

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	extracellular region
D4ABA9	D4ABA9_RAT	Rattus norvegicus	NaN	23.439218	NaN	NaN	38843836	MADAQMQVVHTPTIQMRTGEDLPL PPPSALEGLPPPPKESFSKFQQRQ ANELRRLYKHIHPELRKNLEEA DLAKVLGSEEPTEGDVQCMRWIFEN WSLDAIGDHERPAGKEPVSGGNVQA TSRKFEESFTNSSDLEPGPQLSGG DVRATRQLFETKSLDALTGQEEATQ MTMKEPAVSGDVQGTTRKLFETKPLD MLGSCPSIQDQSPLELRSEIQELKGD VKKTVKLQFTEPLCAIQDAEGAIHEV KAACREEIQSNVRSARWLFETRPL DAFNQDPSQVRVIRGISLEEGALPDV SATRWIFETQPLDAIREILVDEKDFQ PSPDLIPPGPDVQHQHRLFETCALDT LKGERETEAEAPAKEEVIPGDVRS WLFETKPLDAFRDQVQVGHQRVD HQKGEQVIECLSSNGTSALPLSQG VPQNNGPKGDVKTFFKNLFETLPLDS IGQGEPSACGNINRGQRTDSTGLSE GADAPVYAMQDSRGLHALTSVSRE QVVGSDVQGYKWMFETQPLDRLGR SPSTVDVVRGITRQEVVAGDVGTR WLFETQPLEMIHQEQKREEEEG RGPGGPPPELPQKGDVQVIRWLFET YPMSELAEKQSEVTGPVSKAEAQ CTWMFGSQSLNQAEGSGEQHLQTS QVQAGDRQTDHVFETESLSAQGQS SGRKPVRYSRVEIPSGQVSRQKEVF QALEAGKRGVQETVLSPGSIPTGSVH KFTWLFENCPMGLAAESVSGGNL QEEQPKVSADPGTPERQETAERTL RTLHATPGILHHGGILMEARGPGEL CLAKYVLPSPGQGRPCVQKEELVCG ELPRIVRQVLGRDQVQGLLVQED TAGQLRLHPLKLPDGPDPENIEDMD PELQLLACGLGVSMAGTGLVMQE TGQGLVALTAYSLQPQTSRGPERS VQLLASCIDKGLHSLPSLRWEPPTD SSVVPVTEETQRLPPTESIIRVTP LDS TMKMGHLRVSGSTPCPPLPRAAGKV VLPNRPGLQEQARKQVDISHAGQ KGKAASGRPEGTIAPPLGSGAPNLQ E AMQNLRLATAEAQSLHQVLSRHA QGS DPAATSISAQDVLQVPTPTTTTQ G SIMPVAGRARITAVPRKLL	False	False	2.003	2.911	0.859	0.556	False	1.046	0.533