	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	A0A178VGI0	A0A178VGI0_ARATH	Arabidopsis thaliana	histone acetyltransferase	27.144759	NaN	NaN	28154133	MAQGQNRTVLQSGQMHNSVVASAS SVSSQPNMVNGISQDTLLLRQEML NRTYAWLQQRQPSKTDDASKAKLSE VAKRLESAMWRTATSKEDYLDFRSF DVRVESTLKQLLSQRRANPSSSVST MVQTPGVSHGWGQSYTATPMVDTS KFNSSNNLSDATTETGRLLPTNRMY RGTINNGRQELSAVGQMIPTPGFDN SANADVYQSHRNEEYSGDGGKLLAT GSDFGNPSQLQKQRPTGSNDLMGY NLDHQLGGGFRSNIHQNTSGMTSIP LNAGVGMSGNNVHLANVPRSSEGV LSSTHFSTFSQPSQQPVEQLQVSHV NRYSMSNSGTFVSGNLYGVQTSSGS IETAVDMNSMSLNSMRRVDTSFGS QSGLQNNPLLKPHLRHQFENGNFQ SSSNSKENLAQVSHRPLERQFNQQ AHYGQYHQQELLMNNDAYRQSQPA SNLVSQVKNEPRVEYYNEAFQMQAI NKVEPSKPQNQYKQNTVKDEYVGA QSAPVSSSQLKMSPSFPPQTHQTQQ VSQWKDSSSLSAGVQPVSGLGQWH SSSQNLTPISKNSNEEREHFGVRFH KQHEGTNNSSSVRESTNCLTVAPSG TLDVPHLPVGINVLSKQLNGDCGLS YKNQRRWLLFLLHVRKCNAAEDNC ESKYCFTAKTLLKHINCCKAPACAYQ YCHQTRQLIHHYKHCGDEACPVCVF VKNFKEKQKEKFTFLQRAEPSSASL NHGPKESFESMRTSSERDSEAPFVV DDLQPSPKRQKVEKPSQFAYPDTQG NPATISAGVSQAHFSMGLQEKDRLP SDVCKPVRSNVPMNADSSDSSRRLV PVSRELEKPVCKDTHMGRHGVKSAL DGESLRLSKQEKPKRMNEISAPKEE NAEQSLGVVSASNCGKSKIKGVSLIE LFTPEQVEEHIRGLRQWVGQSKTKA EKNKAMGLSMSENSCQLCAVERLA FEPTPIYCTPCGARVKRNAMHYTVV AGESRHYVCIPCYNEARANTVSVDG TPVPKSRFEKKKNDEEVEESWVQCD KCQAWQHQICALFNGRRNHGQAEY TCPNCYIQEVEQGERKPVSQNVILGA KSLPASTLSNHLEQRLFKKLKQERQ ERARLQGKSYEEVPGADSLVIRVVAS VDKILEVKPRFLDIFREDNYSSEFPY KSKAILLFQKIEGVEVCLFGMYVQEF GTDSASPNQRRVYLSYLDSVKYFRP DVRTVSGEALRTFVYHEILIGYLDYC KKRGFSSCYIWACPPLKGEDYILYCH PEIQKTPKTDKLREWYLAMLKKASK EKVVVECTNFYDHFFVQSGECRAKV TAARLPYFDGDYWPGAAEDLIDQMS QEEDGKKSNRKLMPKKVISKRALKA VGQLDLSVNASKDLLLMHKLGEIILP MKEDFIMVHLQHCCKHCCTLMVSG NRWVCNQCKNFQICDK	None	None	None	None	None	None	None	None	None