	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	ex re
0	Q6P4F7	RHGBA_HUMAN	Homo sapiens	Rho GTPase-activating protein 11A	26.417971	S841	S285;T306;S316;S318;T323;S339;S340;S484;T508;S582;S585;S638;S675;S847;T866;S868	35254053;40596516	MWDQRLVRLALLQHLRAFYGIKVKG VRGQCDRRRHETAATEIGGKIFGVPF NALPHSAVPEYGHIPSFLVDACTSLE DHIHTEGLFRKSGSVIRLKALKNKVD HGEGCLSSAPPCDIAGLLKQFFRELP EPILPADLHEALLKAQQLGTEEKNK ATLLLSCLLADHTVHVLRYFFNFLR NVSLRSSENKMDSSNLAVIFAPNLL QTSEGHEKMSSNTEKKLRLQAAVV QTLIDYASDIGRVPDFILEKIPAMLGI DGLCATPSLEGFEEGEYETPGEYKR KRRQSVGDFVSGALNKFKPNRTPSI TPQEERIAQLSESPVILTPNAKRTLPV DSSHGFSSKKRKSIKHNFNFELLPS NLFNSSSTPVSVHIDTSSEGSSQSSL SPVLIGGNHLITAGVPRRSKRIAGKK VCRVESGKAGCFSPKISHKEKVRRS LRLKFNLGKNGREVNGCSGVNRYE SVGWRLANQQSLKNRIESVKTGLLF SPDVDEKLPKKGSEKISKSEETLLTP ERLVGTNYRMSWTGPNNSSFQEVD ANEASSMVENLEVENSLEPDIMVEK SPATSCELTPSNLNNKHNSNITSSPL SGDENNMTKETLVKVQKAFSESGS NLHALMNQRQSSVTNVGKVKLTEP SYLEDSPEENLFETNDLTIVESKEKY EHHTGKGEKCFSERDFSPLQTQTFN RETTIKCYSTQMKMEHEKDIHSNMP KDYLSKQEFSSDEEIKKQQSPKDKL NNKLKENENMMEGNLPKCAAHSK DEARSSFSQQSTCVVTNLSKPRPMR IAKQQSLETCEKTVSESSQMTEHRK VSDHIQWFNKLSLNEPNRIKVKSPL KFQRTPVRQSVRRINSLLEYSRQPTG HKLASLGDTASPLVKSVSCDGALSS CIESASKDSSVSCIKSGPKEQKSMSC EESNIGAISKSSMELPSKSFLKMRK HPDSVNASLRSTTVYKQKILSDGQV KVPLDDLTNHDIVKPVVNNNMGISS GINNRVLRRPSERGRAWYKGSPKHP IGKTQLLPTSKPVDL	True	False	4.668	5.0	1.109	False	False	False	Fa