

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	go ap
Q5SPL2	PHF12_MOUSE	Mus musculus	PHD finger protein 12	38.185612	T472;S616;S622;S626;S630	S131;S134;S555;T557;T570;T670	34887587;40885482;36852467	MWEKMETKTIVYDLDTSGGLMEQI QALLAPPKTDEAEKRSRKEKESRR SGRATNHDSCDSCKEGGDLLCCDH CPAAFHLQCCNPPLEEMLPPGEW MCHRCTVRRKKREKQKELGHVNGL VDKSSKRRTTSPSSDTDLLDRPASKTE LKAIAHARILERRSRPGTPTSNAST ETPTSEHNDVDEDIVDVDEEPVAAE PDYVQPQLRRPFELIAAAMERNPT QFQLPNELTCTTALPGSSKRRRKEE TTGKNVKRTQHELDHNGLVPLPVKV CFTCNRSRVAPLIQDCDYCPLLFHM DCLEPPLTAMPLGRWMCNPHIEHV VLNQNLTLSNRQCVFDRFQDTISQ HVVKVDFLNRHKHPPNRRVLSV KRRSLKVPDAIKSQYQFPPLIAPAAI RDGELICSGVPEESQTHLLNSEHLA TQAEQQEWLCSVVALQCSILKHLA KQMPSPWDSEQTEKADIKPVIVTDS SITSLQTADKAPLPSHYPLSCPSAVS TQNSLGCSPPHQPPTLEDISCSVCVE KSKKAPCGTANGPVNTEIKANGPHL YSSPTDSTDPRRLPGANTPLPGLTHR QGWRPLTPPSAGGLQNHVGIIVKT ENATGPSSCPQRSLVPVPSLPPSIPS SCASIENTSLHRKTQVQSIGPSSSTE SRPLGSPPNATRVLTTPQAAGDSILA TGANQRFCSPAPSSDGKVPGLSIG SALTVPSPFANSTAMVDLTNSLRAF MDVNGEIEINMLDEKLIKFLALQRV HQLFPSRVQASPGNVGTHPLASGGH HPEVQRKEVQARAVFCPLGLGGAV NMCYRTLYIGTGADMVCLTNYGH CNYVSGKHACIFYDENTKHYELLY SEHGTTVDNVLYSCDFSEKTPPTPPS SIVAKVQSVIRRRRHQKQDEEPSEEA AMMSSQAQGPQRPCNCKASSSLI GGSGAGWEGTALLHHGSYIKLGCLO FVFSITEFATKQPKGDASLLQDGVLA EKLSLKP HQGPVLRNSVNP	True	False	2.389	4.71	1.345	1.128	Fal