

[Go to Product page](#)

Datasheet for ABIN7152291

**anti-EXOSC8 antibody (AA 2-276) (FITC)**

## Overview

|                      |  |
|----------------------|--|
| Quantity:            | 100 µg                                     |
| Target:              | EXOSC8                                     |
| Binding Specificity: | AA 2-276                                   |
| Reactivity:          | Human                                      |
| Host:                | Rabbit                                     |
| Clonality:           | Polyclonal                                 |
| Conjugate:           | This EXOSC8 antibody is conjugated to FITC |
| Application:         | Please inquire                             |

## Product Details

|                   |   |
|-------------------|---|
| Immunogen:        | Recombinant Human Exosome complex component RRP43 protein (2-276AA) |
| Isotype:          | IgG   |
| Cross-Reactivity: | Human   |
| Purification:     | >95%, Protein G purified  |

## Target Details

|                   |  |
|-------------------|--|
| Target:           | EXOSC8   |
| Alternative Name: | EXOSC8 ( <a href="#">EXOSC8 Products</a> )   |
| Background:       | Background: Non-catalytic component of the RNA exosome complex which has 3'→5' exoribonuclease activity and participates in a multitude of cellular RNA processing and |

## Target Details

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. EXOSC8 binds to ARE-containing RNAs.

Aliases: bA421P11.3 antibody, CBP interacting protein 3 antibody, CIP3 antibody, EAP2 antibody, EC 3.1.13 antibody, EXOS8\_HUMAN antibody, EXOSC8 antibody, Exosome complex component RRP43 antibody, Exosome complex exonuclease RRP43 antibody, Exosome component 8 antibody, OIP-2 antibody, OIP2 antibody, Opa interacting protein 2 antibody, Opa-interacting protein 2 antibody, OTTHUMP00000042274 antibody, p9 antibody, Ribosomal RNA processing protein 43 antibody, Ribosomal RNA processing protein 43, S. cerevisiae, homolog of antibody, Ribosomal RNA-processing protein 43 antibody, RP11 421P11.3 antibody, RRP43 antibody, Rrp43p antibody

UniProt: [Q96B26](#)

Pathways: [SARS-CoV-2 Protein Interactome](#)

## Application Details

Restrictions: For Research Use only

## Handling

Format: Liquid

Buffer: Preservative: 0.03 % Proclin 300  
Constituents: 50 % Glycerol, 0.01M PBS, PH 7.4

Preservative: ProClin

## Handling

---

|                    |   |
|--------------------|---|
| Precaution of Use: | This product contains ProClin: a POISONOUS AND HAZARDOUS SUBSTANCE which should be handled by trained staff only. |
| Storage:           | -20 °C,-80 °C   |
| Storage Comment:   | Upon receipt, store at -20°C or -80°C. Avoid repeated freeze.   |