

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q8WWM7	ATX2L_HUMAN	Homo sapiens	Ataxin-2-like protein	41.877697	T269;S369;S461;T473;S500;S563;S678;T681;T683;S684	S103;S111;T118;S238;T264;S306;T309;S335;S339;T349;S391;S409;S449;S493;S496;S557;S558;S559;S563;S594;T632;S634;S674;S680;S684	29237092;28657654;31637018;22661428;34725712;30379171;21740066;32574038;31492838;23301498;30397120;32870666;30620550;29351928;34019948;33214551;27114449;20305658;32119511;34229054;30059200;31373491	MLKPQPLQQPSQPQQPPPTQAVAR RPPGGTSPPNGGLPGPLATSAPP PAAASPCCLGPVAAAGSGLRRGAEGIL APQPPPPQHQHQPGRPGAAAIGSARGQ STGKGPPQSPVFEVYNNRMLHFL TAVVVGSTCDVKVKNGTTYEGIFKTL SKFELAVDAVHRKASEPAGGPRRED IVDTMVFKPSDVMLVHFRNVDFNYA TKDKFTDSAIAMNSKVNGEHKEKVL QRWEGGDSNSDDYDLESDMSNGW DPNEMFKFNEENYGVKTTYDSSLSS YTVPLEKDNSEEFQRELRAAQLAR EIESSPQYRLRIAMENDDGRTEEEK HSAVQRQCGSGRESPLASREGKYIPL PQRVREGPRGGVRCSSSRGGRPGLS SLPPRGPHHLNSSPGPGSEARGIN GGPSRMSFKAQRPLRGAKTLSSPSN RPSGETSVPPPPAVGRMYPPRSPKSA APAPISASCPEPPIGSAVPTSASIPVT SSVSDPGVGSISPASPKISLAPTDVKE LSTKEPGRITLPEQELARIAGKVPGLQ NEQKRFQLEELRKFGAQFKLQPS PENSLDFPPRILKEEPKGEKEVDG LLTSEPMGSPVSSKTESVSDKEDKP PLAPSGGTEGPEQPPPCPSQTGSPP VGLIKGEDKDEGPVAEQVKKSTLNP NAKEFNPTKPLLSVKNKSTSTPTSPGP RTHSTPSIPVLTAGQSGLYSPQYISYI PQIHMGPAVQAPQMYYPVNSVPG QQGKYRGAKGSLPPQRSQDQHPASA PPMMQAAAAAGPPLVAATPYSSYIPY NPQQFPGQFAMMQMAHYPSQPVF APMLQSNPRMLTSGSHPQAISSST PQYPSAEQPTPQALYATVHQSYPHH ATQLHAHQPPATTPTGSQPQSQAHA APSPVQHQAGQAPHLGSGQPQNL YHPGALTGTTPSLPPGSAQSPQSSF PQPAAVYAIHHQQLPHGFTNMAHVT QAHVQTGITAAPPHPGAPHPQVM LLHPPQSHGGPPQGAVPQSGVPALS ASTPSYPYIGHPPQGEQPGQAPGFPG GADDRIREFSLAGGIWHGRAEGLQV GQDARVLGGE

0