

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	ex re
Q3UHD9	AGAP2_MOUSE	Mus musculus	Arf-GAP with GTPase, ANK repeat and PH domain-containing protein 2	34.340724	NaN	S113;S128;S149;S632;S744;S746;S802;S921;S979;S1172	22517741;33300544	MSRGAGALQRRRTTYLISLTLVKLES VPPPPSPSAAAAGAPGARGSEPRDP GSPRGSEEPGKKRHERLFHRQDAL WISTSSAGTGGAEPPALSPAPASPAR PVSPAPGRRLSLWAAPPGLSGGLS PDSKPGGAPSSRRPILLSPPSWGGP EPEGRTGGVPGSSSPHPTGTSRRL KVAPPPAPKPFKTVTTSGAKAGGK GAGSRLSWPESEKPRVKGSKSTAG TGASAAAAGGGSAAVTTSGGVVGAG AGTRGKLSPRKSKTLDNSDLHPG PSAGSPPLTVPAPVPAATVATSTQP LGPAPPITLPPAPGLKRGREGGRAS TRDRKMLKFISGIFTKSTGGPPGPGP LPGPQLSSSSGSRELLGAELRTSPK AVVNSQEWTLRSRSEPELRLGVLGDV RSGKSLIHRFLTGSYQVLEKIESEQ YKKEMLVDGQTHLVLIREEAGAPDA KFSGWADAVIFVFSLEDESSQAVS RLHGQLSSLRGEGRGGLALALVGTQ DRISASSPRVGDARARALCTDMKR CSYYETCATYGLNVDVRFQVAQKV VTLRKQQLLAACKSLPSSPSHSAAS TPVAGQASNGGHTSDYSSSLPSSPN VGHRELRAEAAVAGLSTPGSLHRA AKRRTSLFANRRGSDAEKRSLSRG ETTGSGRAIPIKQSFLLKRSGNLKN EWKKKYVTLSSNGFLYHPHSINDYIH STHGKEMDLLRTTVKVPKRPRAI SAFGPSASINGLVKDMSTVQMGEGP EASTPMPSPPSPSSLQLPTDQTSKH LLKPPDRNLARALSTDCTPSGDLSPS REPPSPMVKQRKRLSTPSKTEG SAVQAEAKRKMWKLKSFGLRNIYK AEENFEFLIVSSTGQTWHFEEASFE ERDAWVQAIESQILASLQCESSKVK LRTDSQSEAVAIQAIRNAKGNSTCVD CGAPNPTWASNLGALICIECSGIHR NLGTHLSRVRSLDLDDWPRELTLVL TAIGNDTANRVWESDTRGRAKPTRD SSREERESWIRAKYEQLLFLAPLGT EEPLGRQLWAAVEAQDVAAVLLLLA HARHGPLDTSVEDPQLRSLPLHAAE LAHVVTQLLLWYGADVAARDAQGR TALFYARQAGSQLCADILLQHGCPG EGGSTATTPSAATTPSITATPSPPRRS SAASLGRVDITIALV	True	False	3.691	4.505	2.025	0.94	1.192	4.284	1.1