNHGRRL LLSEEASLNIPAVAAAHVIKRYTAQA PDELSFEVGDIVSVIDMPPTEDRSW WRGKRGFQVGFFPSECVELFTERPG PGLKADADGPPCGIPAPQGISSLTSA VPRPRGKLAGLLRTFMRSRPSRQRL RQRGILRQRVFGCDLGEHLSNSGQD VPQVLRCCSEFIEAHGVVDGIYRLSG VSSNIQRLRHEFDSERIPELSGPAFL QDIHSVSSLCKLYFRELPNPLLTYQL YGKFSEAMSVPGEEERLVRVHDVIQ QLPPPHYRTLEYLLRHLARMARHSA NTSMHARNLAIVWAPNLLRSMELE SVGMGGAAAFREVRVQSVVVEFLLT HVDVLFSDTFTSAGLDPAGRCLLPR PKSLAGSCPSTRLLTLEEAQARTQGR LGTPTEPTTPKAPASPAERRKGERGE KQRKPGGSSWKTFFALGRGPSVPRK KPLPWLGGTRAPPQPSGSRPDTVTL RSAKSEESLSSQASGAGLQRLHRLR RPHSSSDAFPVGPAPAGSCESLSSSS SSESSSSESSSSSSESSAAGLGALSG SPSHRTSAWLDDGDELDFSPPRCLE GLRGLDFDPLTFRCSSPTPGDPAPPA SPAPPAPASAFPPRVTPQAISPRGPTS PASPAALDISEPLAVSVPPAVLELLGA GGAPASATPTPALSPGRSLRPHLIPL LLRGAEAPLTDACQQEMCSKLRGAQ GPLGPDMESPLPPPPLSLLRPGGAPP PPPKNPARLMALALAERAQQVAEQQ SQQECGGTPPASQSPFHRSLSLEVG GEPLGTSGSGPPPNSLAHPGAWVPG PPPYLPRQQSDGSLLRSQRPMGTSR RGLRGPAQVSAQLRAGGGGRDAPEA AAQSPCSVPSQVPTPGFFSPAPRECL PPFLGVPKPGLYPLGPPSFQPSSPAP VWRSSLGPPAPLDRGENLYYEIGASE GSPYSGPTRSWSPFRSMPPDRLNAS YGMLGQSPPLHRSPDFLLSYPPAPS CFPPDHLGYSAPQHPARRPTPPEPLY VNLALGPRGPSPASSSSSSPPAHPRS RSDPGPPVPRLPQKQRAPWGPRTPH RVPGPWGPPEPLLLYRAAPPAYGRG GELHRGSLYRNGGQRGEGAGPPPPY PTPSWSLHSEGQTRSYC	None	None	None	None	None	None	None	None