

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membran
Q14207	NPAT_HUMAN	Homo sapiens	Protein NPAT	26.287343	T364;S389;S402	S207;S554;S599;S775;S779;S1100;S1151;S1200;S1254;T1270;T1350	23301498;38253038	MLLPDVARLVLVGLYQENLISTCQ TFLESSLKEYAEHCTDEGFIPACL LSLFGKNTTILNEVYVAMTKKTSN NVPAIMSSLWKLDHTLSQIRSMQS SPRFAGSQRARTRTGIAEIKRQRKLA SQTAPASAEALLTLPYLSGQFTTPPST GTQVTRPSGQISDPSRSYFVVVNH QSQDVTVTGALNVIPGAQEKKAHA SLMSPGRRKSESQRKSTTLGSPHSTI RNFQDPNFAVEKQMVLENAREKIL SNKSLQEKLAENINKFLTSDNNIAQ VPKQTDNNPTEPETSIDEFLGLPSEI HMSEEAIQDILEQTESDPAFQALFDL FDYGKTKNNKNISQSISSQPMESNP SIVLADETNLAVKGSFETEESDQSG QPAFCTSYQNDPLNALKNSNNHD VLRQEDQENFSQISTSIQKAFKTA PTEQKCDIDITFESVPLNDFNQRG NSNAECNPHCAELYTNQMSSTETEM AIGIEKNLSLSSNVPSESQLOPDQDI PITSFVSLGCEANNENLILSGKSSQL LSQDTSLTGKPSKKSQFCENSNDTV KLKINPHGSKSSDSSEVHKSKIEINV LEPVMSQLSNCQDNCLQSEILPVS VESSHLNVSGQVEIHLGDSLSTKQ PSNDSASVELNHTENEQAQSKSENS QEPSSSVKEENTIFLSLGGNANCEK VALTPPEGTPVENSHSLPPESVCSV GDSDHPESQNTDDKPSNNSAEIDAS NIVSLKVIIISDDPFVSSDTELTSVSS INGENLPTIILSSPTKSPKNAELVKC LSSEETVGAVVYAEVGDASAMEQSL LTFKSEDSAVNNTQNEQDIAFANSV TPCVSKDGGYIQLMPATSTAFGNSN NILIATCVTDPTALGTSVSNVVVL PGNSAPMTAQPLPPQLQTPPRSNSV FAVNQAVSPNFSQGSIIIASPVQPV LQGMVGMIPVSVVQNGNNSFSTPP RQVLHMPLTAPVCNRSIPQFPVPPK SQAQGLRNKPCIGKQVNNLVDSG HVGCHAQKTEVSDKSIATDLGKKS EETTVPPEESIVPAAKPCHRRVLCF DSTTAPVANTQGPNHKMSVQNKER NAVSPNLDSPNVSTLKPPSNNAI KREKEKPLPKILSKSESAISRHTTIR ETQSEKKVSPTEIVLESFHKATANKE NELCSDVERQKNPENSKLSIGQQNG GLRSEKSIASLOEMTKKQGTSSNNK NVLVGTAVKDLKQEQTKSASSLITT EMLQDIQRHSSVSRDLSSDLPVPR TPGSGAGEKHKEEPIDIHKAPSSRRFS EDSSTSKVMVPPVTPDLACSPASET GSENSVNMAAHTLMILSRAAISRTT SATPLKDNTQQFRASSRSTTKKRKIE ELDERERNRPSKSLTNSSIPMKK KKIKKKLPSSFPAGMDVDKFLLSL HYDE	True	False	1.077	5.0	1.19	0.872	0.666	1.373