

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion
Q8WWM7	ATX2L_HUMAN	Homo sapiens	Ataxin-2-like protein	35.312342	S45;T268;T269;S369;S461;T473;S500;S563;S674;S678;T679;S680;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	28657654;31492838;32870666;23301498;34019948;34725712;37217939;20305658;32574038;21740066;35254053;29237092;29351928;31373491;35138101;30397120;30620550;30059200;34229054;30379171;22661428;31637018;35132862;33214551;32119511;27114449;38665916	MLKPOPLQPPSQPQQPPPTQQAVAR RPPGGTSPPNGGLPGPLATSAAPPGP PAAASPCLPVAAAGSGLRRAEGIL APQPPPPQQHQRPGAAAIGSARGQ STGKGGPPQSPVFEVYNNRMLHFL TAVVGSTCDVKVKNGTTYEGIFKTL SKFELAVDAVHRKASEPAGGPRRED IVDTMVFKPSDVMLVHFRNVDFNYA TKDKFTDSAIAMNSKVNGEHKEKVL QRWEGGDSNSDDYDLESDMSNGW DPNEMFKFNEENYGVKTTYDSSLSS YTVPLEKDNSEEFQRELRAAQLAR EIESSFOYRLRIAMENDDGRTEEEK HSAVQRQSGRESPLASREGKYIPL PQVRREGPRGGVRCSSSRGGRPGLS SLPPRGPHHLDNSSPGPGEARGIN GGPSRMSPKAQRPLRGAKTLSSPSN RPSGETSVPPPPAVGRMYPPRSPKSA APAPISASCPPEPPIGSAVPTSSASIPVT SSVSDPGVGSISPASPKISLAPTDVKE LSTKEPGRITLPEQELARIAGKVPGLQ NEQKRFLQLEELRKFQAQFKLQPSSS PENSLDPFPPRILKEPKGKEKEVDG LLTSEPMGSPVSSKTESVSDKEDKP PLAPSGGTEGPEQPPPPCPSQTGSP VGLIKGEDKDEGPVAEQVKKSTLNP NAKEFNPTKPLLSVNKSTSTPTSPGP RTHSTPSIPVLTAGQSGLYSPQYISYI POIHMGPAPVAPQMYPPVNSVPG QQGKYRGAKGSLPPQRSQDQHPASA PPMMQAAAAAGPPLVAATPYSSYIPY NPQQFPGQPAMMQPMAHYPSQPVF APMLQSNPRMLTSGSHQAISSST PQYPSAEQPTPQALYATVHQSYPHH ATQLHAHQPPATPTTGSQPQSQHA APSPVQHQAQAPHLGSGQPQQNL YHPGALTGTPLPPLPGPSAQSPQSSF PQPAAVYAIHHQQLPHGFTNMAHVT QAHVQTGITAAPPPHGPAPHPQVM LLHPPQSHGGPPQAVPQSGVPALS ASTPSYPYIGHPPQGEQPGQAPGFPG GADDRIREFSLAGGIWHGRAEGLQV GQDARVLGGE	None	None	None	None	None