

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
015265	ATX7_HUMAN	Homo sapiens	Ataxin-7	24.512865	S443	NaN	37217939;29351928	MSERAADDVVRGEPRRAAAAAGGAA AAAAARQQQQQQQQPPPPQPPRQ QHPPPPRRTRPEDGGPGAASSTAA AMATVGERRRPLSPPEVMLGQSWNL WVEASKLPGKDGTELDESFKFQGN REVMGLCREDMPIFGFPAHDDDFYL VVCNDCNQVVKPAFQSHYERRHS SSSKPPLAVPPTSVFSPSLSKSKG GSASGNSRSGGVLASSSSSKLLK SPKEKLQLRGNTRPMHQSSRVPH GRIMTPSVKVEKIHKMDGTLKSA VGPTCPATVSSLVKPGLNCPSIPKPT LPSPGQILNGKGLPAPPTLEKKPEDN SNNRKFNLKRLSEREFDPDIHCGVI DLDTKKPCTRLTCKTHSLTORRAV QGRRKRFVLLAEHKNKTRKELIR HPDSQPPQPLRDPHPAPRPTSQEP HQNPHGVIPSESKPFVASKPKPHTP SLPRPPGCPAQGGGAPIDPPPVHES PHPLPATEPASRLSSEEGEDDKE ESVEKLDCHYSGHHPQASFCFTFGS RQIGRGGYVFDNRWNRRLCALNLM VEKHLNAQLWKKIPVPTTSPISTR PHRTNSVPTSQCGVSYLAAATVSTSP VLLSSTCISPNKSVPAHGTTLNAQP AASGAMPVCSMQSRQVSSSSSSPS TPSGLSSVPSSPMSRKPQKLKSSKSL RPKESGNSNQCNAASSTSGGSGK KRKNSSPLLVHSSSSSSSSSSHS MESFRKNCVAHSGPPYPTVTSSH IGLNCVTNKANAVNVRHDQSGRGP PTGSPAESIKRMSVMVNSSDSTLSL GPFHQSNELPVNSHGSFSSHHTPL DKLIGKRRKCPSSSSINNSSSKPTK VAKVPAVNNVHMKHTGTIPGAQGL MNSSLLHQPKARP	True	False	4.305	5.0	1.95	1.975	1.061	1.616	1.459