	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	extracell region
0	Q8BVE8	NSD2_MOUSE	Mus musculus	Histone-lysine N-methyltransferase NSD2	25.385059	NaN	T110;T114;S121;S172;S376;T422	28528544	MEFSIRKSPLSVQKVVKCMKMKQTP EILGSANGKTQNCEVNHECSVFLSK AQLSNSLQEGVMQKFNGHDALPFL PAEKLKDLTSCVFNGEPGAHDTKLC FEAQEVKGIGTPPNTTPIKNGSPEIK LKITKTYMNGKPLFESSICGDGAADV SQSEENEQKSDNKTRRNRKRSIKYD SLLEQGLVEAALVSKISSPADKKIPVK KESCPNTGRDRDLLLKYNVGDLVWS KVSGYPWWPCMVSADPLLHNHTKL KGQKKSARQYHVQFFGDAPERAWIF EKSLVAFEGEEQFEKLCQESAKQAP TKAEKIKLLKPISGRLRAQWEMGIVQ AEEAASMSIEERKAKFTFLYVGDQL HLNPQVAKEAGIVTEPLGEMVDSSG ASEEAAVDPGSVREEDIPTKRRRRTK RSSSAENQEGDPGTDKSTPPKMAEA EPKRGVGSPAGRKKSTGSAPRSRKG DSAAQFLVFCQKHRDEVVAEHPDAS GEEIEELLGSQWSMLNEKQKARYNT KFSLMISAQSEEDSGNGNGKKRSHT KRADDPAEDVDVEDAPRKRLRADKH SLRKRETITDKTARTSSYKAIEAASSL KSQAATKNLSDACKPLKKRNRASAT ASSALGFNKSSSPSASLTEHEVSDSP GDEPSESPYESADETQTEASVSSKKS ERGMAAKKEYVCQLCEKTGSLLLCE GPCCGAFHLACLGLSRRPEGRFTCT ECASGIHSCFVCKESKMEVKRCVVN QCGKFYHEACVKKYPLTVFESRGFR CPLHSCMSCHASNPSNPRPSKGKM MRCVRCPVAYHGGDACLAAGCSVIA SNSIICTGHFTARKGKRHHTHVNVS WCFVCSKGGSLLCCEACPAAFHPDC LNIEMPDGSWFCNDCRAGKKLHFQ DIIWVKLGNYRWWPAEVCHPKNVPP NIQKMKHEIGEFPVFFFGSKDYYWT HQARVFPYMEGDRGSRYQGVRGIG RVFKNALQEAEARFNEVKLQREARE TQESERKPPPYKHIKVNKPYGKVQIY TADISEIPKCNCKPTDENPCGSDSEC LNRMLMFECHPQVCPAGEYCQNQC FTKRQYPETKIIKTDGKGWGLVAKRD IRKGEFVNEYVGELIDEEECMARIKY AHENDITHFYMLTIDKDRIIDAGPKG NYSRFMNHSCQPNCETLKWTVNGD TRVGLFAVCDIPAGTELTFNYNLDCL GNEKTVCRCGASNCSGFLGDRPKTS ASLSSEEKGKKAKKKTRRRRAKGEG KRQSEDECFRCGDGGQLVLCDRKF CTKAYHLSCLGLGKRPFGKWECPW HHCDVCGKPSTSFCHLCPNSFCKE HQDGTAFRSTQDGQSYCCEHDLRA DSSSSTKTEKPFPESLKSKGKRKKRR CWRRVTDGK	None	None	None	None	None	None	None	None	None