

ELAC2 cleaves the 3' end of pre-tRNA

Jarrous, N., Levinger, L., May, B., Motorin, Y.

European Bioinformatics Institute, New York University Langone Medical Center, Ontario Institute for Cancer Research, Oregon Health and Science University.

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03/05/2024

Introduction

Reactome is open-source, open access, manually curated and peer-reviewed pathway database. Pathway annotations are authored by expert biologists, in collaboration with Reactome editorial staff and cross-referenced to many bioinformatics databases. A system of evidence tracking ensures that all assertions are backed up by the primary literature. Reactome is used by clinicians, geneticists, genomics researchers, and molecular biologists to interpret the results of high-throughput experimental studies, by bioinformaticians seeking to develop novel algorithms for mining knowledge from genomic studies, and by systems biologists building predictive models of normal and disease variant pathways.

The development of Reactome is supported by grants from the US National Institutes of Health (P41 HG003751), University of Toronto (CFREF Medicine by Design), European Union (EU STRP, EMI-CD), and the European Molecular Biology Laboratory (EBI Industry program).

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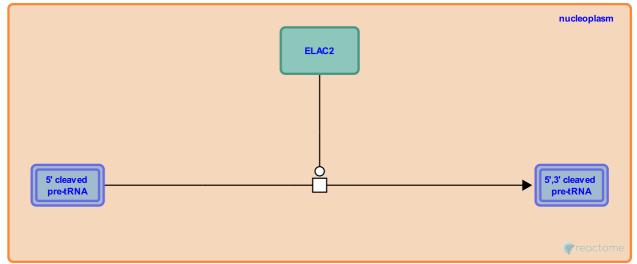
This document contains 1 reaction (see Table of Contents)

ELAC2 cleaves the 3' end of pre-tRNA 7

Stable identifier: R-HSA-5696815

Type: transition

Compartments: nucleoplasm



ELAC2, a RNaseZ activity in humans, endonucleolytically cleaves pre-tRNAs near the 3' end, generating a 3' hydroxyl group to which the nucleotides CCA will be polymerized (Takaku et al. 2004, Yan et al. 2006, Elbarbary et al. 2008). ELAC2 is located in both the nucleus and mitochondria (Brzezniak et al. 2011). ELAC1, a homologue of ELAC2, is located in the cytosol (Rossmanith 2011) and cleaves 3' ends of tRNAs in vitro (Takaku et al. 2003).

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Editions

2015-05-29	Authored, Edited	May, B.
2015-08-11	Reviewed	Levinger, L.
2015-08-25	Reviewed	Motorin, Y.
2015-10-24	Reviewed	Jarrous, N.