

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
D3YZ62	D3YZ62_MOUSE	Mus musculus	NaN	25.161187	NaN	NaN	34887587	MAASELYTKFARVWIPDPPEEVWKSA ELLKDYKPGDKVLLHLLEEGKDLLEY RLDPKTGELPHLRNPDILVGENDLT ALSYLHEPAVLHNLRVRFIDSKLIYT YCGIVLVAINPYEQLPIYGEDIINAYS GQNMGDMDPHIFAVAEAEAYQOMAR DERNQSIIVSGESGAGKTVSAKYAM RYFATVSGSASEANVEEKVLASNPI MESIGNAKTTRNDNSSRFGKYIEIGF DKRYRIIGANMRTYLLKSRVVFQAE EERNYHIFYQLCASAKLPEFKMLRL GNADSFHYTKGGSPMIEGVDDAK EMAHTROACTLLGISESYQMGIFRIL AGILHLGNVGFASRDSDSCTIPPKHE PLTIFCDLMGVVDEEMCHWLCHRK LATATETYIKPISKLOATNARDALAK HIYAKLFNWIVDHNQALHSAVKQH SFIGVLDIYGFETFEINSFEQFCINYA NEKLQQQFNMHVFKLEQEEYMKE QIPWTLIDFYDNQPCINLIESKLGILD LLDEECKMPKGTDDTWAQKLYNTH LNKCALFEKPRMSNKAFIKHFADK VEYQCEGFLEKNKDTVFEEQIKVLK SSKFKMLPELFQDDEKAISPTSATSS GRTPLTRVPVKPTKGRPGQTAKEHK KTVGHQFRNSLHLLMETLNATTPHY VRCIKPNDFKFPFTFDEKRAVQQLR ACGVLETIRISAGFSPSRWTYQEFFS RYRVLKQKQDVLGDRKQTKCNVLE KLILDKDYQFGKTKIFFRAGQVAYL EKLRAADKLRAACIRIQKTIRGWLLRK RYLCMQRAAITVQRYVRYGQARCYA KFLRRTKAATTIQKYWRMYVRRRY KIRRAATVIQSYLRGYLTRNRYRKIL REYKAVIIQKRVRGWLARTHYKRTM KAIVYLQCCFRMMMAKRELKCLKIE ARSVRYKKLHIGMENKIMQLQRKV DEQNKDYKCLMEKLTNLEGVYNSE TEKLRNDVERLQLSEEEAKVATGRV LSLQEEIAKLKRDLEQTRSEKKSIEE RADKYKQETDQLVSNLKEENTLLKQ EKETLNHRIVEQAKEMTETMERKLV EETKQLELDLNDERLRYQNLNNEFS RLEERYDDLKEEMTLMLNVPKPGH KRTDSTHSSNESEYTFSSFAETEDI APRTEEPIEKVPLDMSLFLKQKRV TELEQEKQLMQDELDRKEEQVFRS KAKEEERPOIRGAEEYEYSLKROELE SENKKLNELNELRKALSEKSAPEV TAPGAPAYRVLMEQLTSVSEELDVR KEEVLILRSQLVSKAEIAPKDDKNT MTDSTILLEDVQKMKDKGEIAQAYIG LKETNRLLESQSQKRSHEEAEAE LRGEIQSLKEENNRQQQLLAQNQL PPEARIEASLQHEITRLTNEIDLME QLEKQDKTVRKLKQKLVFAKKIGE LEVGMENISPGIIDEPIRPVNIIPRK EKDFQGMLEYKREDEQKLVKNLILE LKPRGVAVNLIPGLPAYILFMCVRHA DYLNDQKVRSLTSTINSIKKVLK RGDDFETVSWLNTCRFLHCLKQY SGEEGFMKHNTSRQNEHCLTNFDL AEYRQVLSDLAIQIYQQLVVRVLENIL QPMIVSGMLEHETIQVGSVGPVPTGL RKRTSSIADEGTYTLDLRLQNSFH SVMCQHGMDELKQVVKQMFYIV GAILNLLLRKDMCSWSKGMQIRY NVSQLEEWLRDKNLMNSGAKETLE PLIQAAQLQVKKKTDAAAIAICSM CNALTTAQIVKVLNLYTPVNEFEERV SVSFIRTIQMRRLDRKDSPOLLMDA KHIFPVTFPFNPSSLALETIQIPASLG LGFARV	None	None	None	None	None	None	None		