

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
A0A0B4KF23	A0A0B4KF23_DROME	Drosophila melanogaster	Nipped-B protein	10.212237	NaN	NaN	33925313	<p>MGERDNPTVPVTTLAGLTSTSDYSP DQHTANARLEAKKMQIDSYSIPECY ASPTNLSISNEHQLODAQACFNQLT SMTSKEILEEFSQINFKENATEKDKI DVIENSDLNVTKILSQNTLKKKSNTP ERSTVQSIQEQQFFIQQQEANYNLKH DLNQVSTNVGQIQNISDTFPQEPSN FNLNSVNIPNMLYVQSPPFTESEPT QAHHSSIEYLKKKKSQILDVVSILNRQ QVLENTSLHIINKSSTYQTNQNVSTT TSTSTSSSGKSQVRSPELARSMGLL QETCQQQHENLNIIPTSAEECVLES RASSTNNTVKIPIDTFSTIKADEKRSA KRKLAISIGDIPPEQIFSKPKMRRVER ITPLSNTKCVKEEVTRSQTYQQFIRN MDHIIELDDSESPNFDSEDVDETIE CISSKLLNSMSTDVAKLKAKQALDSI PKNKLTLLINYAMRNVYLARNYFAG TEDEDEFVDDEVIEKLLNAMDACLL ICNIYSTVSDLQFLQEDNVSHIIKFTQ FQLRETIFPLHDPVYTAKSIRTTTHR KKIKSHQAQNRSMLFYLKTVELLK VFVTLFDKCVFVDTIVLPLSTLAIEPF FVDNIETLQFVCLELVTTIFRKERYD KIRNSILGDILTSIDRLPSSKKNLRPY KLTNNGGNIQMVTALVLQLIQCATIL PDSLCDNGKFSNKPQEGNTFDEEG KLLQPSQDLLVLQKYDVAVSIGGN FLTTFLNKCKSRSNETDFRPLFENFI HDLLATVKNKPEWPASELLLSLLGTM LVRYVSDKGIEQSIRLVSLDYLGVAA RLRKDTVESRCRVNIIDSMIQSIKLE QEKEGDVTSNNDQFDLEPEEQRTD FLQKILLDFLAVNAQEENLIWDYAR HFYLAQWYRDVIYQRRRINDGKKGL AFRKSIRNNRRTNGDYLDTSDSGS CDESDTDNKKRIHCVDSNDFELNI NIYKALEARKQYFINKIKPFSVFGEQ NHSSNQHIKTYIDYNNAQLIAQYLAT KRPFSQSFDGCLKKIILVVNEPSIAVR TRAMKCLANIVEVDPLVLKRKDMQ MGVNQKFLDTAISVREAAVDLVGKF VLSNQDLIDQYDMLSTRILDGTGVS RKRVIKILRDICLEYPDFSKIPEICVK MIRRVHDEEGIQKLVTEVFMKMWF TPCTKNDKIGIQRKINHIIDVVNTAH DTGTTWLEGLLSIFKPRDNMLRSE GCVQEFIKKNSEPPMDIVACQQLAD GLVDRLIELEDTDNSRMLGCITTLHL</p>

LAKVRPQLLVKHAITIEPYLNIKCHSA
TAAKFICAVADILEKVVPLVNNASES
FLASLEEHLMLLVSRNQAEVTSCV
SCLGALVNKITHNFKLIRDCFOKFYR
VLDVSRSQVIQGNNSVDNIYTPSFRR
SLFTIGILMRYFDFKSPIALGETNDG
LPVSICEDVFHCLMFFCRCTNQEIR
KQALISLGSFCVLNDGYLTRSELKNL
YCEILSSIANDAGFKIICMRNIWIYLT
ESEMFMHNKEKEWEKQSKHEDLK
EMNDVSSGMASRIIQLYLEEILECFL
NRDDTVRLWAVKVIQIVLRQGLVHP
VRMVPYLICLSTDHRIESAHRADALL
KDIDKTYSGFVNMKVQFGLQLCFKL
QKILQINNRGKLEIIRGYASRGPDNT
TTALNDFLYTLLRTTKPQRRALVQTV
TKQFDDQKTSLQQMLYIADNLAYFP
YVVQDEPLYLIHQIDLLISMAGTHLL
ATFKEHIKPSDKEGDVLEDDDDVED
PEVLFNRLPEDLTEIICITSAQACM
LLLILKDHLKEMYAITDSKISRYSPSE
QKLYEKAVTRKSVNDFNPKTTIDVIK
KQMSQEKLSTDINFTLTKEEKLDLV
VKYLDFKQLMLKLPDDGDSDADES
RDKTMLNISASSDGVAFSSSAKNSH
SACDGYSLTVTDVVDVPM SHIAKAS
MLTSKPSGRKTNPVVRTKKRRKIDS
TDETSDAEYA