

FTMT 24mer oxidises 4Fe2+ to

4Fe(3+)O(OH)

D'Eustachio, P., Jassal, B.

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

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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.

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Literature references

- Fabregat, A., Sidiropoulos, K., Viteri, G., Forner, O., Marin-Garcia, P., Arnau, V. et al. (2017). Reactome pathway analysis: a high-performance in-memory approach. *BMC bioinformatics, 18,* 142. 7
- Sidiropoulos, K., Viteri, G., Sevilla, C., Jupe, S., Webber, M., Orlic-Milacic, M. et al. (2017). Reactome enhanced pathway visualization. *Bioinformatics*, 33, 3461-3467. A
- Fabregat, A., Jupe, S., Matthews, L., Sidiropoulos, K., Gillespie, M., Garapati, P. et al. (2018). The Reactome Pathway Knowledgebase. *Nucleic Acids Res, 46*, D649-D655. ↗
- Fabregat, A., Korninger, F., Viteri, G., Sidiropoulos, K., Marin-Garcia, P., Ping, P. et al. (2018). Reactome graph database: Efficient access to complex pathway data. *PLoS computational biology*, *14*, e1005968. *オ*

This document contains 1 reaction (see Table of Contents)

FTMT 24mer oxidises 4Fe2+ to 4Fe(3+)O(OH) 7

Stable identifier: R-HSA-5691107

Type: transition

Compartments: mitochondrial matrix



Mitochondrial ferritin (FTMT) is specifically taken up by the mitochondria and processed to a mature protein that assembles into functional ferritin shells. It is a homooligomer of 24 subunits, is roughly spherical and contains a central cavity into which the mineral iron core is deposited. FTMT possesses ferroxidase activity. Iron is taken up in the ferrous form (Fe2+) and deposited as ferric hydroxide (Fe(3+)O(OH)) after oxidation. FTMT may play an important role in the regulation of iron homeostasis in the mitochondrion (Levi et al. 2001, Langlois d'Estaintot et al. 2004).

Literature references

Drysdale, J., Bosisio, M., Arosio, P., Levi, S., Sanford, D., Corsi, B. et al. (2001). A human mitochondrial ferritin encoded by an intronless gene. J. Biol. Chem., 276, 24437-40.

Santambrogio, P., Précigoux, G., Gallois, B., Arosio, P., Levi, S., Granier, T. et al. (2004). Crystal structure and biochemical properties of the human mitochondrial ferritin and its mutant Ser144Ala. J. Mol. Biol., 340, 277-93.

Editions

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