	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	B7ZNA4	B7ZNA4_MOUSE	Mus musculus	NaN	29.185005	NaN	NaN	22645316	MALGTQLMLLLWKNYTYRRRQPIQL LVELLWPLFLFFILVAVRHSHPPLEH HECHFPNKPLPSAGTVPWLQGLVC NVNNSCFQHPTPGEKPGVLSNFKD SLISRLLADTRTVLGGHSIQDMLDAL GKLIPVLRAVGGGVRPQESDQPTSQ GSVTKLLEKILQRASLDPVLGQAQDS MRKFSDAIRDLAQELLTLPSLMELR ALLRRPRGSAGSLELVSEALCSTKGP SSPGGLSLNWYEANQLNEFMGPEV APALPDNSLSPACSEFVGTLDDHPV SRLLWRRLKPLILGKILFAPDTNFTR KLMAQVNQTFEELALLRDLHELWG VLGPQIFNFMNDSTNVAMLQRLLD VGGTGQRQQTPRAQKKLEAIKDFLD PSRGGYSWREAHADVGRLAGILGQ MMECVSLDKLEAVPSEEALVSRALE LLGERRLWAGIVFLSPEHPLDPSELS SPALSPGHLRFKIRMDIDDVTRTNKI RDKFWDPGPSADPFMDLRYVWGGF VYLQDLLEQAAVRVLGGGNSRTGLY LQQMPHPCYVDDVFLRVLSRSLPLF LTLAWIYSVALTVKAVVREKETRLRE TMRAMGLSRAVLWLGWFLSCLGPF LVSAALLVLVLKLGNILPYSHPVVIFL FLAAFAVATVAQSFLLSAFFSRANLA AACGGLAYFALYLPYVLCVAWRERL HLGGLLAASLLSPVAFGFGCESLALL EEQGDGAQWHNLGTGPAEDVFSLA QVSAFLLLDAVIYGLALWYLEAVCPG QYGIPEPWNFPFRRSYWCGPGPPKS SVLAPAPQDPKVLVEEPPLGLVPGVS IRGLKKHFRGCPQPALQGLNLDFYE GHITAFLGHNGAGKTTTLSILSGLFP PSSGSASILGHDVQTNMAAIRPHLGI CPQYNVLFDMLTVEEHVWFYGRLK GVSAAAMGPERERLIRDVGLTLKRD TQTRHLSGGMQRKLSVAIAFVGGSR VVIMDEPTAGVDPASRRGIWELLLKY REGRTLILSTHHLDEAELLGDRVAM VAGGSLCCCGSPLFLRRHLGCGYYL TLVKSSQSLVTHDAKGDSEDPRREK KSDGNGRTSDTAFTRGTSDKSNQAP APGAVPITPSTARILELVQQHVPGAQ LVEDLPHELLLVLPYAGALDGSFAM VFQELDQQLELLGLTGYGISDTNLEE IFLKVVEDAHREGGDSRPQLHLRTC TPQPPTGPEASVLENGELAKLVLDP QAPQGLAPNAAQVQGWTLTCQQLR ALLHKRFLLARRSRRGLFAQVVLPAL FVGLALFFSLIVPPFGQYPPLQLSPA MYGPQVSFFSEDAPGDPNRMKLLE ALLGEAGLQEPSMQDKDARGSECT HSLACYFTVPEVPPDVASILASGNWT PESPSPACQCSQPGARRLLPDCPAG AGGPPPPQAVAGLGEVVQNLTGRNV SDFLVKTYPSLVRRGLKTKKWVDEV RYGGFSLGGRDPDLPTGHEVVRTLA EIRALLSPQPGNALDRILNNLTQWA LGLDARNSLKIWFNNKGWHAMVAF VNRANNGLLHALLPSGPVRHAHSIT TLNHPLNLTKEQLSEATLIASSVDVL VSICVVFAMSFVPASFTLVLIEERITR AKHLQLVSGLPQTLYWLGNFLWDM CNYLVAVCIVVFIFLAFQQRAYVAPE NLPALLLLLLLYGWSITPLMYPDSFF FSVPSTAYVVLTCINLFIGINSSMATF VLELLSDQNLQEVSRILKQVFLIFPH FCLGRGLIDMVRNQAMADAFERLG DKQFQSPLRWDIIGKNLLAMMAQG PLFLLITLLLQHRNRLLPQSKPRLLP PLGEEDEDVAQERERVTKGATQGDV LVLRDLTKVYRGQRNPAVDRLCLGIP PGECFGLLGVNGAGKTSTFRMVTGD	None	None	None	None	None	None	None	None	None

								TLPSSGEAVLAGHNVAQERSAAHRS MGYCPQSDAIFDLLTGREHLELFAR LRGVPEAQVAQTALSGLVRLGLPSYA DRPAGTYSGGNKRKLATALALVGDP AVVFLDEPTTGMDPSARRFLWNSLL SVVREGRSVVLTSHSMEECEALCTR LAIMVNGRFRCLGSSQHLKGRFGAG HTLTLRVPPDQPEPAIAFIRITFPGAE LREVHGSRLRFQLPPGGRCTLTRVF RELAAQGRAHGVEDFSVSQTTLEEV FLYFSKDQGEEEESSRQEAEEEEVS KPGRQHPKRVSRFLEDPSSVETMI								