	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	extracellular region
0	Q69ZH9	RHG23_MOUSE	Mus musculus	Rho GTPase-activating protein 23	29.885389	NaN	S361;S372;S421;S515;S579;S607;S619;T652;S655;S658;S673	22517741	MNGVAFCLVGIPPRPEPRPPQLPLGP RDGCSSGRPLPWPGPRTLLLRKSLQ DGFGFTLRHFIVYPPESAVHCILKEE ENGGRGGGPSPRHRLEPMDTIFVKN VKDGGPAHRAGLRTGDRLVKVNGE SIIGKTYSQVIGLIQNSDDTLELSIMP KDEDILQLAYSQDAYLKGNEPYSGE ARSIPEPPPLCYPRKTYAPPTRAPAW ATMVPEPISALPPDPRSPAAWSDPGS RVPSATRAHLDNSSLGMSQPRPSPG AFPHLPSESRTPRAFPEPGSRVLPSR LECQQALSHWLSNQIPRRAGERRCP AMPPRARSASQDRLEDVTTHHPWP CSTSQDALSQLGQEGWHRARSDDY LSRATRSAEALGPGALVSPRLERCG WASQRPSARTSACPSRDLTQAPPPS GLQGLDDIGYIGYRSYSPSFQRRTGL LHALSFRDSPFGGLPTFSLAQSPASF PPEASEPPRVVRPDPSTRALEPPAED HRDEVVLRQKPPTGRKVQLTPARQ MNLGFGDESPEPEARGERLGRKVAP LATTEDSLASIPFIDEPTSPSIDLQAK HVPASAVVSSAMNSAPVLGTSPSSP TFTFALSRHYSQDCSSIKAGRRSSYL LAITTERSKSCDDGLNTFRDEGRVL RRLPSRVPSLRVLRSFFTDGSLDSW GTSEDADAPSKRHSTSDLSDATFSDI RREGWLYYKQILTKKGKKAGGGLRQ WKRVYAVLRARSLSLSKERREPGPA AAGAAAAGAGEDEAAPVCIGSCLVDI SYSETKRRHVFRLTTADFCEYLFQAE DRDDMLGWIRAIRENSRAEGEDPGC ANQALISKKLNDYRKVSHSSGPKAD SSPKGSRGLGGLKSEFLKQTAVRGL RTQEQPPGSKEDSVAAPKTPWGINII KKNKKAAPRAFGIRLEECQPATENQ RVPLIVAACCRIVEARGLESTGIYRVP GNNAVVSSLQEQLNRGPSDINLQDE RWQDLNVISSLLKAFFRKLPEPLFTD DKYNDFIEANRIEDSRERMKTLRKLI RDLPGHYYETLKFLVGHLKTIADHSE KNKMEPRNLALVFGPTLVRTSEDN MTDMVTHMPDRYKIVETLIQHSDW FFSDDEDKGERTPVDDKEPQSVPNI EYLLPNIGRTVPPSDPGSDSTTCSSA KSKGSWVPKKEPYAREMLAISFISAV NRKRKKRREARGLGSSTDDDSEQEA HKAAVGTQEPPEGQLPGPAAEEAPG RLSPPTAPDERPAADTRSIVSGYSTL STMDRSVCSGTGGRRAGAGDEADD ERSELSHVETDTEGGAGPGGRLSRR PSFSSHHLMPCDTLARRRLSRSRAE AEGPGAGTARACPRGPEPPGSASSS SQESLRPPAAAPALGSRPSRVEALRL RLRGTADDMLAVRLRRPLSPETRRR RSSWRRHTVVVQSPLTDLNFNEWK ELGGGGPQESVGVRPHSDNKDSGL SSLESTKARASSAASLPSGDLGALQG LPQRRSAAARLHQCL	None	None	None	None	None	None	None	None	None