

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
A1Z6H7	PO210_DROME	Drosophila melanogaster	Nuclear pore membrane glycoprotein 210	11.65895	NaN	NaN	33925313	MDSILCMILLILVRNHASEAARLNH PRVLLPIFEDKAINFTLEVDEPNKYK WSSSRQDLISVMPIYKGFSECAYQAV VTVRTHDRRRNTAIVFAEEVQTGET LRSDVIVDVIAASLNVRTATRQLYLEE APAMFELHAFDEQNEFFTLEGIEF DWEILEPGSKRPTAMRYLTFTDSPY HTVPPTIEKFEADGKKGHMILLEGIN TGTAKVTIAMPQAEYKHVRPVEVYIS VLANIIIPESEVTIMAGDSVSFRILQL KMDRLHVIDNNQYILEVEDSSIAYL RGNSATGAALGRTQVFLRDRNMAD SDEVQKGPSALLTVAYPNRLSISLLP HLNWVTVQGEHHAIALDLFAADGQ KITMGTKYSINSEVDESFFAIVDRTR NGSRLFGQAKKEGITQVYGSYKDL VQAEIQIFEELQLAPTKVVLVYDPNS LKPLKLQFHASGGDNNYAWFSGNP QVIQIDTQGQATTEIRDVKSAYVNQE VLKDGKLTAAHTTVKVALSKNQKIS RVAHIYFLPPERLQITRSNFETALKD FVHVHVGVIYARINNSEVPYTSCDNL HFQLDFSQPILQLEGNEGAEAAHEA CHVLRRLRATAVGTTSLRVSYMMDK VLYDIIDLIVFEPLVVLNPIENEVLP VGSSRNIIYANGPQRSFTVAEIIQST AFDEKILKVSLEFDTQNLITAFVTL CRELGETQFTYRVHNSLPTSSFALY QSEVTTKVHCVRPRFLKLYARHNL DSCPLEKRTSLLFLKDPENKIEIEIEV HDSNNRRLMNISSLGLDWEFSAGE ERYQKNIIPHRQISELEFNHGVTLPS RDLLVTLSEVATNFRIKGTVSQYND KLLAQHGIHAERPPFGIKNPQTGLIY TPLIENEIRLHAVNSTLLPKDYMSIF LASGYSERIPIAQSGYLQLELSEAGI VQVEYNENTRILVLTPLRLGHVRLEL TDRCLMNEPSHLSISVVGIGAIEVVS MDRLERTTRIEAIVRLFDTNDNLLL DQSKLSAYDLSEVVADQSILSVRLGE QENVGPGIEIRYTITGNQVGETKILFQ SGKGIYKVASDPLNIQVFAPIRLFRD STLVVGSSIQVYFHGGPHNTNMIIS VEKEQVATISSTVVTAKLGTTKIVG KLLKNPVTGKDEVVSQDSVEVHV ALKGVQIRTPLVRIHSGAVMPATLW GLSDLSPMILGTLQNTKISWKVSQP QVVEIFNVFTTAGIEYQSGDLISVRV RALNPGKATITASVTLADGTILPPATV

