	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	A0A1I9LPG3	A0A1I9LPG3_ARATH	Arabidopsis thaliana	histone acetyltransferase	27.144759	NaN	NaN	28154133	MWRTATSKEDYLDFRSFDVRVESTL KQLLSQRRANPSSSVSTMVQTPGVS HGWGQSYTATPMVDTSKFNSSNNL SDATTETGRLLPTNRMYRGTINNGR QELSAVGQMIPTPGFDNSANADVYQ SHRNEEYSGDGGKLLATGSDFGNPS QLQKQRPTGSNDLMGYNLDHQLGG GFRSNIHQNTSGMTSIPLNAGVGMS GNNVHLANVPRSSEGVLSSTHFSTF SQPSQQPVEQLQVSHVNRYSMSNS GTFVSGNLYGVQTSSGSIETAVDMN SMSLNSMRRVDTSFGSQSGLQNNP LLKPHLRHQFENGNFQSSSNSKEN LAQVSHRPLERQFNQQAHYGQYHQ QELLMNNDAYRQSQPASNLVSQVK NEPRVEYYNEAFQMQAINKVEPSKP QNQYKQNTVKDEYVGAQSAPVSSSQ LKMSPSFPPQTHQTQQVSQWKDSS SLSAGVQPVSGLGQWHSSSQNLTPI SKNSNEEREHFGVRFHKQHEGTNN SSSVRESTNCLTVAPSGTLDVPHLPV GINVLSKQLNGDCGLSYKNQRRWLL FLLHVRKCNAAEDNCESKYCFTAKT LLKHINCCKAPACAYQYCHQTRQLI HHYKHCGDEACPVCVFVKNFKEKQ KEKFTFLQRAEPSSASLNHGPKESF ESMRTSSERDSEAPFVVDDLQPSPK RQKVEKPSQFAYPDTQGNPATISAG VSQAHFSMGLQEKDRLPSDVCKPVR SNVPMNADSSDSSRRLVPVSRELEK PVCKDTHMGRHGVKSALDGESLRL SKQEKPKRMNEISAPKEENAEQSLG VVSASNCGKSKIKGVSLIELFTPEQV EEHIRGLRQWVGQSKTKAEKNKAM GLSMSENSCQLCAVERLAFEPTPIYC TPCGARVKRNAMHYTVVAGESRHY VCIPCYNEARANTVSVDGTPVPKSRF EKKKNDEEVEESWVQCDKCQAWQ HQICALFNGRRNHGQAEYTCPNCYI QEVEQGERKPVSQNVILGAKSLPAS TLSNHLEQRLFKKLKQERQERARLQ GKSYEEVPGADSLVIRVVASVDKILE VKPRFLDIFREDNYSSEFPYKSKAILL FQKIEGVEVCLFGMYVQEFGTDSAS PNQRRVYLSYLDSVKYFRPDVRTVS GEALRTFVYHEILIGYLDYCKKRGFS SCYIWACPPLKGEDYILYCHPEIQKT PKTDKLREWYLAMLKKASKEKVVVE CTNFYDHFFVQSGECRAKVTAARLP YFDGDYWPGAAEDLIDQMSQEEDG KKSNRKLMPKKVISKRALKAVGQLD LSVNASKDLLLMHKLGEIILPMKED FIMVHLQHCCKHCCTLMVSGNRWV CNQCKNFQICDKCHEVEENRVEKE KHPVNQKEKHVLYPVAIDNIPTEIKD NDDILESEFFDTRQAFLSLCQGNHY QYDTLRRAKHSSMMILYHLHNPTVP AFAMACAICQQELETAQGWRCEVCP DYDVCNACYSKGINHPHSIISRPSAT DSVVQNTQTNQIQNAQLREVLLHV MTCCTAQCQYPRCRVIKGLIRHGLV CKTRGCIACKKMWSLFRLHSRNCR DPQCKVPKCRELRAHFSRKQQQADS RRRAAVMEMVRQRAADTTASTPE	None	None	None	None	None	None	None	None	None