

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q7TQH0	ATX2L_MOUSE	Mus musculus	Ataxin-2-like protein	31.853967	T436	S27;T45;S109;T116;S236;T262;S304;T307;S333;S337;T347;S389;S407;S453;S496;S499;S560;S561;S562;S597;T635;S637;S677;S683;S687	22645316;25153642;24788674;28528544;22517741;28135057;22826440	MLKPQQPQTSQPQQPPPTQQAVAR RSPGGTSPPNGGLPGPLTATAAPP GPPAAVSPCLGPAAAAAGSGLRRAESIL AASAPPQHQRPGAVAIGSVRGQTT GKGPPQSPVFEQVYNNRMLHFLTA VVGSTCDVKVKNGTTYEGIFKTLSSK FELAVDAVHRKASEPAGGPRREDIV DTMVFKPSDVLVHFRNVDFNYATK DKFTDSAIAMNSKVNGEHKEKVLQR WEGGDSNSDDYDLESDMSNGWDP NEMFKFNEENYGVKTTYDSSLSSYT VPLEKDNSEEFRQRELRAAQLAREI ESSPQYRLRIAMENDDGRTEEEKHS AVQRQSGRESPLVSREGKYIPLPQ RVREGPRGGVRCSSSRGGRPGLSSL PPRGPHELLDNSSPGPGSEARGINGG PSRMSPKAQRPLRGAKTLSSPSNRP SGEASVPPTSAALPFLPVGRMYPPRS PKSAAPAPVSASCPEPPIGSAVASSA SIPVTSSVDPGAGSISPASPKLSLTP TDVKELPTKEPSRNLEAQELARIAGK VPGQLONEQKRFQLEELRKFQGAQFKL QPSSSPETGLDPFPSRILKEEAKGKE KEVDGLLTSDPMGSPVSSKTESILDK EDKVP MAGVGGTEGPEQLPAPCPSQ TGSPVGLIKGDEKEEGPVTEQVKKS TLNPNAKEFNPTKPLLSVNKSTSTPT SPGPRTHSTPSIPVLTAGQSGLYSPQ YISYIPQIHMGPVAVQAPQMYPPVSN SVPGQQGKYRGAKGSLPPQRSDQH QPASAPPMMQAAAAAAGPPLVAATP YSSYIPYNPQQFPGQPAMMQPMAH YPSQPVFAPMLQSNPRMLTSGSHPQ AIVSSSTPQYPAAEQPTPQALYATVH QSYPHHATQLHGHQPQPATTPTGSQ PQSQAAPSPVQHQAQAPHLGSG QPQQNLYHPGALTGTTPSLPPGPSA QSPQSSFPQAAVYAIHPHQQLPHG FTNMAHVTQAHVQTGVTAAPPPHP GAPHPQVMLLHPPQGHGGPPQGA VPPSGVPALSASTPSYPYIGHPVQVQ SHPSQQLPFHPPGN