

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q99700	ATX2_HUMAN	Homo sapiens	Ataxin-2	29.465981	S213;S216;S218;S220;S528;T529;S533;S541;T666;S776	S248;S393;S466;S478;S508;S554;S624;S642;S684;S728;T741;S772;S784;S856;S857;S861;S865;S867;S888;S889	31492838;32119511;29351928;34725712;34019948;21740066;29237092;30620550;33214551;35254053;30397120;35132862;35138101;30059200;28604694;20305658;30379171;28657654	MRSAAAAPRSPAVATESRRFAAARW PGWRSLQRPARRSGRGGGGAAPGP YPSAAPPPPGPGPPSRQSSPPSASD CFGSNGNGGGAFRPGSRRLGLGG PPRPFVLLPLASPGAPPAAPTRAS PLGARASPPRSGVSLARPAPGCP CEPVYGPLTMSLKPQQQQQQQQQ QQQQQQQQQQQQPPAAANVRKP GSGLLASPAAPSPSSSVSSSSAT APSSVVAATSGGGRPGLGRGRNSK GLPOSTISFDGIYANMRMVHILTSV GSKCEVQVKNGGIYEGVFKTYSKPC DLVLDAAHEKSTESSGPKREEIME SILFKCSDFFVVVQFKDMSSYAKRD AFTDSAISAKVNGEHKEKDLEPWDA GELTANEELEALENDVSNWDPND MFRYNEENYGVVSTYDSSLSSYTV LERDNSEEFKREARANQLAEIEE SAQYKARVALENDDRSEEEKYTA RNSSEREGHSINTRENKYIPPGQRN REVISWGSQRQNSPRMGQPGSGSM PSRSTSHTSDFNPNSSGDQRVVNGG VPWPSPCPSPSRPPSRYSQGPNSLP PRAATPTRPPSRPPSRPPSHPSA HGSPAPVSTMPKRMSSEGPPRMSP KAQRHPRNHRVSAGRGSISSGLEFV SHNPPSEAATPPVARTSPSGGTWSS VVSGVPRLSPKTHRPRSPRQNSIGN TPSGPVLASFPQAGIIPTEAVAMPIAA SPTPASASNRAVTPSSEAKDSRLQD QRQNSPAGNKENIKPNETSPSFSKA ENKGISPVVSEHRKQIDDLKKFKND FRLQPSSTSEMDQLLNKNREGEKS RDLIKDKIEPSAKDSFIENSSNCTS GSSKPNSPSISPSILSNTEHKRGPEV TSQGVQTSSPACKQEKDDKEEKDA AEQVRKSTLNPNAKEFNPRSFQPK PSTTPTSPRQAQPSMVGHQPT PVYTPVCFAPNMMYPVPVSPGVQP LYPIPMTPMPVNQAKTYRAVPNMPQ QRQDQHHQSAMMHPASAAGPPIAA TPPAYSTQYVAYSPQQFPNQLVQH VPHYSQHPHVYSPVIQGNARMMA PPTHAPGLVSSSATQYGAHEQTHA MYACPKLPYNKETSFSFYFAISTGSL AQQYAHPNATLHPHTPHPOPSATPT GQQSQHGGSHAPSPVQHHQHQA AQALHLASPPQQSAIYHAGLAPTPPS MTPASNTQSPQNSFPAAQQTVFTIH PSHVQPAYTNPPHMAHVPAHVQS GMVPSHPTAHAPMMLMTTQPPGGP QAALAQSALQPIPVSTTAHFPMTHP SVQAHHQQL

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