	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	D3Z5K8	D3Z5K8_MOUSE	Mus musculus	NaN	25.775171	NaN	NaN	34678516	MPRSPTSSEDEMAQSFSDYSVGSES DSSKEETIYDTIRATTEKPGGVKMEE LQGNTLVIRVVIQDLQQTKCIRFNPD ATVWVAKQRILCTLNQGLKDVLNYG LFQPASNGRDGKFLDEERLLREYPQ PMGPGVPSLEFRYKKRVYKQSNLDE KQLARLHTKTNLKKFMDHAQHRSV EKLAKLLDRGLDPNFHDLETGETPL TLAAQLDDSMEVIKALRNGGAHLDF RSRDGMTALHKAARMRNQVALKTL LELGASPDYKDSYGLTPLYHTAIVGG DPYCCELLLHEHASVCCKDENGWH EIHQACRYGHVQHLEHLLFYGADM SAQNASGNTALHICALYNQDSCARV LLFRGGNKELKNYNSQTPFQVAIIAG NFELAEYIKNHKETDIVPFREAPAYS NRRRRPPNTLAAPRVLLRSNSDNNL NAGAPEWAVCSAATSHRSLSPQLLQ QTPSKPDGATKSLGSYTPGPRSRSPS LNRLGGTAEDGKRTQPHWHVGSPF TPGANKDSLSTFEYPGPRRKLYSAVP GRLFVAVKPYQPQVDGEIPLHRGDR VKVLSIGEGGFWEGSARGHIGWFPA ECVEEVQCKPRDSQAETRADRSKKL FRHYTVGSYDSFDAASDCIIEDKTVV LQKKDNEGFGFVLRGAKADTPIEEF TPTPAFPALQYLESVDEGGVAWQAG LRTGDFLIEVNNENVVKVGHRQVVN MIRQGGNHLILKVVTVTRNLDPDDT ARKKAPPPPKRAPTTALTLRSKSMTA ELEELDKPEEIVPASKPSRTAENVAI ESRVATIKQRPTSRCFPAASDVNSVY ERQGIAVMTPTVPGSPKGPFLGLPR GTMRRQKSIDSRIFLSGITEEERQFL APPMLKFTRSLSMPDTSEDIPPPPQS VPPSPPPPSPTTYNCPRSPTPRVYGTI KPAFNQNPVVAKVPPATRSDTVATM MREKGMFYRRELDRFSLDSEDVYS RSPAPQAAFRTKRGQMPENPYSEVG KIASKAVYVPAKPARRKGVLVKQSNV EDSPEKTCSIPIPTIIVKEPSTSSSGKS SQGSSMEIDPQATEPGQLRPDDSLT VSSPFAAAIAGAVRDREKRLEARRNS PAFLSTDLGDEDVGLGPPAPRMQAS KFPEEGGFGDEDETEQPLLPTPGAA PRELENHFLGGGEAGAQGEAGGPLS STSKAKGPESGPAAPLKSSSPAGPEN YVHPLTGRLLDPSSPLALALSARDRA MQESQQGHKGEAPKADLNKPLYIDT KMRPSVESGFPPVTRQNTRGPLRRQ ETENKYETDLGKDRRADDKKNMLI NIVDTAQQKSAGLLMVHTVDVPMA GPPLEEEEDREDGDTKPDHSPSTVP EGVPKTEGALQISAAPEPAVAPGRTI VAAGSVEEAVILPFRIPPPPLASVDLD EDFLFTEPLPPPLEFANSFDIPDDRA ASVPALADLVKQKKNDTSQPPTLNS SQPANSTDSKKPAGISNCLPSSFLPP PESFDAVTDSGIEEVDSRSSSDHHL ETTSTISTVSSISTLSSEGGESMDTCT VYADGQAFVVDKPPVPPKPKMKPIV HKSNALYQDTLPEEDTDGFVIPPPAP PPPPGSAQAGVAKVIQPRTSKLWGD VPEVKSPILSGPKANVISELNSILQQ MNRGKSVKPGEGLELPVGAKSANLA PRSPEVMSTVSGTRSTTVTFTVRPGT SQPITLQSRPPDYESRTSGPRRAPSP VVSPTELSKEILPTPPPPSATAASPSP TLSDVFSLPSQSPAGDLFGLNPAGR SRSPSPSILQQPISNKPFTTKPVHLW TKPDVADWLESLNLGEHKETFMDN EIDGSHLPNLQKEDLIDLGVTRVGH RMNIERALKQLLDR	None	None	None	None	None	None	None	None	None