	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	e: re
0	Q9H2M9	RBGPR_HUMAN	Homo sapiens	Rab3 GTPase-activating protein non-catalytic subunit	28.070169	NaN	S39;S450;T901;S916;S978	23301498;40914422;38665916	MACSIVQFCYFQDLQAARDFLFPHL REEILSGALRRDPSKSTDWEDDGWG AWEENEPQEPEEEGNTCKTQKTSW LQDCVLSLSPTNDLMVIAREQKAVF LVPKWKYSDKGKEEMQFAVGWSGS LNVEEGECVTSALCIPLASQKRSSTG RPDWTCIVVGFTSGYVRFYTENGVL LLAQLLNEDPVLQLKCRTYEIPRHPG VTEQNEELSILYPAAIVTIDGFSLFQS LRACRNQVAKAAASGNENIQPPPLA YKKWGLQDIDTIIDHASVGIMTLSPF DQMKTASNIGGFNAAIKNSPPAMSQ YITVGSNPFTGFFYALEGSTQPLLSH VALAVASKLTSALFNAASGWLGWKS KHEEEAVQKQKPKVEPATPLAVRFG LPDSRRHGESICLSPCNTLAAVTDDF GRVILLDVARGIAIRMWKGYRDAQIG WIQTVEDLHERVPEKADFSPFGNSQ GPSRVAQFLVIYAPRRGILEVWSTQQ GPRVGAFNVGKHCRLLYPGYKIMGL NNVTSQSWQPQTYQICLVDPVSGSV KTVNVPFHLALSDKKSERAKDMHLV KKLAALLKTKSPNLDLVETEIKELILD IKYPATKKQALESILASERLPFSCLRN ITQTLMDTLKSQELESVDEGLLQFC ANKLKLLQLYESVSQLNSLDFHLDT PFSDNDLALLLRLDEKELLKLQALLE KYKQENTRTNVRFSDDKDGVLPVKT FLEYLEYEKDVLNIKKISEEEYVALGS FFFWKCLHGESSTEDMCHTLESAG LSPQLLLSLLLSVWLSKEKDILDKPQ SICCLHTMLSLLSKMKVAIDETWDS QSVSPWWQQMRTACIQSENNGAAL LSAHVGHSVAAQISNNMTEKKFSQT VLGADSEALTDSWEALSLDTEYWKL LLKQLEDCLILQTLLHSKGNTQTSKV SSLQAEPLPRLSVKKLLEGGKGGIAD SVAKWIFKQDFSPEVLKLANEERDA ENPDEPKEGVNRSFLEVSEMEMDL GAIPDLLHLAYEQFPCSLELDVLHAH CCWEYVVQWNKDPEEARFFVRSIE HLKQIFNAHVQNGIALMMWNTFLV KRFSAATYLMDKVGKSPKDRLCRRD VGMSDTAMTSFLGSCLDLLQILMEA DVSRDEIQVPVLDTEDAWLSVEGPIS IVELALEQKHIHYPLVEHHSILCSILY AVMRFSLKTVKPLSLFDSKGKNAFF KDLTSIQLLPSGEMDPNFISVRQQFL LKVVSAAVQAQHSATKVKDPTEEAT PTPFGKDQDWPALAVDLAHHLQVS EDVVRRHYVGELYNYGVDHLGEEAI LQVHDKEVLASQLLVLTGQRLAHAL LHTQTKEGMELLARLPPTLCTWLKA MDPQDLQNTEVPIATTAKLVNKVIEL LPEKHGQYGLALHLIEAVEAISLPSL	True	True	4.641	2.284	1.096	4.288	1.397	4.443	0.