

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q8SWR8	ATX2_DROME	Drosophila melanogaster	Ataxin-2 homolog	27.156312	NaN	S208;S211;S221;S266	28604694	MNNSKRKRTRPTGGGASGGISRYN SNDNSLRPTNNKAGAGGGNGGAAV RPSAQGVYNNFTFMHSATALVGSVV EVLRLSGNIYEGVFRITFSGNFDIALE LPACIKSKNLP EEGKVPKHIIIFPADTV VTIVAKDFDSQYATAGAFQTDGAISD KCNNGARPDEKELEPWDSGANGDIDI ELDSAANGWDPNEMFRKNENTFGV TSTFDDSLASYTVPLDKGDSLEFKEA EAKAEKLA AEIENNPTCRDRLDLEN GDEEALFAAVERPSTEQDQRGDRGD RERNDRDREEREERDRDRDRDRGNK PRGAGDFQLRETMSSDRYITKQTRSI TGPQLSHVGMSSQSGRDRDTRGD GSMMSQSGGGSGGGSTOSTAAL MLAGGLKGVGPAPSANASADSSSKY SGGSMVKRKTVPQGGKVMRNNVPT GGSNVSVSQGGNGNSVQGNKGGYQ PSMGMPQSQSYQGNQIMHGSSQY RNQSHMGGANKLNGDSNANTNKP LPQRQMRQYQGSQSNSSLNYGGEP QSLGKPVHGHGGHHPGQNSNSPPL QTAGPQQQQQQQQQQQQQQQQQ PPQQQHQNIQPGQNTQPARQVR TRDNQMQLRQFGQDFQLAPSNTS PPQQQQQQQQQQHQVQQQQQR ALQQSASPPQQQQQQQQQQHVVL HQVPQTHLHQAAALSQPHYVPPQQP QQAPLPQQQHVPVPHMQKAQQQQ LVETQHQHVQKQHQSQPQVQPPPP QLLQDPSQQLPIYHTMPPPTSPVV VTSPVLEQPPPQMPVVQQQTQQ LATPKPEVSPAPPSSNTTPTGIASP TAGVIASAGSEKTTAAFTPTSNSAT VPTGTAATAGGATGTPVVKKHLN PSAKPFTPRGPSTPNPSRPHTPQTPV PMTNIYTTTGGHVPPAANQPIYVMQ POHPFPQTHPQAGQPPRLRRSNYP PMAASQMHSVASAATGQPLITAGPIP QFIQYGHAPHQQQFQSHTYAPMQM RVYDPQPQLQFMTQTPQSTTSPSG QPHQQFHPPPQSPAGGGPQPAFTP PTQAATYQLMCVHPQSLLANHYFPP PTPQHPQQNQYQIVMQQHQPQ	None	None	None	None	None	None	None		

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