

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extrac region
Q9Z180	SETBP_MOUSE	Mus musculus	SET-binding protein	25.019509	T1098;S1271;S1276;S1277;S1326;S1343	NaN	34418053	MEPREMLSSCRQGESEFLQSSSRSPAPGCSGEPLKGISVGGEMPEEEDELGSGRDVDCNSNADSEKVVAGDGLLEEQEFFSIKEANFTEGSLKLIKIQTTRAKKPPKNLENYICPPEIKITIKQSGDQKVSRTGKNSKATKEDERNHKKKLLTAGDPTASDLKAFQTQAYERPQKHSTLQYDPGHSQGFSDTLKPKHQKSSSQSHMEWSNSSDSGPATQNCFISPEAGRDTASTSKVPALPVA SFAKAQSKKGSTGGAWSQLSSSSKDLLGVSVPSPSHNSPATPSSSAECNGLQPLGDQDGGSTKDLPEPPTLSSK KKSCKDMISQTLPNSDLWVKSAAQKAFETTEGKREAYSADSAQEASPARQSISVSNPENDSSHVRITPIKTPSLDPSNHRKRRQSIKAVVEKIVPEKALASGISMSSEVVNRILSNSEGSKKDPRVPKLGKMIENETPSVLETGGNAEKIVPGGASKQRKPPMVMVMTSPTRTEHAPSGKLEIQHPKFAAKRRCSKAKPPAMLREAVLATAEKLMEVPPSAYPIITPSPLYTNTDSLTVITPVKKRGRPKKQPLLTVEIHEGTSTSPVSPISREFPGTKRKRNRNLAKLAQLVPGEDKPMSEMFKHKVKVGLVLDKKTIKTINKMKTLLKRKNILNQLSCSSVALKAKAPPETSPGAASIESKLGKQINVSKRGTIYGKKRGRKPRTELPPPSEEPKTAIKHPRPVSSQPDVPAVPSFQSPVASSPAA MHPLSTQLGGSNGNLSPASTETNFS ELKTMPNLQPISALPTKTQKGIHGGT WKLSPRMLMANSPSHLCEIGSLKEITLSPVSESHSEETIPSDSGIGTDNNSTSDQAEKSSERRRYSFDFCSLDNPEAIPSDTSTKNRHGHRQKHLIVDTFLAHESLKKPKHKRKRKSLQNRDDLQFLAELEELITKFQVFRISHRGYTFYHENPYPISIFRINFQYYVPYIQYDPLLYLRRTSDLKSKKKRGRPAKNTDMTKVPFLQGFYSYPIPSGSYYAPYGMPTSM PPMNLGYGQYPAPLYLSHTLGAASPFMRPTVPPPPQHASSHVKISGATKHAKHGVHLQGTVMGLGDIQPSL NPPKVGATLSSSRLHKRKHKKRKHKEKEDRILGTHDNLGSLFAGKATGFS SHLLSERLSGSDKELPLVSEKSKHKERQKHQHGGEASHKVSKNNFEVDLSTLSLSDAQHWTOAKDKGDLSSPEV ESCAKRYSGGGDSTRSEGLDVFSEMNPSSDKWSDMGGSKRRSFEGFGTYREKDIQAFKMNRRKERGSYESSMSPGMPSPHLKVDQTAHHSKSEGSISAMMARKKPTAVDSVAIPSAFVLSLLAASAATSDAASSLKKRFRREIEAIOCEVRKMCHYTKLLSTKKNLDHVNKILKAKRLQRQSKTGNFVKRRRGRPRKQPSQFDEDSRDQMPVLEKCIDLPSKRGQKPSLSPLALEPASGQDVMATIEAVIHMAREAPLPPPPPPPLPPPPP PPPPPPLPKTARGGKRKRHPQPPA QPAQPTPQPLPQEEVKAKRPRKSRASESDVLP	True	False	2.983	4.738	1.304	0.648	0.515	1.23	1.311