

## Product datasheet for RC218985L3V

## OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com CN: techsupport@origene.cn

## Ataxin 3 (ATXN3) (NM\_030660) Human Tagged ORF Clone Lentiviral Particle

**Product data:** 

**Product Type:** Lentiviral Particles

**Product Name:** Ataxin 3 (ATXN3) (NM\_030660) Human Tagged ORF Clone Lentiviral Particle

Symbol: Ataxin 3

**Synonyms:** AT3; ATX3; JOS; MJD; MJD1; SCA3

Mammalian Cell

Selection:

Puromycin

**Vector:** pLenti-C-Myc-DDK-P2A-Puro (PS100092)

Tag: Myc-DDK
ACCN: NM 030660

ORF Size: 918 bp

**ORF Nucleotide** 

okr Nucleotide

Sequence:
OTI Disclaimer:

Cytogenetics:

The ORF insert of this clone is exactly the same as(RC218985).

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 accurring variations (e.g., polymorphisms), each with its own valid evictors. This

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

**OTI Annotation:** This clone was engineered to express the complete ORF with an expression tag. Expression

varies depending on the nature of the gene.

**RefSeg:** NM 030660.2

 RefSeq Size:
 2533 bp

 RefSeq ORF:
 921 bp

 Locus ID:
 4287

 UniProt ID:
 P54252

Domains: UIM, Josephin

**Protein Families:** Druggable Genome, Transcription Factors

14q32.12





**MW:** 34.8 kDa

**Gene Summary:** 

Machado-Joseph disease, also known as spinocerebellar ataxia-3, is an autosomal dominant neurologic disorder. The protein encoded by this gene contains (CAG)n repeats in the coding region, and the expansion of these repeats from the normal 12-44 to 52-86 is one cause of Machado-Joseph disease. There is a negative correlation between the age of onset and CAG repeat numbers. Alternatively spliced transcript variants encoding different isoforms have been described for this gene. [provided by RefSeq, Jul 2016]