

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q868Z9	PPN_DROME	Drosophila melanogaster	Papilin	25.496522	NaN	NaN	33925313	MDLSRRLCSTALVAFIVLASIHDSQS RFPGLRQKRQYGANMYLPSSVTPG GEGNDPDEWTPWSSPSDCSRTCCG GVSQOTRECLRRDDRGEAVCSGGSR RYFSCNTQDCPEEESDFRAQQCSRF DRQQFDGVPFYEWVYPYTNAPNCEL NCMPKGERFYRQREKVVVDGTRCN DKDLDCVNGECMPVGCMMMLGS DAKEDKCRKCGDGGSTCKTIRNTITT KDLAPGYNDLLLLPEGATNIRIEETV PSSNYLACRNHSGHYLNGDWRIDF PRPMFFANSWWNYQRKPMGFAAP DQLTCSGPISSESLFVMLVQEKNISL DYEYSIPESLSHSQQDHTHTWHHF NACSASCGGGSQSRKVTNNRITLA EVNPSLDCQKSKPVEEQACGTEPCA PHWVEGEWSKCSKCGSDGFQNRS ITCERISSSGEHTVEEDAVCLKEVGN KPKATQECNRDVKNCPKYHLGPWT PCDKLCGDGKQTRKVTCFIEENGHK RVLPEEDCVEEKPETEKSCLLTPCEG VDWISQWSGCNACGQNTETRTAIC GNKEGKVPPEEFCEPEVPTLSRCK SPKCEAQWFSSEWSKCSAPCGKGV KSRVICGEFDGKTVTPADDDSKCNK ETKPESEQDCEGEEKVCPGEWFTGP WGKCSKPCGGGERVREVLCLSNGT KSVNCDDEEKVEPLSEKCNSEACTED EILPLTSTDKPIEDDEEDCEDGIELI SDGLSDDEKSEDVIDLEGTAKTETTP EAEDLMQSDSPTPYDEFESTGTTFE GSGYDSESTTDSGISTEGSGDDEETS EASTLSSSTDSGSTSSDSTSSDSSS SISSDATSEAPASSVSDSSDSTDAST ETTGVSDSDTDVSSSTEASASESTDV SGASDSTGSTNASDSTPESSTEASSS TDDSTDSSDNSSNVSESSTEASSSS VSDSNDSSDGDGVSSTTENSSDS TSDATSDSTASSDSTSDQTTETT PESSTDSTESSTLDASSTTDASSTSE SSSESSTDGSSTTNSASSETTGLSS DGSITDATTAAASDNTDITDGSSTDES TDGSSNASTEGSTEGASEDTTISTES SGSTESTDAIASDGSSTTEGSTVEDLS SSTSSDVTSDSTITDSSPSTEVSGST DSSSSTDGSSTDASSTEASSTDVTES TDSTVSGGTSDDTESGPTTEESTTEGS TESTTEGSTDSTQSTDLSTSDIWS TSDKDDESESSTPYSFDFSEVTKSKPR KCKPKKSTCAKSEYGCCPDGKSTPK GPFDEGCPIAKTCADTKYGCCLDGV SPAKGKNNKGCPSQCAETLFGCCP DKFTAADGENDEGCPETITVPPPTT TEETQPETTTEIEGSGQDSTTSEPDT KKCSFSEFGCCPDAETSAKGPDFE GCGLASPVAKGCAEENGCCPDGQT PASGPNNEGCSGCTRERFGCCPDSQ TPAHGPNKEGCCLDTQFGCCPDNIL AARGPNNEGCECHYTPYGCCPDNK SAATGYNQEGCACETTQYGCCPDKI TAAKGPKHEGCPETTQFGCCPDGL TFAKGPHHHGCHCTQTEFKCCDDE KTPAKGPNGDGCTCVESKFGCCPDG VTKATDEKFGGCENVQEPQKACGL PKETGTCNNYSVKYYFDTSYGCCAR FWYGGCDGNDNRFESAECKDTCQ DYTGKHVCLLPKSAGPCTGFTKKWY FDVDRNRCEEFQYGGCYGTNNRFD SLEQCQGTCAASENLPTCEQPVESG PCAGNFERWYDNETDICRPFTYGG CKGNKNNYPTHEACNYNCRQPGVL KDRCALPKQTGDCSEKLAKWHFSES EKRCVFFYYSGCGGNKNNFPLESC EDHCPQVAKDICEIPAIEVGECCANY	None	None	None	None	None	None	None		

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ENRFPTEESCLARCDRKEPTTTTPA  
TRQPQSRQDVCDEEPAPGECSTWVL  
KWHFDRKIGACRQFYGNCGGNGN  
RFETENDCQQRCLSQEPPAPTPRA  
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GGNDNRFATEEESARCSVNIDIRIG  
ADPVEHDTSKCFLAFEPGNCYNNVT  
RWFYNSAEGLCDEFVYTGCGGNAN  
NYATEEEECQNECNDAQTTICALPPVR  
GRCSDLRWRWYFDERSGECHEFEFT  
GCRGNRRNFVSQSDCLNFCIGEPVV  
EPSAPTYSVCAEPPEAGECDNRRTA  
WFDYSENMACTAFTYTGCGGNGNR  
FETRDQ CERQCGEFKGVDVCNEPVT  
TGPCTDWQTKYYFNTASQACEPFTY  
GGCDGTGNRFSDLFECQTVCLAGRE  
PRVGSKEICLLPVATGRCNGPSVH  
ERRWYDDEAGNCVSIYAGCSGNQ  
NNFRSFEACTNQCRPEPNKQDNEI  
GQNPCTFD AECQELRCYGVRRVA  
ARSQPECTQCICENPCEGYSCPEGO  
QCAIDVASSDDRQFAPVCRDIYKPGE  
CPALSANASGCARECYTDADCRGDN  
KCCSDGCGQLCVH PARPTQPPRTQA  
PVVSYPGDARAALPEKAHELDVQT  
AIGGIAVLRFCFATGNPAPNITWSLKN  
LVINTNKG RYVLTANGDLTIVQVRQT  
DDGTYYCVASNGLGEPVRRVALQV  
TEPVSQPAYIYGDKNVTQIVELNRPA  
VIRCPAGGFPEPHVSWWRNGQMFG  
LKNNLMARDYSLVFNSIQLSDLGLY  
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LSP EEEQYMQYVLPATRPVTRQRP  
YPYRPTRPAYVPEPTVNVHAVLALEP  
KNSYTPGSTIVMSCSVQGYPEPNVT  
WIKDDVPLYNNERVQITYQPHRLVL  
SDVTSADSGKYTCRASNAYTYANGE  
ANVSIQSVVPVSP ECVDNPYFANCK  
LIVKGRYCSNPYYTQFCRCSCTLAGQ  
VASPPLHPNAV