	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	A0A1P8ARU4	A0A1P8ARU4_ARATH	Arabidopsis thaliana	histone acetyltransferase	27.144759	NaN	NaN	28154133	MNVQAHMSGQRSGQVPNQGTVPQ NNGNSQMQNLVGSNGAATAVTGAG AATGSGTGVRPSRNIVGAMDHDIMK LRQYMQTLVFNMLQQRQPSPADAA SKAKYMDVARRLEEGLFKMAVTKE DYMNRSTLESRITSLIKGRQINNYNQ RHANSSSVGTMIPTPGLSQTAGNPN LMVTSSVDATIVGNTNITSTALNTGN PLIAGGMHGGNMSNGYQHSSRNFS LGSGGSMTSMGAQRSTAQMIPTPGF VNSVTNNNSGGFSAEPTIVPQSQQQ QQRQHTGGQNSHMLSNHMAAGVR PDMQSKPSGAANSSVNGDVGANEK IVDSGSSYTNASKKLQQGEGYSTTNP DPFDGAITSAGTGTKAHNINTASFQP VSRVNSSLSHQQQFQQPPNRFQQQ PNQIQQQQQQFLNQRKLKQQTPQQ HRLISNDGLGKTQVDSDMVTKVKCE PGMENKSQAPQSQASERFQLSQLQ NQYQNSGEDCQADAQLLPVESQSDI CTSLPQNSQQIQQMMHPQNIGSDS SNSFSNLAVGVKSESSPQGQWPSKS QENTLMSNAISSGKHIQEDFRQRIT GMDEAQPNNLTEGSVIGQNHTSTIS ESHNLQNSIGTTCRYGNVSHDPKFK NQQRWLLFLRHARSCKPPGGRCQD QNCVTVQKLWSHMDNCADPQCLYP RCRHTKALIGHYKNCKDPRCPVCVP VKTYQQQANVRALARLKNESSAVGS VNRSVVSNDSLSANAGAVSGTPRCA DTLDNLQPSLKRLKVEQSFQPVVPK TESCKSSIVSTTEADLSQDAERKDHR PLKSETMEVKVEIPDNSVQAGFGIKE TKSEPFENVPKPKPVSEPGKHGLSG DSPKQENIKMKKEPGWPKKEPGCP KKEELVESPELTSKSRKPKIKGVSLT ELFTPEQVREHIRGLRQWVGQSKAK AEKNQAMENSMSENSCQLCAVEKL TFEPPPIYCTPCGARIKRNAMYYTVG GGETRHYFCIPCYNESRGDTILAEGT SMPKAKLEKKKNDEEIEESWVQCD KCQAWQHQICALFNGRRNDGGQAE YTCPYCYVIDVEQNERKPLLQSAVLG AKDLPRTILSDHIEQRLFKRLKQERT ERARVQGTSYDEIPTVESLVVRVVSS VDKKLEVKSRFLEIFREDNFPTEFPY KSKVVLLFQKIEGVEVCLFGMYVQE FGSECSNPNQRRVYLSYLDSVKYFR PDIKSANGEALRTFVYHEILIGYLEYC KLRGFTSCYIWACPPLKGEDYILYCH PEIQKTPKSDKLREWYLAMLRKAAK EGIVAETTNLYDHFFLQTGECRAKVT AARLPYFDGDYWPGAAEDIISQMSQ EDDGRKGNKKGILKKPITKRALKASG QSDFSGNASKDLLLMHKLGETIHP MKEDFIMVHLQHSCTHCCTLMVTG NRWVCSQCKDFQLCDGCYEAEQKR EDRERHPVNQKDKHNIFPVSCIGLF WVRNCDEVLSLSLMRFSILLYRKR	None	None	None	None	None	None	None	None	None