	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	Q9P267	MBD5_HUMAN	Homo sapiens	Methyl-CpG-binding domain protein 5	18.894839	S530	NaN	34019948;35254053	MNGGKECDGGDKEGGLPAIQVPVG WQRRVDQNGVLYVSPSGSLLSCLEQ VKTYLLTDGTCKCGLECPLILPKVFN FDPGAAVKQRTAEDVKADEDVTKLC IHKRKIIAVATLHKSMEAPHPSLVLT SPGGGTNATPVVPSRAATPRSVRNK SHEGITNSVMPECKNPFKLMIGSSN AMGRLYVQELPGSQQQELHPVYPRQ RLGSSEHGQKSPFRGSHGGLPSPAS SGSQIYGDGSISPRTDPLGSPDVFTR SNPGFHGAPNSSPIHLNRTPLSPPSV MLHGSPVQSSCAMAGRTNIPLSPTL TTKSPVMKKPMCNFSTNMEIPRAM FHHKPPQGPPPPPPPSCALQKKPLTS EKDPLGILDPIPSKPVNQNPVIINPTS FHSNVHSQVPMMNVSMPPAVVPLP SNLPLPTVKPGHMNHGSHVQRVQH SASTSLSPSPVTSPVHMMGTGIGRIE ASPQRSRSSSTSSDHGNFMMPPVG PQATSSGIKVPPRSPRSTIGSPRPSM PSSPSTKSDGHHQYKDIPNPLIAGIS NVLNTPSSAAFPTASAGSSSVKSQPG LLGMPLNQILNQHNAASFPASSLLS AAAKAQLANQNKLAGNNSSSSSNS GAVAGSGNTEGHSTLNTMFPPTAN MLLPTGEGQSGRAALRDKLMSQQK DALRKRKQPPTTVLSLLRQSQMDSS AVPKPGPDLLRKQGQGSFPISSMSQ LLQSMSCQSSHLSSNSTPGCGASNT ALPCSANQLHFTDPSMNSSVLQNIP LRGEAVHCHNANTNFVHSNSPVPN HHLAGLINQIQASGNCGMLSQSGM ALGNSLHPNPPQSRISTSSTPVIPNSI VSSYNQTSSEAGGSGPSSSIAIAGTN HPAITKTTSVLQDGVIVTTAAGNPLQ SQLPIGSDFPFVGQEHALHFPSNSTS NNHLPHPLNPSLLSSLPISLPVNQQ HLLNQNLLNILQPSAGEGDMSSINN TLSNHQLTHLQSLLNNNQMFPPNQ QQQQLLQGYQNLQAFQGQSTIPCPA NNNPMACLFQNFQVRMQEDAALL NKRISTQPGLTALPENPNTTLPPFQD TPCELQPRIDPSLGQQVKDGLVVGG PGDASVDAIYKAVVDAASKGMQVVIT TAVNSTTQISPIPALSAMSAFTASIGD PLNLSSAVSAVIHGRNMGGVDHDG RLRNSRGARLPKNLDHGKNVNEGD GFEYFKSASCHTSKKQWDGEQSPR GERNRWKYEEFLDHPGHIHSSPCH ERPNNVSTLPFLPGEQHPILLPPRNC PGDKILEENFRYNNYKRTMMSFKE RLENTVERCAHINGNRPRQSRGFGE LLSTAKQDLVLEEQSPSSSNSLENSL VKDYIHYNGDFNAKSVNGCVPSPSD AKSISSEDDLRNPDSPSSNELIHYRP RTFNVGDLVWGQIKGLTSWPGKLVR EDDVHNSCQQSPEEGKVEPEKLKTL TEGLEAYSRVRKRNRKSGKLNNHLE AAIHEAMSELDKMSGTVHQIPQGDR QMRPPKPKRRKISR	True	False	2.34	5.0	0.882	0.668	0.702	0.743	4.0