

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q3V0G7	GARL3_MOUSE	Mus musculus	GTPase-activating Rap/Ran-GAP domain-like protein 3	36.530658	T901;S905	S68;S449;S455;T851;S6	22517741;36852467	MDPLTKGSCGSQLAQTLLWKASSL SFGIQPLQTWPTKDPPELESQVNLV SEDLGCRRGDFSRKHYGSVELLIS DADGAIQRAGRFRVENGSTDESAAA LPGTWRRTDVHLENPEYHTRWYFK YFLGQVHQNYIGNDAEKSPFFLSVT LSDQNNQRVPOYRAILWRKTGTQKI CLPYSPKTLVSVKILSAMNLDKFEK GPREIFHPEIQDLLVLEEQEGSVNF KFGVLFADGQQLTDDDEMFSNEIGSE AFQKFLNLLGDITLKGWTGYRGGL DTKNNTTGINSVYTVYQGHVEMFHV STMLPYSKENRQQVERKRHIGNDIV TIVFOEGEESPFAFKPSMIRSHFTHI FALVRYDQQNDNYRLKIFSEESVPLF GPPLPSPVFTDHDQEFRDFFLVKLIN GEKATLETPTFAQKRRRLDMLIRSL YQDLMPDLHKNMLNRRSFSDVLP SPKSARKKEEARQAEFVRIGQALKK SIVRGDAPSSLAASGMCKKEPWEPO CFCCNFPEHAVCADPWGQALLVST DAGVLLVDDDLPSVPVFDRTLVPKQI HVLETDLLVLRADKKGKDARLFVFR LSAVQKGLDGRQTGRSRSDCRENKL EKTGCHLYAINTHHSRELRIVVAIR NKLLITRKHKPHNKPSGVPVGSLLSPL SESPVEEFQYIREICLDCSPAVMALV DGPTEESDNLICVAYRHQFDVVNES TGEAFRLHHVEANKVNFVAIDVYE DGEAGLLLCYNYSCYKKVCPFNNG SFLQPSASDFQFCWNQAPYAIVCA FPYLLAFTTDSMEIRLVVNGNLVHT AVVPQLQLVASRDIYFTAATTVHEG SSGGSSKGASAHTSPOTPPARDTPLF PSSLGEGEIQSKNLYKIPLRNLVGRSI ERPLKSPVSKVITPPTSIGLVAAIP VTHSLSLSRMEIKEIASRTRRELLGL SDDGGTKTEGAPRAKSKTRKREES QGGPKPETVRSASSDRIPSGILESPA SEANPEGHNHWASSEQDAGVDKEG SPGSGSPFQLMASSEDIIDLK	None	None	None	None	None	None	None	None	

0