

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	extracellular region
000291	HIP1_HUMAN	Homo sapiens	Huntingtin-interacting protein 1	14.128268	T50	S338	30379171;37217939;28510447	MDRMASMKQVNPPLPKVLSRRGV GAGLEAAERESFERTQTVSINKAINT QEVAVKEKHARTCILGTHHEKGAQT FWSVVNRLPLSSNAVLCWKFCHVF HKLLRDGHPNVLKDSLRYRNELESD MSRMWGHLSSEGYGQLCSIYLLKLR KMEYHTKNPRFPGNLQMSDRQLDE AGESDVNFFQITVEMFDYLECEL NLFQTVFNSLDMRSRSVSTAAGQCR LAPLIQVILDCSHLYDYTVKLLFKLH SCLPADTLQGHDRDFMEQFTKLKDL FYRSSNLQYFKRLIQIPQLPENPPNF LRASALSEHISPVVVIPAEASSPDSEP VLEKDDLMDMDASQQLFDNKF DIFGSSFSDDPFNFNSQGVNKDEK DHLIERLYREISGLKAQLENMKTES QRVVLQKLGHVSELEADLAEQQHLR QQAADDCEFLRAELDELRRQREDTE KAQRSLSEIERKAQANEQRYSKLKE KyselVQNHADLLRKNAEVTQVVS MARQAQVDLEKKELEDLSLERISD QGQRKTOEQLEVLVLESLKQELATSQR ELQVLQGSLETSQAQSEANWAAEFAE LEKERDSLVSAAHREELLSALRKE LQDTQLKLASTEESMCQLAKDQRK MLLVGSRKAAEQVIQDALNQLEPP LISCAGSADHLLSTVTSISSCIEQLEK SWSOYLACPEDISGLLHSITLLAHLT SDAIAHGATTCLRAPPEPADSLTEAC KQYGRETLAYLASLEEEGSLENADS TAMRNCLSKIKAIGEELPRGLDIQ EELGDLVDKEMAATSAAIETATARIE EMLSKSRAGDTGVKLEVNERILGCC TSLMQAIQVLIVASKDLQREIVESGR GTASPKFYAKNSRWTEGLISASKAV GWGATVMVDAADLVVQGRGKFEEL MVCHEIAASTAQLVAASKVKADKD SPNLAQLQASRGVNOATAGVVAST ISGKSQIEETDNMDFSSMTLTQIKR QEMDSQVRVLELENELQKERQKLG ELRKKHYELAGVAEGWEEGTEASPP TLQEVVTEKE	True	True	4.581	5.0	0.941	0.281	5.0	2.754	0.924