

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochr
Q15652	JHD2C_HUMAN	Homo sapiens	Probable JmjC domain-containing histone demethylation protein 2C	44.405871	T650;S652;S655;T691;T699;T912;S913;S969;S1073;S1083;S1087;S1091;S1173;S1185;T1190;S1192;S1193;T1194;T1196;T1351;S1376;S1379;S1437;S1451;T1452;T1453;T1457;T1822;T2031;T2052;S2057;T2073;S2099	S317;S320;S373;S376;S475;S501;T505;S601;S617;S638;S639;S641;S652;S943;S1989	28657654;31492838;37340703;34019948;20305658;32574038;21740066;35254053;29237092;21158410;29351928;20068230;35138101;30397120;30620550;30059200;30379171;22661428;31637018;33214551;35289036;38253038;38665916	MAVETRAELVVGKRFLLCVAVGDEARSERWESGRGWSRWRAGVIRAVSHRDRSRNPDLAVYVEFDLLEWDKREWVKVYEDFSTFLVEYHLIWAQRNDPSQTQGSKSKQIQWPALTFKPLVERNIPSSVTAVEFLVDKQLDFLTEDSAFQPYQDDIDSLNPVLRDNPQLHEEVKVVVKEQKVQEIFMQGPYSLNGYRVRYRQDSATQWFTGITHHDLFTRTMIVMNDQVLEPQNVDPMSVMQMTFLDDVVHSLKGENIGITSRRRSRANQNVNAVHSHYTRAQANSRPRAMNSQAAPKQNTHQQQQORSIRPNKRKGSDDSSIPDEEKMKEEKYDIYSRGENPKGKKNKHLMNKRRRPEEDEKKNMRLRTD NVSDFSESSDSENSNKRIDNSSEQKPENELKNKNTSKINGEEGKPHNNEKAGEETLKNQPPWDQIQEDKKHEEAEKRKSVDTQLQEDMIIHSSEQSTVSDHNSNDLLPQECNMDKTHTMELLPKEKFSRPPPTPKCVIDITNDTNLEKVAQENSSTFGLQTLQKMDPNVSDSKHSIANAKFLETAKKDSQSWVSDVVKVDLTQSSVFNASSGNDHLNMEKEKYVSYISPLSAVSVMEDKLHKRSPPETIKSKLNTSVDTHTKIKSSPPEVVKPKITHSPDSVKSKATYVNSQATGERRLANKIEHELRSRCSFHPIPTRSSTLETTKSPLIIDKNEHFTVYRDPALIGSETGANHISPFLLSQHPFPLHSSSHRTCLNPGTHHPALTPAPHLLAGSSSQPTLPTINTHPLTSGPHHAAVHHPHLLPTVLPGVPTASLLGGHPRLESAHASSLSHLALAHQQQQQLLOHQSPHLLGQAHPASYNQLGLYPHIWQYPNGTHAYSGLGLPSSKWVHPENAVNAEASLRRNSPSPWLHQPTPVTSAAGIGLLSHIPVRPSSAEPHRPLKITAHSSPPLTKTLVDHHEEELERKAFMEPLRSVASTSAKNDDLNRSTGKDCHELRHFVDPVLNQLQRPPQETGERLNKYKEEHRRILQESIDVAPFTTKIKGLEGERENYSRVASSSSPKSHIHKQMDMVERSVDLYKMKHSVPQSLPQSNYFTLSNSVVENPPRSYPSKEVSNYIGDKQSNALA AAAANPQTLTSTFITSLSKPPPLIKHQPESEGLVGIPEHLPHQIASHSVTTFRNDCRSPHTLTVSSTNLRSMPALHRAPVFHPPIHHSLERKEGSYSSLSPTLTPVMPVNAVAGGKQVQSQKPTLPEPKDSQANFKSSSEQSLTEMWRPNNLSKEKTEWHVEKSSGKLAAMASVIVRPSSTKTDSMPAMQLASKDRVSERSSAGAHKTDCLKLAEAGETGRRIIPNVNSDSVHTKSEKNFQAVSQGSVPSSVMSAVNTMCNTKTDVITSAADTTSVSSWGGSEVISLNTILASTSSECVSSKSVSQPVAQKQECKVSTTAPVTLASSKTGSVVQPSGFGSTDFIHLKKHKAALAAAQYKSSNASETEPNAIKNQTLASLPLDSTVICSTINKANSVGNQASQTSQPNYHTKLLKAWLTRHSEEDKNTNKMENSGNSVSEIIPC SVNLIASSTSDIQNSVDKIIVDKYVKDDKNRRKAKRITYESGESGDSDES ESKSEQRTRKQPKPTYKKQNDLQKRKGEIEEDLKPNGVLSRSAKERSKLLQSNSTNGIPRSVLKDRWKVKKLQ TGESFLQDDSCCEIGPNLQKRECR LIRSKGEEPAPHSVFCRFYFRRLSFSKNGVVRIDGFSSPDQYDDEAMSLWTFENFEDDELDIETSKYILDIIIGDKFCQLVTSEKTLASVWKDADAKIAWKR AVRGVREMCDACEATLFNIHWVCQKCGFVCLDCYKAKERKSSRDKELYAWMKCVKQPHDHHMLMPTQIIPG SVLTDLLDAMHTLREKYGIKSHCHCTNKQNLQVGNFPTMNGVSOVLQNV LNHSNKISLCMPESQQQNTPPKSEKNGGSSPESDVGTDNKLTPPESQSP	True	False	2.428	4.615	1.481

