

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
B7FAU9	B7FAU9_MOUSE	Mus musculus	NaN	16.84821	NaN	NaN	25153642;34887587	MSSSHSRCGQSAAVASPGGSIDSRD AEMPATEKDLAEDAPWKKIQNTFT RWCNEHLKCVSKRIANLQTDLSGDL RLIALLEVLVSQKKMHRKHNQRPTFR QMQLENVSVALEFLDRESIKLVSIDS KAIVDGNLKLILGLIWTLILHYSISMP MWDEEEDDEEAKKQTPKQRLLGWIQ NKLPLPITNFSRDWQSGRALGALV DSCAPGLCPDWDSWDASKPVNNAR EAMQQADDWLGIPQVITPEEIVDPN VDEHSVMTYLSQFPKAKLKPGAPLR PKLNPKKARAYGPGIEPTGNMVKKR AEFTVETRSAGQGEVLVYVEDPAGH QEEAKVTANNDKNRRTFSVWYVPEV TGTHKVTVLFAGQHIAKSPFEVYVDK SQGDASKVTAQGPGLPSGNIANKT TYFEIFTAGAGMGEVEVVIQDPTGQK GTVEPQLEARGDSTYRCSYQPTMEG VHTVHVTFAGVPIPRSPYTVTVGQAC NPAACRAIGRGLQPKGVRVKETADF KVYTKGAGSGELKVTVKGPKGEERV KQKDLGDGVYGFYYPTIPGTYTVTI TWGGQNIGRSPFEVKVGTECGNQK VRAWGPGLEGGIVGKSADFVVEAIG DDVGTLGFSVEGPSQAKIECDDKGD GSCDVRYWPQEAGEYAVHVLNSEN DIRLSPFMADIREAPQDFHPDRVKA RGPGLKGTGAVNKPAEFTVDAKHA GKAPLRVQVQDNEGCSVEATVKDN GNGTYSYVPRKPKHTAMVSWG GVSIPNSPFRVNVGAGSHPNKVKVY GPGVAKTGLKAHEPTYFTVDCTEAG QGDVSIKICAPGVVGPTEADIDFDII RNDNDTFTVKYTPCGAGSYTIMVLF ADQATPTSPIRVKVEPSHDASKVKA GPGLNRTGVELGKPTHFTVNAKTAG KGKLDVQFSGLAKGDVAVRDVDIIDH HDNTYTVKYIPVQQGPVGNVNTYGG DHIPKSPFSVGVSPSLDLSKIKVSGL GDKVDVKGDKQEFVKSAGAGGQK VASKIVSPSGAAVPCKVEPGLGADNS VVRFPREEGPYEVEVTDGVPVPGS PFPLEAVAPTKPSKVKAFGPGLOGG NAGSPARFTIDTKGAGTGGLGLTVE GPCEAQLECLDNGDGTCSVSYVPTE PGDYNINILFADTHIPGSPFKAHVAP CFDASKVKCSGGLERATAGEVGQF QVDCSSAGSAELTIEICSEAGLPAEV YIQDHGDGTHITTYIPLCPGAYTVTIK

YGGQVPNFP SKLQVEPAVDTSGVQ
CYGPGIEGQGVFREATTEFSVDARAL
TQTGGPHVKARVANPSGNLTDTYVQ
DCGDGTYKVEYTPYEEGVHSVDVTY
DGSPVPSSPFQVPVTEGCDPSRVRV
HGPGIQSGTTNKP NKFTVETRAGAT
GGLGLAVEGPSEAKMSCMDNKDGS
CSVEYIPYEAGTYS LNVTYGGHQVPG
SPFKVPVHDVTDASKVKCSGPGLSP
GMVRANLPQS FQVDTSKAGVAPLQV
KVQGP KGLVEPVDVVDNADGTQTV
NYVPSREGSYSISVLYGEEEVPRSPF
KVKVLP THDASKVKASGPGLNTTGV
PASLPVEFTIDAKDAGEGLLAVQITD
PEGKPKKTHIQDNHDGTYTVAYVPD
VPGRYTILIKYGGDEIPFSPYRVRAVP
TGDASKCTVTGAGIGPTIQIGEETVIT
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GTVTVRYS PSEAGLHEMDIRYDNM
HIPGSPLQFYVDYVNCGHITAYG PGL
THGVVNKPATFTVNTKDAGEGGLSL
AIEGPSKAEISCTDNQDGTCSVSYLP
VLPGDYSILVKYNDQHIPGSPFTARV
TGDDSMRMSHLKVGSAADIPINISE
TDL SLLTATVPPSGREEPCLLKRLR
NGHVGISFVPKETGEHLVHVKKNGQ
HVASSPIPVVISQSEIGDASRVRVSG
QGLHEGHTFEPAEFIIDTRDAGYGGL
SLSIEGPSKVDINTEDLEDGTCRVTY
CPTEPGNYIINIKFADQHVPGPSF SV
KVTGEGRVKESITRRRRAPSVANIGS
HCDLSLKIPEISIQDMTAQVTSPSGK
THEAEIVEGENHTYCIRFVPAEMGM
HTVSVKYKGQHVPGPSPFQFTVGPLG
EGGAHKVRAGGGLERA EVGVPAEF
GIWTREAGAGGLAIAVEGPSKAEISF
EDRKDGSCGVAYVVQEPGDYEVSVK
FNEEHIPDSPFVVPV ASPSGDARRLT
VSSLQESGLKVNQPASFAVSLNGAK
GAIDAKVHSPSGALEECYVTEIDQDK
YAVRFIPRENGIYLIDVKFN GTHIPGS
PFKIRVGE PGHGGDPGLVSAYGAGL
EGGVTGSPA EFIVNTSNAGAGALSVT
IDGPSKVKMDCQECPEGYRVTYTPM
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GPRLVSNHSLHETSSVFVDSLTKVA
TVPQHATSGPGPADVSKVVAKGLGL

									SKAYVGQKSNFTVDCSKAGNNMLL VGVHGPRTPCEEILVKHMGSRLYSV SYLLKDKGEYTLVVKWGDEHIPGSP YRIMVP
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