

Product datasheet for **SC118310**

ATX2 (ATXN2) (NM_002973) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	ATX2 (ATXN2) (NM_002973) Human Untagged Clone
Tag:	Tag Free
Symbol:	ATX2
Synonyms:	ATX2; SCA2; TNRC13
Mammalian Cell Selection:	None
Vector:	<u>pCMV6-XL4</u>
E. coli Selection:	Ampicillin (100 ug/mL)
Fully Sequenced ORF:	>NCBI ORF sequence for NM_002973, the custom clone sequence may differ by one or more nucleotides

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ATGCGCTCAGCGCCGCGAGTCTCCGGAGTCCCGCGGTGGCCACCGAGTCTCGCCGCTTCGCCGAGCCA  
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TCCGGCCGCGCAACCCGCGCTCCCGCTCGGCGCCCGCGGTCCCGCGCGTTCGGCGTCTCCTTG  
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GAATTCTGGTTCAGACCAAAGAGTAGTTAATGGAGGTGTTCCCTGGCCATCGCCTTGCCCATCTCCTTCC
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5' Read Nucleotide Sequence: >OriGene 5' read for NM_002973 unedited
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 ACCGAGTCTCGCCGCTTCGCCGCAGCCAGGTGGCCCGGTGGCGCTCGCTCCAGCGGCCG
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3' Read Nucleotide Sequence: >OriGene 3' read for NM_002973 unedited
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 TTCAAGGGTTATTAATAAATAAATAACTCCAGTTTCGGCAAGCAGAGCTGGGGTACCTG
 CGGGACTCTGAAACAGCATATGGAATTATGGAATAGCCCCAAGTTCCTAAATGCCTCTA
 CTCGGTCCAAGTATCTTCACTGCAAGTGAAGTGTAGCATTCTATTGGATGTTACAAG
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 TAAATGGCTGACTGCTGTGGACTGGCCANATGGAGAGCCTGGGCGCCCTGGTGTGA
 TGGTGCT

Restriction Sites: NotI-NotI
ACCN: NM_002973
Insert Size: 4500 bp

OTI Disclaimer: Due to the inherent nature of this plasmid, standard methods to replicate additional amounts of DNA in E. coli are highly likely to result in mutations and/or rearrangements. Therefore, OriGene does not guarantee the capability to replicate this plasmid DNA. Additional amounts of DNA can be purchased from OriGene with batch-specific, full-sequence verification at a reduced cost. Please contact our customer care team at custsupport@origene.com or by calling 301.340.3188 option 3 for pricing and delivery.

The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing variants is recommended prior to use. [More info](#)

Components: The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

Reconstitution Method:

1. Centrifuge at 5,000xg for 5min.
2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.
3. Close the tube and incubate for 10 minutes at room temperature.
4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.
5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.

RefSeq: [NM_002973.2](#), [NP_002964.2](#)

RefSeq Size: 4712 bp

RefSeq ORF: 3942 bp

Locus ID: 6311

UniProt ID: [Q99700](#)

Cytogenetics: 12q24.12

Gene Summary:

This gene belongs to a group of genes that is associated with microsatellite-expansion diseases, a class of neurological and neuromuscular disorders caused by expansion of short stretches of repetitive DNA. The protein encoded by this gene has two globular domains near the N-terminus, one of which contains a clathrin-mediated trans-Golgi signal and an endoplasmic reticulum exit signal. The encoded cytoplasmic protein localizes to the endoplasmic reticulum and plasma membrane, is involved in endocytosis, and modulates mTOR signals, modifying ribosomal translation and mitochondrial function. The N-terminal region of the protein contains a polyglutamine tract of 14-31 residues that can be expanded in the pathogenic state to 32-200 residues. Intermediate length expansions of this tract increase susceptibility to amyotrophic lateral sclerosis, while long expansions of this tract result in spinocerebellar ataxia-2, an autosomal-dominantly inherited, neurodegenerative disorder. Genome-wide association studies indicate that loss-of-function mutations in this gene may be associated with susceptibility to type I diabetes, obesity and hypertension. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Nov 2016] Transcript Variant: This variant (1) represents the longest transcript and encodes the longest isoform (1). Sequence Note: The RefSeq transcript and protein were derived from genomic sequence to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on alignments.