

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
A0A0A0MQ79	A0A0A0MQ79_MOUSE	Mus musculus	NaN	15.841233	NaN	NaN	37507081	MSEKSGQSTKAKDGGKYATLSLFNT YKGSLETQKTTARHGLQSLGKVGIS RRMPPANLPSLKAENKGNPNVNI VPKDG TGWASKQEQHEEEKAPEVS PAQPKPGVAAPPEVAPAPKSWASNK QGGQGDGIQVNSQFQQEFPSLQAA GDQEKEKEANDENYGPGPSLRPP NVACWRDGGKSAGSPSSDQDEKQL GDESTAITSEQNDILKVVEKRIACG PPQAKLNGQQPALASQYRAMMPY MFQQYPRMAYPPLHGPMRFPSPSL EANKSLRGRGPPPSWASEPERPSILS ASELKELDKFDNLDAEADGWAGA QMEVDYTEQLNFSDDDEQGSTSPK ESSSEDQTAKTPESTENRKEVDEAS STKSSSQIPAQPPVTKSPYKGGPPFN QERGSSHLPPPKLLAQQHPPPPD RQIPGRQGPFPSKPPVDPNDDEIWKQ RRKQQSEISAVERARKRREEEERR MEEQRKAACAELKQLDEKLGIEK QPSPEELREREREKERERELEKEKE RELEKEQEKQREMERARQOKELE QREKEQELQRLREQEKEGEPKEQ EKEEKVEPQEPVVEPATENQSENN CKKEEPIFTRQDSNRSEKETTQVV QEAEPESGAQPRPGYFKQFQKSLPP RFQRQEQMKQQQWQQQQQQQQ QGVLPQTVPSQPSNGSVPPPHRPL YQPMQPHPQHILASMGFDPRWLMM QSYM DPRMISGRPAMDIPPIHPGMI PPKPLIRRDQMEGSPNSSESFEHIAR SARDHGISLSEPRMMWGSDPYHAE PQQAATPKSAEETGDARPETAMDHE HMTAAYPVEHSQLETHSKTDVARDS TETEEQKFLRSLEDVKPRHVD TNT QSACFDVIDQKSLPTSAERISALES QPARKRSVSHGSNHAQNAEEQRNE PSVSIPKVINRCMDSKETVEKPEEK RKDGF LRSSEGPKPEKVYKSKSETR WGPRPSSNRREEGNDRPVRRS GPIK KPVLRDMKEEREQRKEKEGEKLEKV TEKVVKA EKPEKKDLPLPLPPAPAQ PQPQLVSPVQPEPEKPPSTETSTL TQKPSQDEKPLEPVGSVQVEPVVKT VNQQSVAAPT VKEEKPEKVINKDV GIERSRPDSRLAVKKDSSLPTRTYWK EARDRDWFPDQGYRGRGRGEYYSR GRSYRGSYGGRRGRGRGHTREYPQY RDNKPRTEHVPSGPLRQRESETRS ESSDFEVVPKRRRQRGSETD TDSEV

HE S A S D K D S V S K G K L P K R E E R P E N K
K P V K P Q S S F K P E N H V R I D N R P L E K P
Y I R E E D K S K P G F L P K G E P T R R G R G G T
F R R G G R D P G G R P S R P A T L R R P A Y R D
N Q W N T R Q A E P P K P E D G E P P R R H E Q
F M P I P A D K R P P K F E R K F D P A R E R P R
R Q R P T R P P R Q D K P P R F R R L R E R E A A
S K T S E V L V P S N G T A N N V V Q E P V N P P
A D I S G N K T P D L S N Q N S S D Q A N E E W
E T A S E S S D F N E R R E R D E K K N A D K S S
Q A V V K A G E S V L P P K R E I A K R S F S S Q R
P G V D R Q N R R G N N G P P K S G R N F S G P
R N E R R N G P P S K G G K R G P F D D Q A S G
T A G A D P V S G N S A H H Q E G V P N G A G Q
K N S K D A A G K K R E D T K P G P K K P K E K V
D A L S Q F D L N N Y A S V V I I D D H P E V T V I
E D P Q S N L N D D G F T E V V S K K Q Q K R L
Q D E E R R K K E E Q V V Q V W S K K N I G E K
G R S Q T S K L P P R F A K K Q A T G T Q Q I Q A P
P S A P V P V S S S A P G L T A P A A A A P A S T P
A P V P I L A S A T A L V P V S T P A P V L T S C P A
P V P T S A S A P V P A S T S S P V T A S S S S Q P
S V P A P T P V L A S A S T T V S V P I L T S A S I P I
L A S A L A P A T V S S P T P V V S A T A V P S I S T
P A V P A S A P T A S V P L A P A S A A S T V P P P
A S T V Q T Q T Q T Q T H K P V Q S P L P P S A P S
S K Q P P P S I R L P S A Q T S N G T D F V A A G K
S M P T S Q S H G S L T A E L W D S K V A A P A V
L N D I S K K L G P I S P P Q P P S V S A W N K P L
T S F G S A T S S E G T R N G Q E S G V E I G I D T
I Q F G A P A S N G N E N E V V P V L S E K A T D
K V P E P K E Q R Q K Q P R A G P I K A Q K L P D
L S L V E N K E H K P G P I G K E R S L K N R K V
K D A Q Q V E P E G Q E K P S P A V V R S T D P E
T A K E T K A V S E M S A E I G A M I S V S S A E Y
G S D A K E S V T D Y T T P S S S L P N T V A T N
N A K M E D T L V N N V P L P N T L P L P K R E T
I Q Q S S S L T S V P P T T F S L T F K M E S A R K
A W E N S P N L R E K G S P V T S T A P P I V S G
V S S S A S G P S T A N Y S S F S S A S M P Q I P V
A S V T P T A S L S G A G T Y T T S S L S T K S T T T
S D P P N I C K V K P Q Q L Q T S S L P S A S H F S
Q L S C M P S L I A Q Q Q S P Q V Y V S Q S A A
A Q I P A F Y M D T S H L F N T Q H A R L A P P S
L A Q Q Q G F Q P G L S Q P T S V Q Q I P I P I Y A
P L Q G Q H Q A Q L S L G A G P A V S Q A Q E L F
S S S I Q P Y R S Q P A F M Q S S L S Q P S V V L S
G T A I H N F P A V Q H Q E L A K A Q S G L A F Q
Q T S N P Q I P I L Y D H Q L G Q A S G L G S S Q
L I D T H L L Q A R A N L T Q A S N L Y S G Q V Q
Q P G Q T N F Y N T A Q S P S A L Q Q V T V P L P
A S Q L S L T N F G S T G Q P L I A L P Q T L Q P Q

							LQHTTPQAAQSLSRPAQVSQPFRG LIPAGTQHSMMATTGKMSEMELKA FGSGIDIKPGTPIGGRSTTPTSSPFR ATSTSPNSQSSKMNSVVYQKQFQSA PATVRMAQPFPAQFAPQILSQPNLV PPLVRAPHTNTFFPAPVQRPPMALAS QMPPPLTTGLMSHARLPHVARGPC GSLSGVRGNQAQAALKAEQDLKAK QRAEVLQSTQRFFSEQQNKQIGGK TORVSDTSDTSDNPETLSDPPGTCPEKV EEKPPPAPTITTKPVRTGPIKQAIKT EETKS
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