

EXOSC10 antibody C-terminal region Rabbit Polyclonal Antibody Catalog # Al11705

## **Specification**

## **EXOSC10** antibody C-terminal region - Product Information

Application Primary Accession Other Accession Reactivity

Predicted Host Clonality Calculated MW WB <u>Q01780</u> <u>NM\_001001998</u>, <u>NP\_001001998</u> Human, Mouse, Rat, Rabbit, Pig, Horse, Bovine, Dog Mouse, Rabbit, Horse, Bovine, Dog Rabbit Polyclonal 101kDa KDa

## EXOSC10 antibody C-terminal region - Additional Information

Gene ID 5394

Alias Symbol

PM-Scl, PM/Scl-100, PMSCL, PMSCL2, RRP6, Rrp6p, p2, p3, p4

### **Other Names**

Exosome component 10, 3.1.13.-, Autoantigen PM/Scl 2, P100 polymyositis-scleroderma overlap syndrome-associated autoantigen, Polymyositis/scleroderma autoantigen 100 kDa, PM/Scl-100, Polymyositis/scleroderma autoantigen 2, EXOSC10, PMSCL, PMSCL2, RRP6

### Format

Liquid. Purified antibody supplied in 1x PBS buffer with 0.09% (w/v) sodium azide and 2% sucrose.

### **Reconstitution & Storage**

Add 50 ul of distilled water. Final anti-EXOSC10 antibody concentration is 1 mg/ml in PBS buffer with 2% sucrose. For longer periods of storage, store at 20°C. Avoid repeat freeze-thaw cycles.

**Precautions** 

EXOSC10 antibody C-terminal region is for research use only and not for use in diagnostic or therapeutic procedures.

## EXOSC10 antibody C-terminal region - Protein Information

### Name EXOSC10 (HGNC:9138)

### Function

Catalytic component of the RNA exosome complex which has 3'->5' exoribonuclease activity and participates in a multitude of cellular RNA processing and degradation events. In the nucleus, the RNA exosome complex is involved in proper maturation of stable RNA species such as rRNA, snRNA and snoRNA, in the elimination of RNA processing by-products and non-coding 'pervasive' transcripts, such as antisense RNA species and promoter-upstream transcripts (PROMPTs), and of



mRNAs with processing defects, thereby limiting or excluding their export to the cytoplasm. Part of the small subunit (SSU) processome, first precursor of the small eukaryotic ribosomal subunit. During the assembly of the SSU processome in the nucleolus, many ribosome biogenesis factors, an RNA chaperone and ribosomal proteins associate with the nascent pre-rRNA and work in concert to generate RNA folding, modifications, rearrangements and cleavage as well as targeted degradation of pre-ribosomal RNA by the RNA exosome (PubMed:<a

href="http://www.uniprot.org/citations/34516797" target=" blank">34516797</a>). The RNA exosome may be involved in Ig class switch recombination (CSR) and/or Ig variable region somatic hypermutation (SHM) by targeting AICDA deamination activity to transcribed dsDNA substrates. In the cytoplasm, the RNA exosome complex is involved in general mRNA turnover and specifically degrades inherently unstable mRNAs containing AU-rich elements (AREs) within their 3' untranslated regions, and in RNA surveillance pathways, preventing translation of aberrant mRNAs. It seems to be involved in degradation of histone mRNA. EXOSC10 is required for nucleolar localization of C1D and probably mediates the association of MTREX, C1D and MPHOSPH6 with the RNA exosome involved in the maturation of 5.8S rRNA. Plays a role in the recruitment of replication protein A complex (RPA) and RAD51 to DNA double-strand breaks caused by irradiation, contributing to DNA repair by homologous recombination (PubMed:<a href="http://www.uniprot.org/citations/25632158" target="\_blank">25632158</a>, PubMed:<a href="http://www.uniprot.org/citations/31086179" target=" blank">31086179</a>). Regulates levels of damage-induced RNAs in order to prevent DNA-RNA hybrid formation at DNA double-strand breaks and limit DNA end resection after damage (PubMed:<a href="http://www.uniprot.org/citations/31086179" target=" blank">31086179</a>). Plays a role in oocyte development, maturation and survival (By similarity). Required for normal testis development and mitotic division of spermatogonia (By similarity). Plays a role in proper embryo development (By similarity). Required for global protein translation (PubMed:<a href="http://www.uniprot.org/citations/26857222" target=" blank">26857222</a>, PubMed:<a href="http://www.uniprot.org/citations/36912080" target=" blank">36912080</a>). Required for cell proliferation (PubMed:<a href="http://www.uniprot.org/citations/36912080" target=" blank">36912080</a>). Regulates metabolism of C9orf72- derived repeat RNA that can be translated into toxic dipeptide repeat proteins (PubMed: <a href="http://www.uniprot.org/citations/32830871" target=" blank">32830871</a>).

## **Cellular Location**

Cytoplasm. Nucleus. Nucleus, nucleolus. Nucleus, nucleoplasm Note=Strongly enriched in the nucleolus and a small amount has been found in cytoplasm supporting the existence of a nucleolar RNA exosome complex form (PubMed:20531386, PubMed:34516797). Arginine-rich dipeptide repeat proteins expressed from C9orf72-derived repeat RNA cause diffuse nuclear misdistribution of EXOSC10 (PubMed:32830871) Relocates to the DNA double-strand breaks in response to irradiation (PubMed:31086179).

# EXOSC10 antibody C-terminal region - Protocols

Provided below are standard protocols that you may find useful for product applications.

- <u>Western Blot</u>
- <u>Blocking Peptides</u>
- Dot Blot
- Immunohistochemistry
- Immunofluorescence
- Immunoprecipitation
- Flow Cytomety
- <u>Cell Culture</u>

## EXOSC10 antibody C-terminal region - Images





van (2007) RNA 13 (7), 1027-1035 Reconstitution and Storage:For short term use, store at 2-8C up to 1 week. For long term storage, store at -20C in small aliquots to prevent freeze-thaw cycles.