	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	extracellula region
0	Q9HCH0	NCK5L_HUMAN	Homo sapiens	Nck-associated protein 5-like	23.686208	NaN	S440;S451;S470;S477;S493;S496;S498;S571;S577;T659;S767;S1194	35254053	MSEAMDQPAGGPGNPRPGEGDDGS MEPGTCQELLHRLRELEAENSALAQ ANENQRETYERCLDEVANHVVQAL LNQKDLREECIKLKKRVFDLERQNQ MLSALFQQKLQLTTGSLPQIPLTPLQ PPSEPPASPSLSSTEGPAAPLPLGHC AGQREVCWEQQLRPGGPGPPAAPPP ALDALSPFLRKKAQILEVLRALEETD PLLLCSPATPWRPPGQGPGSPEPING ELCGPPQPEPSPWAPCLLLGPGNLG GLLHWERLLGGLGGEEDTGRPWGP SRGPPQAQGTSSGPNCAPGSSSSSS SDEAGDPNEAPSPDTLLGALARRQL NLGQLLEDTESYLQAFLAGAAGPLN GDHPGPGQSSSPDQAPPQLSKSKGL PKSAWGGGTPEAHRPGFGATSEGQ GPLPFLSMFMGAGDAPLGSRPGHP HSSSQVKSKLQIGPPSPGEAQGPLLP SPARGLKFLKLPPTSEKSPSPGGPQL SPQLPRNSRIPCRNSGSDGSPSPLLA RRGLGGGELSPEGAQGLPTSPSPCY TTPDSTQLRPPQSALSTTLSPGPVVS PCYENILDLSRSTFRGPSPEPPPSPL QVPTYPQLTLEVPQAPEVLRSPGVPP SPCLPESYPYGSPQEKSLDKAGSESP HPGRRTPGNSSKKPSQGSGRRPGDP GSTPLRDRLAALGKLKTGPEGALGS EKNGVPARPGTEKTRGPGKSGESAG DMVPSIHRPLEQLEAKGGIRGAVAL GTNSLKQQEPGLMGDPGARVYSSH SMGARVDLEPVSPRSCLTKVELAKS RLAGALCPQVPRTPAKVPTSAPSLGK PNKSPHSSPTKLPSKSPTKVVPRPGA PLVTKESPKPDKGKGPPWADCGSTT AQSTPLVPGPTDPSQGPEGLAPHSAI EEKVMKGIEENVLRLQGQERAPGAE VKHRNTSSIASWFGLKKSKLPALNR RTEATKNKEGAGGGSPLRREVKME ARKLEAESLNISKLMAKAEDLRRAL EEEKAYLSSRARPRPGGPAPGPNTG LGQVQGQLAGMYQGADTFMQQLLN RVDGKELPSKSWREPKPEYGDFQPV SSDPKSPWPACGPRNGLVGPLQGCG KPPGKPSSEPGRREEMPSEDSLAEP VPTSHFTACGSLTRTLDSGIGTFPPP DHGSSGTPSKNLPKTKPPRLDPPPG VPPARPPPLTKVPRRAHTLEREVPGI EELLVSGRHPSMPAFPALLPAAPGH RGHETCPDDPCEDPGPTPPVQLAKN WTFPNTRAAGSSSDPLMCPPRQLEG LPRTPMALPVDRKRSQEPSRPSPTP QGPPFGGSRTPSTSDMAEEGRVASG GPPGLETSESLSDSLYDSLSSCGSQG	None	None	None	None	None	None	None	None	None