

**EXOSC1 Antibody (C-term)**  
**Affinity Purified Rabbit Polyclonal Antibody (Pab)**  
**Catalog # AP18442b**

**Specification**

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**EXOSC1 Antibody (C-term) - Product Information**

Application	WB,E
Primary Accession	<a href="#">O9Y3B2</a>
Other Accession	<a href="#">O9DAA6</a> , <a href="#">NP_057130.1</a>
Reactivity	Human
Predicted	Mouse
Host	Rabbit
Clonality	Polyclonal
Isotype	Rabbit IgG
Calculated MW	21452
Antigen Region	165-191

**EXOSC1 Antibody (C-term) - Additional Information**

**Gene ID** 51013

**Other Names**

Exosome complex component CSL4, Exosome component 1, EXOSC1, CSL4

**Target/Specificity**

This EXOSC1 antibody is generated from rabbits immunized with a KLH conjugated synthetic peptide between 165-191 amino acids from the C-terminal region of human EXOSC1.

**Dilution**

WB~~1:1000

E~~Use at an assay dependent concentration.

**Format**

Purified polyclonal antibody supplied in PBS with 0.09% (W/V) sodium azide. This antibody is purified through a protein A column, followed by peptide affinity purification.

**Storage**

Maintain refrigerated at 2-8°C for up to 2 weeks. For long term storage store at -20°C in small aliquots to prevent freeze-thaw cycles.

**Precautions**

EXOSC1 Antibody (C-term) is for research use only and not for use in diagnostic or therapeutic procedures.

**EXOSC1 Antibody (C-term) - Protein Information**

**Name** EXOSC1

## Synonyms CSL4

**Function** Non-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. 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. The catalytic inactive RNA exosome core complex of 9 subunits (Exo-9) is proposed to play a pivotal role in the binding and presentation of RNA for ribonucleolysis, and to serve as a scaffold for the association with catalytic subunits and accessory proteins or complexes. EXOSC1 as peripheral part of the Exo-9 complex stabilizes the hexameric ring of RNase PH-domain subunits through contacts with EXOSC6 and EXOSC8.

## Cellular Location

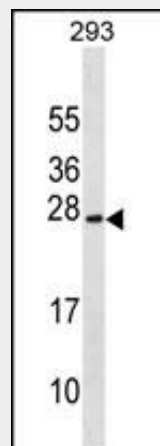
Nucleus, nucleolus. Nucleus. Cytoplasm

## EXOSC1 Antibody (C-term) - Protocols

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

- [Western Blot](#)
- [Blocking Peptides](#)
- [Dot Blot](#)
- [Immunohistochemistry](#)
- [Immunofluorescence](#)
- [Immunoprecipitation](#)
- [Flow Cytometry](#)
- [Cell Culture](#)

## EXOSC1 Antibody (C-term) - Images



EXOSC1 Antibody (C-term) (Cat. #AP18442b) western blot analysis in 293 cell line lysates (35ug/lane). This demonstrates the EXOSC1 Antibody detected the EXOSC1 protein (arrow).

**EXOSC1 Antibody (C-term) - Background**

This gene encodes a core component of the exosome. The mammalian exosome is required for rapid degradation of AU rich element-containing RNAs but not for poly(A) shortening. The association of this protein with the exosome is mediated by protein-protein interactions with ribosomal RNA-processing protein 42 and ribosomal RNA-processing protein 46.

**EXOSC1 Antibody (C-term) - References**

Andersen, J.S., et al. Nature 433(7021):77-83(2005)  
Lehner, B., et al. Genome Res. 14(7):1315-1323(2004)  
Deloukas, P., et al. Nature 429(6990):375-381(2004)  
Raijmakers, R., et al. J. Mol. Biol. 323(4):653-663(2002)  
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