

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
G5E866	G5E866_MOUSE	Mus musculus	NaN	25.161187	NaN	NaN	34887587	MAKIAKTHEDIEAQIREIQGKKAALD EAQGVGLDSTGYDQEIYGGSDSRF AGYVTSIAATELEDDDDYSSSTSL GQKKPGYHAPVALLNDIPQSTEQYD PFAEHRPPKIADREDEYKKHRRTMII SPERLDPFADGGKTPDKMNNARTY MDVMREQHLTKEREIRQQLAEKA KAGELKVVNGAAASQPPSKRKRWW DQTADQTPGATPKKLSWDDQAETPG HTPSLRWDETPGRAKGETPGATPG SKIWDPTPSHTPAGAATPRGDTGPG HATPGHGGATSSARKNRWDETPKT ERDTPGHGSGWAETPRTRDGGDSIG ETPTPGASKRKSrWDETPASQMGGS TPVLTGKTPIGTPAMNMATPTPGHI MSMTPEQLQAWRWEREIDERNRPL SDEELDAMFPEGYKVLPPPAGYVPIR TPARKLTATPTPLGGMTGFHMOTED RTMKS VNDQPSGNLPFLKPDIDQYF DKLLVDVDESTLSPEEQKERKIMKL LLKIKNGTTPMRKALRQITDKAREF GAGPLFNQILPLMSPTLEDQERHL LVKVIDRILYKLDLVRPYVHKILVVI EPLLEDYARVEGREHISNLAKAAG LATMISTMRPDIDNMDEYVRNTAR AFAVVASALGIPSLPFLKAVCKSKK SWQARHTGKIVQQAAILMGCAILPH LRSLVEIEHGLVDEQKQVRTISALAI AALAEATPYGIESFDSVLKPLWKG RQHRGKGLAAFLKAIGYLIPLMDAEY ANYYTREVMLILIREFQSPDEEMKKI VLKVVKCCCGTDGVEANYIKTEILPP FFKHFWQHRMALDRRNYRQLVDTT VELANKVGAAEHSRIVDDLKDEAEQ YRKMVMETIEKIMGNLGAADIDHKL EEQLIDGILYAFQEQTEDSVMLNG FGTVVNALGKRVKPYLPQICGTVLW RLNNSAKVRQQAADLISRTAVVMK TCQEEKLMGHLGVVLYEYLGEEYPE VLGSILGALKAI VNVIGMHKMT PPIK DLLPRLTPILKNRHEKVQENCIDLVG RIADRGAEYVSAREW MRICFELLELL KAHKKAIRRATVNTFGYIAKAIGPHD VLATLLNKLKQERQNRVCTTVAIAI VAETCSPFTVLPALMNEYRVPENLV QNGVLKSLSFLFEYIGEMGKDIYAV TPLEDALMDRDLVHRQTASAVVQH MSLGVYFGCEDSLNHLNLYVWPV VFETSPHVIQAVMGALEGLRVAIGPC RMLQYCLQGLFHPARKVRDVYWKIY NSIYIGSQDALIAHYPRYNDKNTYI RYELDYIL	True	False	1.556	5.0	1.459	1.032	0.523	3.155	1.227