

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion
Q9UPT8	ZC3H4_HUMAN	Homo sapiens	Zinc finger CCCH domain-containing protein 4	27.098827	S835;T898;S899;S904;S907;S908	T72;T75;S76;S92;S94;T155;S807;S808;S904;S907;S908;S1104;T1106;S1108;S1110;S1114;T1118;S1269;S1275	31492838;35132862;34019948;35289036;29237092;37217939;29351928	MEAAPGTPPPPSEPPPPSPPPPST PSPPPCSPDARPATPHLLHRLPLPD DREDGELEEGELEDGAEETQDTS GPERSRKEKGEKHSDSDEEKSHR RLKRRKKEKEREKRRSKRRKSKH KRHASSDDFSDSDDSDSDFSPSEK HRKYREYSPYPASHQQYPPSHATPL PKKAYKMDKSYGMYEDYENEQY GEYEGDEEEDMGKEDYDDFTKELN QYRRAKEGSSRGRGRGRGRGYRG RGSRRGSRGRGMGRGRGRGRGS MGGDHPEDEEDFYEEEMDYGESEE PMGDDDYDEYSKELNQYRRSKDSR GRGLSRGRGRGRGRGKGMGRGRG RGSRRGMNKGGMNDDDFYDED MGDGGGGSYRSRDHDKPHQSDKK GKVICKYFVEGRCTWGDHCNFSHDI ELPKKRELCKFYITGFCARAENCPY MHGDFPCKLYHTGNCINGDDCMF SHDPLTEETRELLDKMLADDAEAGA EDEKEVEELKKQGINPLKPPPGVGL LPTPPRPPGQAPTSNCRPMQGGP PPPPPPPPPPGPPQMPMPVHEPLS PQQLQQQDMYNNKIPSLFEIVVRPT GQLAEKLGVRFPGGPPGPMGPGP NMGPPGPMGGPMHPDMHPDMHP DMHPDMHADMHADMPMGPMNP GPPMGPPGPPMMPYGPDSPHSG MMPPIPPAQNfyENFYQQEGMEM EPGLLGAEDYGHYEELPGEPEHL FPEHLEPDSFSEGGPPGRPKPGAG VPDFLPSAQRALYLRIQQKQEEEE RARRLAESSKDRENEEGDTGNWY SSDEDEGGSSVTSILKTLRQOTSSRP PASVGELESSGLDPRQLQKGHPTGS RLADPRLSRDPRLTRHVEASGGSGP GDSGSPDPRALARALPTSKPEGSLHS SPVGPSSSKGSGPPPTTEEEGERALR EKAVNIPLDPLPGHPLRDRSRLQ FSHIKDVTLKPSFARTVLWNPED LIPLPIPKQDAVPPVPAALQSMPTLD PRLHRAATAGPPNARORPGASTDSS TQGANLPDFELLSRILKTVNATGSSA APGSSDKPSDPRVRKAPTDPRLQKP TDSTASSRAAKGPAEAPSPTASPSG DASPPATAPYDPRVLAAGGLGQGGG GGQSSVLSGISLYDPRTPNAGGKATE PAADTGAQPKGAEGNGKSSASKAKE PPFVRKSALEQPETGKAGADGTTPT DRYNSYNRPRKAAAAPAAATATPPP EGAPPQPGVHNLFPVPTLFGTVKQTP KTGSGSPFAGNSPAREGEQDAASLK DVFKGFDPASPFCCQ	None	None	None	None	None

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