

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
A0A1B0GS91	A0A1B0GS91_MOUSE	Mus musculus	NaN	10.025162	NaN	NaN	34678516	MFRSPGLSSCLGAFRFSEKQDFNKK NLGFQGFPHQRKAASFQKEFSFDDK EEPANHRRGLDVKCLAVLPKVAELR KIFEPKRKDFLEMKRKERIARRLEGI ENDSQPILLQSQCTGLVTHRLLLEEDTP RYMRATDPASPHIGRSKEEEDTPGS SLEKQTPSKYCIETSGIHSSGSMDTH SLESKAERIARYKAERRRQLAEKYGL TLDPEADSEYLSRYAKSRKDPDVTER RGKSDKQEEQSKDANSRHSRTESG PRTSLVASQDCTPLGSNMSDQEQLL NVENQRRVQDPPLGEDGSSAFFSER SISFPEVPRSPKQIPSSPLQQPASP HPGDSPLPTEARASTGKPTHEWFLQ RDSEGDTPSLINWPSRVKVRKLVK EESARSSPELTSESLTQRROQPAPA HFLPIQSESSTFDRVTSKAVSSLQPS QSGVLPDTPVHAIKLVMTDTPESTSE FSWVGSATPKVIKSTTLKILEGGSRD APVLHICESKAEDWLSPEPLERSPKS LLTSEDDRLVRGHKDPGKNDLKA IICSIDVESERERQVQHLPTQRTGRS EMLLYVQSGPVSQDATLTSHTKEAS PKKRKVLARSLSDYTGPPQLQVPRH KDEAPSQELELQSSRAEGPGAESV LDTRVSAQLRNIFMESTRASKKPE LQSRVERSAEGIGLPMERERGSRKP RRYLSPGESRKTSEFRTPITSAER KESDRYPSGSEIPVVEDEEKVDERAK LSVAAKRLLFREMEKSFDEHTVPKR HSRNAAVEQRLRRLQDRSHTQPITT EEVVIAATEPIPASCSGVTHPVTARLP SPTVARSSVQPARLQASAHQKALAR DQANEGRESAEPGEPDSSTLSLAEK LALFNKLSQPVSKAISTRNRIDVRQR RMNARYQTQPVTLGEVEQVQSGKLI SFSPTVNTSVSIMASAVPTYAGDLR KLSVDNNTSATDYKSPAENS DSPV RSILKPQAWRPLVEHSGSKGMPGES GKTESKNALTVAAEDSGVQTRGAFE EEEEPSYPILGRVREGDGQKEPKHV VLRGSLLELGNPSAAHLGDELKEVS TAKSSLQENLDLKDQASEENTDVE TVMRKFSLKEFGETTSEQTEVAARK ASVQMATPGAWKQQESSEQLAEKL FKNPCAMFASGEVKVPVGDSDLSP SKTMSIKERLALLKKSPEEDWKNRL IRKQEYKATGGLHTQEVEQSLKKK RVTESRESQMTIEERKHLITVREEA WKTGKRGGAANDSTQFTVAGRMVKK

GLASPTSITPISSPLCSKSRGTTTPVSK
PLEDIEARPDMQLESDLKLDRLLETFL
RRLNKNVAGIQETVLTVTGKSVEV
MKLDDDETFKAFYRSVDHSIPRSPV
ELEEDFDVIFDPYAPKLTSSVAEHKR
QVRPKRRVQASKNPLKLLAARDLL
QEYTEQRLNVAFMESKRMKVEKMS
SNSNFSEVTLAGLASRENFSNINLR
SVNLMEQNSNNSAMPYKKLMLLQI
KGRRHVQTRLVEPRASSLNSGDCFL
LLSPQYCFLWVGEFSNVIEKAKASEL
ATLIQTKRELGCRATYIQTIEGINTH
THAAKDFWKLLGGQTSYQSAGDPKE
DELYETAIETNCVYRLTDDKLPDD
DYWGKIPKCSLLQSKEVLVDFGSE
VYVWHGKEVTLAQRKIAFQLAKHLW
NGTFDYENCDINPLDPGECNPLIPR
KGQGRPDWAIFGRVTEHNETILFKE
KFLDWTELKRPTEKNSGEVVQKD
DPRADV KPYDVTRMVATPQITAGTIL
DGVNVGRGYGLVEGDDRRQFEIATV
SVDVWHILEFDYSRLPRQSIGQFHE
GDAYVVKWKYMASTAVGSRQKGEH
LVRVAGKEKCVYFFWQGRHSTVSEK
GTSALMTVELDEERGAQVQVLQGKE
PPCFLQCFQGGMVVHSGRREEEEE
NVQSEWRLYCVRGEVPMEGNLLEV
ACHCSSLRRTSMVVLNINKALIYL
WHGCKAQGHTKEVGRTAANKIKEE
CPLEAGLHSSSNVTIHECDEGSEPL
GFWDALGRRDRKAYDCMLQDPGSF
NFAPRLFILSSSGDFSATEFVYPAQ
APSAVSSMPFLQEDLYSAPQPALFLV
DNHHEVYLWQGWPTENKITGSAR
IRWASDRKSAMETVLQYCRGKNLKR
PPPKSYLIHAGLEPLTFTNMFPSWE
HREDIAEITEMDTEVSNQITLVEDVL
AKLCKTTYPLADLLARPLPEGVDPLK
LEIYLTDEDFEFALDMSRDEFNALPT
WKQVNLKKSGLF