

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q5SPL2	PHF12_MOUSE	Mus musculus	PHD finger protein 12	16.335304	T472;S616;S622;S630	S131;S134;S555;T557;T570;T670	34887587;36852467	MWEKMETKTIVYDLDTSGGLMEQI QALLAPPKTDEAEKRSRKPEKESRR SGRATNHDCSDSCKEGGDLLCCDH CPAAFHLQCCNPPLSEEMLPPEGEW MCHRCTVRRKKREQKELGHVNGL VDKSSKRRTTSPSSDSDLDRPASKTE LKAIAHARILERRASRPGTPTSNAST ETPTSEHNDVDEDIVDVDEEPPVAEE PDYVQPQLRRPFELLIAAAMERNPT QFQLPNELTCTTALPGSSKRRRKEE TTGKNVKRTOHELDHNGLVPLPVKV CFTCNRSCRVAPLIQCDYCPPLLFHM DCLEPPLTAMPLGRWMCNPHIEHV VLNQKNLTLNRCQVDFDRFQDTISQ HVVKVDFLNRIHKKHPPNRRVLQSV KRRSLKVPDAIKSQYQFPPLIAPAAI RDGELICSGVPEESQTHLLNSEHLA TQAEQQEWLCSVALQCSILKHLA KQMPSPWDSEQTEKADIKPVIVTDS SITTSLQADKAPLPSHYPLSCPSAVS TQNSLGCSPPHQPPTLEDISCSSCVE KSKKAPCGTANGPVNTEIKANGPHL YSSPTDSTDPRLPGANTPLPGLTHR QGWPRPLTPPSAGGLQNHVGIIVKT ENATGPSSCPQRSVLPVPSLPPSIPS SCASIENTSTLHRKTVQSQIGPSSTE SRPLGSPPNATRVLTPPQAAGDSILA TGANQRFCSPAPSSDGKVSPGTLSIG SALTVPSPFANSTAMVDLTNSLRAF MDVNGEIEINMLDEKLIKFLALQRV HQLFPSRVQASPGNVGTHPLASGGH HPEVQRKEVQARAVFCPLLGLGGAV NMCYRTLTYGTGADMDVCLTNYGH CNYVSGKHACIFYDENTKHYELLNY SEHGTTVDNVLYSCDFSEKTPPTPPS SIVAKVQSVIRRRRHQKQDEEPPSEEA AMMSSQAQGPQRRPCNCKASSSLI GGSGAGWEGTALLHHGSYIKLGCLQ FVFSITEFATKQPKGDASLLQDGVLA EKLSLKP HQGPVLRNSVP