

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
O95613	PCNT_HUMAN	Homo sapiens	Pericentrin	16.435419	T57;T345;T2439;T3000;T3005;S3006	S44;S188;T191;S610;S682;S1245;S1653;S1712;S2044;S2177;S2192;S2225;S2226;S2327;S2352;S2355;S2477;S2486;S3302	34019948;37340703;23301498;31492838;29351928;30379171	MEVEQEQRRRKVEAGRTKLAHFRRQ RKTKGDSHSEKKTAKRKGSAVDAS VQEEESPVTKEDSALCGGGDICKSTS CDDTPDGAGGAFAAQPEDCDGKEK EDLEQLQQKQVNDHPPEQCGMFTV SDHPPEQHGMFTVGDHPPEQRGMF TVSDHPPEQHGMFTVSDHPPEQRG MFTISDHQPEQRGMFTVSDHTPEQ RGIFTISDHPAEQRGMFTKECEQEC ELAITDLESGREDEAGLHQSQAVHG LELEALRLSLSNMHTAQLELTQANL QKEKETALTELEMLNSRRAQELAL LQSRQQHELELLREQHAREKEEVVL RCGQEAELKEKLOSEMKNQIVK TLKEDWESEKDLLENLRKELSAKH QSEMEDLQNFQKELAEQRAELEKI FQDKNQAERALRNLESHHQAAIEKL REDLQSEHGRCLLEDLEFKFKESEKE KQLELENLQASYEDLKAQSQEEIRR LWSQLDSARTSRQELSELHEQLLAR TSRVEDLEQLKQREKTOHESELEQL RIYFEKKLRDAEKTYQEDLTLQORL QGAREDALLDSVEVGLSCVGLLEKP EKGRKDHVDELEPERHKESELPRFQA ELEESHRHQLEALESPLCIQHEGHV SDRCCVETSALGHEWRLEPSEGHS QELPWWHLQGVQDGDLEADTERAA RVLGLETEHKVQLSLLQTELKEEIEL LKIENRNLYGKLQHETRLKDDLEKV KHNLIEDHQKELNNAKQKTELMKQ EFQRKETDWMKQEEELQREAEEL TLMLELREKAESEKQTIINKFELRE AEMRQLQDQAAQILDLESLTEQQ GRLQOLEQDLTSDDALHCSQCGREP PTAQDGELAALHVKEDCALQLMLAR SRFLEERKEITEKFSAEQDAFLQEAQ EQHARELQLLQERHQQLLSVTAEL EARHQAAALGELTASLESKQGALLAA RVAELQTKHAADLGALET'RHLSSLD SLESCYLSEFQTIREEHRQALELLRA DFEEQLWKKDSLHQTLTQELEKLK RKHEGELQSVRDHLRTEVSTELAGT VAHELQGVHQGFEGSEKKTALHEK EETLRLQSAQAQPFHQEKEKESLSLQ LQKKNHQVQQLKDQVLSLSHEIEEC RSELEVLQRRERENREGANLLSM LKADVNLSHSERGALQDALRLLGL FGETLRAAVTLRSRIGERVGLCLDDA GAGLALSTAPALEETWSDVALPELD RTLSECAEMSSVAEISSHMRESFLM SPESVRECEQPIRRVFSLSLAVDGL MEMALDSSRQLEEARQIHSRFEKEF SFKNEETAQVVRKHQELLECLKEES AAKAELELELHKTQGTLEGFKVETA DLKEVLAGKEDSEHRLVLELESRR QLQAAQEQAAALREECTRLWSRGE ATATDAEAREAALRKEVEDLTKEQS ETRKQAEKDRSALLSQMKILESELE EQLSQHRGCAQAEAVTALEQQVAS LDKHLRNQRQFMDEQAAEREHERE EFQQEIQRLEGQLRQAAKPOPWGPR DSQQAPLDGEVELLQQLREKLDEF NELAIQKESADRQVLMQEEIEIKRLE

EMNINIRKKVAQLQEVEEKQKNIVK  
GLEQDKEVLKKQMQSSLLASTLQS  
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VLTVCQRQLQSELLVKNEMRSLSE  
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AQGKAPRPGPRARQPQSPPTRESP
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HSGGRATPSPNSRLERSLTASQDPE
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