

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum
Q7TQH0	ATX2L_MOUSE	Mus musculus	Ataxin-2-like protein	48.823845	T436;S437;T672;S677;S683	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	22826440;36852467;22645316;24788674;25153642;22517741;28135057;28528544;35822049	MLKPPQPPQTSQPQQPPPTQQAVAR RSPGGTSPPNGGLPGPLTATAAPP PAAVSPCLGPAAGSGLRRAAESIL AASAPPQHQRPGAVAIGSVRGQTT GKGPPQSPVFEVYNNSRMLHFLTA VVGSTCDVKVKNGTTYEGIFKTLSSK FELAVDAVHRKASEPAGGPRREDIV DTMVFKPSDVLVHFRNVDFNYATK DKFTDSAIAMNSKVNGEHKEKVLQR WEGGDSNSDDYDLESDMSNGWDP NEMFKFNEENYGVKTTYDSSLSSYT VPLEKDNSEEFQRELRAAQLAREI ESSPOYRLRIAMENDDGRTEEEKHS AVORQSGRESPLSVSREGKIPLPQ RVREGPRGVRCSSSRGGRPGLSSL PPRGPHHLDNSSPGPGSEARGINGG PSRMSPKAQRPLRGAKTLLSSPNRP SGEASVPP TSAALPFLPVGRMYPPRS PKSAAPAPVSAASCEPPIGSAVASSA SIPVTSSVVDPGAGSISPAKLSLTP TDVKEKLPKPSRNLEAQELARIAGK VPGLONEQKRFQLEELRKFQAQFKL QPSSSPETGLDPFPSRILKEEAKGKE KEVDGLLTSDFMGSPVSSKTESILDK EDKVP MAGVGGTEGPEQLPAPCPSQ TGSPPVGLIKGDEKEEGPVTEQVKKS TLNPNAKEFNPTKPLLSVKNSTSTPT SPGPRTHSTPSIPVLTAGQSGLYSPQ YISYIPIQHMGPAVQAPQMYPPVSN SVPGQGGKYRGAKGSLPPQRSDQH QPASAPPMMQAAAAAGPPLVAATP YSSYIPYNPQQFFPGQAMMQPMAH YPSQPVFAPMLQSNPRMLTSGSHPQ AIVSSSTPQYPAEQPTPQALYATVH QSYPHHATQLHGHQPQATTPTGSQ POSQHAAPSPVQHAGQAPHLGSG QPQONLYHPGALGTGTPPSLPPGPSA QSPQSSFPQAAVYAIHPHQQLPHG FTNMAHVTQAHVQTGVTAAAPPHP GAPHPQVMLLHPQGHGGPPQGA VPPSGVPALSASTPSYPYIGHPVQVQ SHPSQQLPFHPPGN	True	False	3.343	4.608	0.965	0.92