

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P32521	PAN1_YEAST	Saccharomyces cerevisiae	Actin cytoskeleton-regulatory complex protein PAN1	9.365641	NaN	T241;T570;S747;S757;T993;T995;S1003;S1180;S1250;S1253;S1281;T1321	33229814	<p> MYNPNYQQQGMGYQQQQQQQQQP NGFYPPQQQGGQSSNQPGQPQPQQ QMAFNQPQATGIGGMPQSFQNSFS SMPQQPQTGYNNNGNNGSVYGNQ NFGQQPQQQQQAKPQHTGYVPNS SMPMMNTTGTMPPPNPAQQPQLQS IQPQGTGYQAANTANVHSVQPLQS QGTGYVSTPNLISSNQTTQPLQAQ GTGYYSQPQQVPPQQAQSLQPLK PQQTGFYLPQNPQAPLEPLKPTATG FVNSFANNGLNNDIKIPAIRLSFITA QDQAKFETLFRSIVTNGSNTVSGAN CRKILMRSGLPPSQLARIWTLCDTSK AGELLFPEFALAMHLINDVLQGDTP YELDSKTKNEVSSFIDAINLSIANQD SSANDAPKTPFDEFITAGVQNLQPQ PTGYMPQTSFGIPLQSQITGGGVASA LNPQSTGFMPTTFNMSMNTGTPG LNPQITGGAPASMQPNTGNALQPQ TTGMMPQTTGMMPQTTGMMPQTS FGVNLGPQLTGGALQSQYGGYGSV MPQQSGPASPMPNLSFNQQGLQSQL TGLQPQPTGFLPPSNFSATMPLTAQ KTGFGNNEIYTKSNFNNDLIDNSSQ DKISTEEKSLFYKIFETFDTONKGLL DSPTAVEIFRKSGLNRADLEQIWNL CDINNTGQLNKQEFALGMHLVYGK LNGKPIPNVLPSSLIPSSTKLLDNLK NQLKTEPTTTKEKPSFGKIDALS YKN NDDDVLPNYRNRKRVYSAKNEEQS SFSSPSAKSVNHSSSTLQDDISVDK TVEKKTAKPKYAGFSREINLKNIASL ENEIKNISNPENCYDSSIPSDLTSRF DAIIAKLPNLFNEISTIDNEITNAKIQ LYRKNPSSIIGSGPNGEITENDRKK AKSRALLRARMSALTGKSTESDLSL SMEDEQQSAEIKRIQQENGKNOEII KDIRSSISDISASLKSTMTGSMNISM QEFERWEFGIGLEDGVREFLDDLKS NSNKSVTESSPFVPSSTPTPVDDRSS SPSYSQFKTAEERAAYLKEQAKKRM KEKLAKFDKNRRNVTQSSRSISEN SREQPQQIAGSSNLVEPRATPFQEE KYVEVAQPTQPVQSTQPVQPTQPVQ PTQPVQPTQPVQPTQPVQPTQPVQN VYNAKQESDDEDEDDEEKRLQEELK RLKLLKKADKEKRLAALRKQIEDAQ NESDEEETNGKDNFGGHVNVPPQAA PVAPSAAFSQNSTNAPRSVHAAVTP </p>

