

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9NZM1	MYOF_HUMAN	Homo sapiens	Myoferlin	3.7625	T617;S727	S174;S729;S1915	35083852;34019948;34725712;37217939;26853435;33214551;33465208;30379171	MLRVIVESASNIPKTKFGKPDPIVSVI FKDEKKKTKKVDNELNPVWNEILEF DLRGIPLDFSSSLGIIVKDFETIGQNK LIGTATVALKDLTGDQSRSLPYKLISL LNEKGQDTGATIDLVIIGYDPPSAPHP NDLSGSPVPGMGGDGEEDEGEDR LDNAVRGPGPKGPVGTVSEAQLARR LTKVKNSRRMLSNNKQDFQIRVRVI EGRQLSGNNIRPVVKVHVCQTHRT RIKRGNNPFFDELFFYNVNMTPSEL MDEIISIRVYNHSLRADCLMGFEKI DVGfVYDEPGHAVMRKWLLLNDPE DTSSGSKGYMKVSMFVLGTGDEPPP ERRDRDNDSDVESNLLLPAIALR WVTFLLKIYRAEDIPQMDDAFSQT KEIFGGNADKKNLVDPFVEVSFAGK KVCTNIIKKNANPEWNQVNLQIKF PSVCEKIKLTIYDWRDLTKNDVVGTT YLHLSKIAASGGEVEDFSSSGTGAAS YTVNTGETEVGFVPTFGPCYLNLYGS PREYTGFPDPYDELNTGKGEVAYR GRILVELATFLEKTPPKLEPISND DLLVVEKYQRRRKYSLSAVFHSATM LQDVGEAIQFEVSIGNYGNKFDTTT KPLASTTQYSRAVFDGNYYYLPWA HTKPVVTLTSYWEDISHRLDAVNTL LAMAERLQTNIEALKSGIQGKIPANQ LAELWLKLIDEVIEDTRYTLPLTEGK ANVTVLDTQIRKLRSRSLSQIHEAAV RMRSEATDVKSTLAEIEDWLDKLMQ LTEEPQNSMPDIIWMIRGEKRLAYA RIPAHQVLYSTSGENASGKYCGKTQ TIFLKYPQEKNNGPKVPVELRVNIWL GLSAVEKKFNSFAEGTFTVFAEMYE NQALMFGKWGTSGLVGRHKFSDVT GKIKLKREFFLPPKGWEWEGEWIVD PERSLLTEADAGHTEFTDEVYQNES RYPGGDWKPAEDTYTDANGDKAASP SELTCPGWEWEDDAWSYDINRAV DEKGWEYGITIPPDHKPKSWVAAEK MYHTRRRRLVRKRKKDLTQTASST ARAMEELQDQEGWEYASLIGWKFH WKQRSSDTFRRRRWRRKMAPSETH GAAAIFKLEGALGADTTEDGDEKSLE KQKHSATTVFGANTPIVSCNFDRVYI YHLRCYVYQARNLLALDKDSFSDPY AHICFLHRSKTTEIIHSTLNPTWDQT IIFDEVEIYGEPQTVLQNPVKVIMELF DNDQVGKDEFGLRSIFSPVVKLNSE

MDITPKLLWHPVMNGDKACGDVLV  
TAEILRGKDGSNLPILPPQRAPNLY  
MVPQGIRPVVQLTAIEILAWGLRNM  
KNFQMASITSPSLVVECGGERVESV  
VIKNLKKTPNFPSSVLFMKVFLPKEE  
LYMPPLVIKVIDHRQFGRKPVVGQCT  
IERLDRFRCDPYAGKEDIVPQLKASL  
LSAPPCRDIEMEDTKPLLASKLTE  
KEEEIVDWWSKFYASSGEHEKCGQY  
IQGYSKCLKIYNCELENVAEFGLTD  
FSDTFKLYRGKSDENEDPSVVGEFK  
GSFRIYPLPDDPSVPAPPRQFRELPD  
SVPQECTVRIYIVRGLELQPQDNGL  
CDPYIKITLGKKVIEDRDHYIPNTLNP  
VFGRMYELSCYLPQEKDLKISVYDYD  
TFTRDEKVGETIIDLENRFLSRFGSH  
CGIPEEYCVSGVNTWRDQLRPTQLL  
QNVARFKGFPPQILSEDGSRIRYGGR  
DYSLDEFEANKILHQHLGAPEERLA  
LHILRTQGLVPEHVETRTLHSTFQP  
NISQGKLQMWVDVFPKSLGPPGPPF  
NITPRKAKKYLRVWIWNTKDVILDEK  
SITGEEMSDIYVKGWIPGNEENKQK  
TDVHYRSLDGEGNFNWRVFVFPFDY  
LPAEQLCIVAKKEHFWSIDQTEFRIP  
PRLIIQIWDNDKFSLDDYLGFLDL  
RHTIIPAKSPEKCRDLMIPDLKAMNP  
LKAKTASLFEQKSMKGWWPCYAEK  
DGARVMAGKVEMTLEILNEKEADE  
RPAGKGRDEPNMNPKLDLPNRPET  
SFLWFTNPCKTMKFIVWRRFKWVII  
GLLFLILLFFVAVLLYSLPNYLSMKI  
VKPNV