	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	Q9H9B1	EHMT1_HUMAN	Homo sapiens	Histone-lysine N-methyltransferase EHMT1	21.731224	NaN	S435;S483;S1004;S1048	23301498	MAAADAEAVPARGEPQQDCCVKTEL LGEETPMAADEGSAEKQAGEAHMA ADGETNGSCENSDASSHANAAKHT QDSARVNPQDGTNTLTRIAENGVSE RDSEAAKQNHVTADDFVQTSVIGSN GYILNKPALQAQPLRTTSTLASSLPG HAAKTLPGGAGKGRTPSAFPQTPAA PPATLGEGSADTEDRKLPAPGADVK VHRARKTMPKSVVGLHAASKDPREV REARDHKEPKEEINKNISDFGRQQL LPPFPSLHQSLPQNQCYMATTKSQT ACLPFVLAAAVSRKKKRRMGTYSLV PKKKTKVLKQRTVIEMFKSITHSTVG SKGEKDLGASSLHVNGESLEMDSD EDDSEELEEDDGHGAEQAAAFPTED SRTSKESMSEADRAQKMDGESEEE QESVDTGEEEEGGDESDLSSESSIK KKFLKRKGKTDSPWIKPARKRRRRS RKKPSGALGSESYKSSAGSAEQTAP GDSTGYMEVSLDSLDLRVKGILSSQ AEGLANGPDVLETDGLQEVPLCSCR METPKSREITTLANNQCMATESVDH ELGRCTNSVVKYELMRPSNKAPLLV LCEDHRGRMVKHQCCPGCGYFCTA GNFMECQPESSISHRFHKDCASRV NNASYCPHCGEESSKAKEVTIAKAD TTSTVTPVPGQEKGSALEGRADTTTG SAAGPPLSEDDKLQGAASHVPEGFD PTGPAGLGRPTPGLSQGPGKETLESA LIALDSEKPKKLRFHPKQLYFSARQG ELQKVLLMLVDGIDPNFKMEHQNK RSPLHAAAEAGHVDICHMLVQAGA NIDTCSEDQRTPLMEAAENNHLEAV KYLIKAGALVDPKDAEGSTCLHLAAK KGHYEVVQYLLSNGQMDVNCQDDG GWTPMIWATEYKHVDLVKLLLSKGS DINIRDNEENICLHWAAFSGCVDIAE ILLAAKCDLHAVNIHGDSPLHIAARE NRYDCVVLFLSRDSDVTLKNKEGET PLQCASLNSQVWSALQMSKALQDS APDRPSPVERIVSRDIARGYERIPIPC VNAVDSEPCPSNYKYVSQNCVTSPM NIDRNITHLQYCVCIDDCSSSNCMC GQLSMRCWYDKDGRLLPEFNMAEP PLIFECNHACSCWRNCRNRVVQNG LRARLQLYRTRDMGWGVRSLQDIPP GTFVCEYVGELISDSEADVREEDSYL FDLDNKDGEVYCIDARFYGNVSRFI NHHCEPNLVPVRVFMAHQDLRFPR IAFFSTRLIEAGEQLGFDYGERFWDI KGKLFSCRCGSPKCRHSSAALAQRQ ASAAQEAQEDGLPDTSSAAAADPL	None	None	None	None	None	None	None	None	None