

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9VXE6	NU153_DROME	Drosophila melanogaster	Nuclear pore complex protein Nup153	16.334393	NaN	NaN	33925313;28604694	MEDAQE QRESSQAELAKKHPLTPLK ELPEESEEEEEDESSAGALRES SNNSIMGKMKMRVSSILPASLSGWF SPSSKDGNDALSSPANLRQSQPRQS NGRLTTKRKRGRRRIMLAEVDADAA DDLDDGSDAKGLNYEEVALADNIAE HDLAAEDEQTRRSEYNVFLLRKRAG AVAAAGGDEDEAEDELEEDDEDG DEEDDDEEQENLQSSAAVQTKRRR LELETPVNLPMRRLPLLSSTPAAPL AAATSSSSSQMYKGVSHIAPHRRNH LNLYGSQRQREPAYNFFTGNAAEG STGDLPHSIRSLNIPFGSSTATSY NNSLSSLPNHKRPSLIGKQTHRRDL TMDETGTGPAMSSEEHLNHLLRISR TNNNTSNNNNNNNNNNNNVIETKT RRSELSAAAGCGDSQSESDMNEY HDNGEGHDGLRPSHYNSNSNLEFY GNLQSSKSIFNRSNTAAQQSHRNST WSLNSLTQRRRFNASIYGSTSALSD SRLLSGSASNSGSASASSSPFYQGR TFGGNSGNNRFLFSRNLSSSAASSM LGLNSAGSSPAHQLHASMTGGIGYG MKAVDMRPSDSGSLAETSVGQGGG KKPGTGLSNTTMRILNLESYSTPLI DAKRMGSSIKEHQSSRQQRQGPAT PYLRSTSASRNVSPNHINELAELRS NKLLVPTMQQLLERRRLHRVTQNS RDVVHSQNVVRAGGENNQEKPKPTA PYVAPIDQSANHTQHTNKMRSLSH QTRNKETRTAEAEAPPPLDLPQISFP DMASAPKFDLIKPTVPVSKPSTTD PIQSSKSSNTNLSTTNSKQMPNFLA NPQPAAPIVNFAANGNVSAISKPSKR TFTFSEPTPLSNFQENCIPKPKINRK YTFSAPAPLDDLITNKQSQPTINGT PSSKEWECDCMVRNKPEINKCVA CETAKPVASAAPVQAPLPPSTAAIDT QSFVGFGRFDRFKSTTAWECDACML SNKAEASKCIACETPRKTVPKVN FSPLITNAKSNEWECVCLVRNKVE VSKCVACESAKPGATMALPATSNIAV ATPSIITDGFDRFKKSATAWECDAC MLSNKAEASKCIACETPRKSSTPIAN SSYPSINNNLPAGSGFDISFTRKAN MWECQTCLVMNKSSDEECIACQTP NSQARNSNSEALISSISSSSASFSG SLSRPSSRSSSGSTSTCGSVCSGSIV SISSTTESAKLSAKKVPKPDAGFQ

QLVAAQKTSTWECEACLAKNDMSR
KTCICCEQMMPEAFNPAATTANSAA
SSVPKFRFGFSHVKEVVKPSVETTTT
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TTSTSTANAIASSFAPTSTPQLFGNW
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WSSNGDAAKSNSAAVGSAAVPSSSA
STMATPIFGSSSMFGPSSSSNNTTST
STTSLPFGSAATTAATTPAGGNAALT
GLFGNVGNLAGVGAPVATTPAATA
AAPLTNIFGNPTPVAAAAPVFGSGST
IPSAGFGAPAAAAPLAAPALPGAFNF
GGATAATPAASSAPFVFGSSTNEPLA
KPSFNFTGSAASSTAPAPAFNFTANT
AATNNPSGGDSTPNANALFQFSATS
TAPANIFAFNPPAAGNSAQSSQTAR
RKIRAPVRRLLPPR