

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
A0A1N9NPH8	A0A1N9NPH8_MOUSE	Mus musculus	NaN	25.775171	NaN	NaN	34678516	MTEVTRAGSWLSRRYPNGCWVAGR VRVPCREARPHLSDPLYLTYSEM KRKGMNTKRLRLSPSEEAFFILKEDYE RRRKLRLQVREQERGIQAFQIREGIK QRRSQVSHLAEELRAKWEEAQSQ KIQNLEKLYLASLRHMGDGHROAKE NEPDLDALSRRAAERKRKAQVVRHKE ALKVQKNQKEMLMKQKTRHIKARK EAVLVEKQRSAMKARLPPVPSPF NIDVNRIPSLKTNSTYHHISTFVSR QMGTKQPDHLLAAEEARRVERLR KQAAQERVKQFERAHVRSQAMKK IHIAQNERLMEELKQLQKEDLARR RQTVAQMPQMELPYRRSEMKEKED RQRELEFAFEDMYNADRKVGNLIL HLKPEPLPTISDQLQDEELDSMEQ ENQVPLAAKIQQIPSRILFKRLLNKIR SQKSLWTIKSVSEDEGEVTSIIIEIES KVPSVDSGAIITEERTASFEQEQVT DSDRLTIESGPLSSEDKPLYKAGTG REQAMAVSPATAVAQSSVLLHPQE EAVRIRMSLRRKQIMEIEEQKQKQL ELLEQIEQQKRLRETDCAFRAQLEEQ RKQADQPEVCCAPMSHAMISDEDS HRQMIRNYQHQLLQQRNLKQKQKQ TARKRLLEYQTVLKERSPSLSASALV PDSVVSGPPQSYKPAASDSDWDP QRLKLSPSKYQVQSPQIPALEQSHI QVPRHGHITQRQGMVAVSEMLGKQ PVESQERQWQFSQVETHQGDYEFV LKDSHLSRSLSYVRPOTLQDAREV SKPPRVICQSLDSQQISSEDSSENISS KPSESPFPLPLVPERPFTSLPVKFFHS GTIHKPFTTINOSVISQMHQDPLSSS ETTAAQGGDLRFLQEQLELQKQVLO ARQEAAREKLLCTQKELGQQTGLPV FLPSPAGNIFSSLPSASAESGNFQTS STKSDATVSSDNMDRLWDSQPISS QOHLQEQSSVETDNLQARREA QEVFLAHTQNTLEKIVRSEQAGSSLP HQVAQSFSSLTADTQSKKIQKQP LPANKKGLLPSQSEVSKAQDGSSEGF LQQTLPQNTLKLQEQQLTRQSRMI PPRRDGOETLLLYKESCEDSEAGP VESLSSVVQHADASRAVSEVPKRL QDVYSSEENRVLSSHLITHGFPOH SLQRQEHFTPLQEEETHIQRLLGARK NNEEFAPKQNELEKGLCSQQTDALS SPSQVTDWGTSRGVSVRSRDRDPL RHFKIPAFRERLVRVSOHTFPLQDN LQEHQEWVDTEKESFQSSPLTPENP SSQQTGFSSFKASRLRPSVSLPSAD SGITQHPLSTESDSKVKSSHLQIPEL QHRLSKISQLIPPQDLSKALQEQLA TQREAIHSRQEAHEETLREWKEKIF PEQVGFPSPLIPQHSASFPVSDTER AQELCSTNSDTISSGYPEMELPDRT LGLSHTALPQNNLTAHPEHLHAQT NFFHSTEKAEGLVFRPCQFEEMS AEHFIQPDHDLKALQQLDMQRE AIRSGQEMQEKMLLQRLNKLEORIS SKQISSSLFSSQVALPIANSDGTLOS FPKTSNETELLGSQDEYLSFSQPRLP LQNNMTEQLDLEKVFHKEILLHKQ KSQNKSESSEHSLPPLFLSKEIEHPF ISLPAESKSKSICELYLSDKHAAP NDAVIPRLQDRLLSCSQPVLTOQDN MSLQKQLNLQRETLHSRQKAQEEL LVORQTSLQQQIQRHRETLLKNFFNV SQARNPTDENDLEMQRKREQLGGWF PHTQGLTWGDAGQGSANGEQPRAD VHAENGESEKELSGRASKPPVSK VKCVLDLNQHELSTIQEVESPASGRI SMPGKAIFYQDRDPLRVSVSREQSF LESPLAHDPFQCHQPPAQENSKSHD DNAEAVKVKSDVEDHAVLSHAVSK EEACTNLGKPLGKPDDEAETQEISQE	None	None	None	None	None	None	None		

PLSSVTVSTGSLFYSYEITDLSLTDPEP
FSEQTEHLEQESTNKQEETDPLSIAV
PSVIYQQQHSLGAHNLLPMEEEST
SDHTHVQQIMDNDVNEANLIPDKR
DFQVPAVDLDFRELEHIFPHLHRQL
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LQHSTPCASVSSECSVKLLESREEVL
GFEELSRRAVTMSQRLTEDEVVLP
INPHVGRVEKEASVQGSNPLSIQNE
KPIQNFIEDTTEAVGNVCQLAQAE
HILKSCPFRSPIWETDTGYGIMEE
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EDNEAQFFQAGVVLP TSSMETSVCG
AVSEPYVDQPTVAPSATSGSLQEAF
MTRQTLTERSQRQREIWNKTRLPO
TKVSKEKLPTGCTGS