

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
Q8R4I1	ATX7_MOUSE	Mus musculus	Ataxin-7	24.992371	S697	NaN	36852467	MSERAADDVVRGEPRAAGGAAAAAR QQQQQPQLQQRQHPPLRPRAE DGGTGDITTSAAAMATVGERRPLPS PEAMLGQSWNLWVEASKLPGKDG ELDESFKEFGKNREVMGLCREDMPI FGLCPAHDDFYLVVNCDCNQVVKP QAFQSHYERRHSSSSKPALAVPHTS VFSLLPSLSKSKGSGAGGSSRPPSGG VLCASSSSKLLRLPKEKPLRGNMK PMHPVQQIKVPHGRVMTPSVKVEK MHPKMDGTLKSTVGPACPATMSS AVKPGLNCPSPKPTLPSGQILNGK GLPAMPTEKKSEDSSNNRKFNLKR LSEREFDPDIHCGVIDLDTKKPCTRS LTCKTHSLTORRAVOGRRKRFDVLL AEHKNKAREKELIRHDSQVPHPLR DPHPTPPRTPQEPQLPAESKPFLASK PKPQTPSLPRPPGCPAQGGSTPIDP PPGQESPHPLPATEPASRLSSEEGE GDDREESVEKLDCHYSGRHPQPASF CTFGSRQIGRGGYVFDNRWNRLRCA LNLNVEKHLNAQLWKKIPPVPCTTS PVSARVPHRTNSVPTSQGGISYLAAT TVSAPPVLLSSTCISPNSKSVPAHGT TLNAQPAGSGAMPVCSVQSRQVSA SSSPSTPSGLSSVSSPLSRKPQKW KPSKSIRPKESALSTNCHNASSSTS GGSGKKRKNSSPLLVPSSSSSSSSS SSSHSVNSFRKNCVAHSGTPYLSTA PSSHIGLNCVTNKTHSVSLRHEQA GRGPAGVSSAEPKRMVSVNNSDS TSLGPFHQASELVNPVNSHTPLDK LIGKKRKCSPGSSVGNVSGSKPTKVA KLPAMNNVHMKHTGNISGAQGLTN NSLLHQPKARP	True	False	3.127	5.0	1.832	1.742	1.006	1.515	1.319