	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	A8WHL8	A8WHL8_DROME	Drosophila melanogaster	[histone H4]-N-methyl-L-lysine(20) N-methyltransferase	28.06426	T888	NaN	40245499	MNIRFRPLKVDTQQLKAIVDDFIHTQ NYDIAIQRIYEGPWIPRHLKNKNKIA TKRLHDHIVRYLRVFDKDSGFAIEAC YRYTLEEQRGAKISSTKRWSKNDKIE CLVGCIAELTEAEEAALLHSGRNDFS VMYSCRKNCAQLWLGPAAYINHDC RANCKFLATGRDTACVKVLRDIEVG EEITCFYGEDFFGDSNRYCECETCE RRGTGAFAGKDDGLMLGLSMGLGL ASSGPGNNGGYRLRETDNRINRIKS RANSTNSTSNSNSNTNDSTGPSETS STNGLVASGGAGGATGAAMLPTPSQ QSTGGKEATAAVSLLEKKLPNVVVS PLTMKELRQKGMTKYDAEMIMANA AYQQQHHHQHHFHHHHHHHHHH HNHGQHASTGAEATAAVQQMAAM QKPGVGGTGAAGNAGATTVSSVAAG AGSEVNGGRSTSLRKSMRVNSTSSS ISTASADEVIAPVVAASISLPSKAPVV LMPRCKPAQMAIAALHQSQQRQLR RSERQKEKLTDGESSDTSSEQQKKE QKQQDHQLPQKMFSLAEEPQPEKS EEKQQEQQKRVTRNSAGRVGLVARL ATAHNNNIATTTNSSSSSNKATTITN CNNHNSNNSSGINHNSNLSSRLSV KSRKPAPSEASSIPSSTSSENQQQQA TRRSCSPTPAYKKNLLASFDPDPPST QGIKEQLKDESVTYSPVKQKRSRRA AALAAAQSIHCEALGGFPTGSTGSQ RKRAQAGEPTTSCSSTTISNVEPLLK TPERRLKLTLRMKRSPILDEVIELGT SLSNGGAGRGAPGSHREGTAGEGSV RSALNLTGSSSNGIEYEILRMEGISE HGNDDDEDEEEDDEEPAAEEEEEP PPKEELQLVNKKQRKKQRSRSRSSQ RRSPAPSSVYGTPQKKRLRLIFGNES HTIDIPPAAAEGSGSGLDDLNSSGGG GDESFNASYASSTSLTVNTSSSSTSS SSGVGGATSTSAEPVDSSTVGPPIAA QSPSSTTSSSFQSACTSTTNSNSYFP NGKQRAAGEDSYAMHYYQLGKFAG TPSPGQGQAIVSSSSGSSGGGGSGA GFLSMPKHTFGTCALLAPTSFACLQ NQPQISQQKTSSGGGAGVVPTSTST GAVTSHHHTNNHHGQK	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	None