

MT3 binds cadmium

Atrian, S., May, B.

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

The contents of this document may be freely copied and distributed in any media, provided the authors, plus the institutions, are credited, as stated under the terms of [Creative Commons Attribution 4.0 International \(CC BY 4.0\) License](#). For more information see our [license](#).

02/05/2024

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).

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. [↗](#)
- Sidiropoulos, K., Viteri, G., Sevilla, C., Jupe, S., Webber, M., Orlic-Milacic, M. et al. (2017). Reactome enhanced pathway visualization. *Bioinformatics*, 33, 3461-3467. [↗](#)
- 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. [↗](#)

Reactome database release: 88

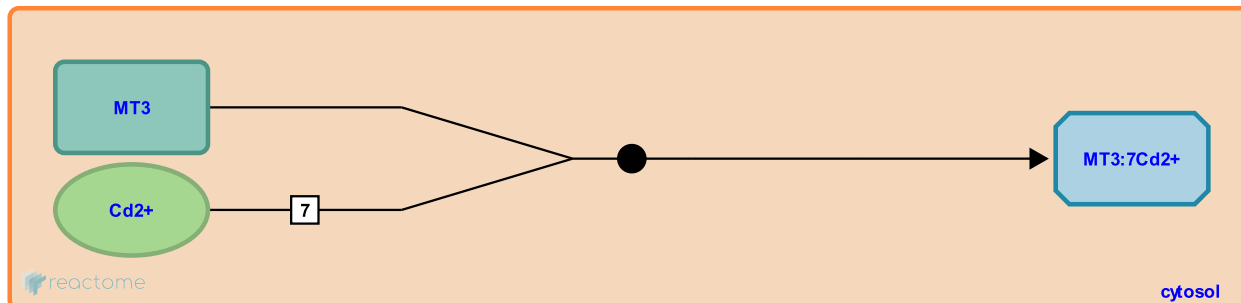
This document contains 1 reaction ([see Table of Contents](#))

MT3 binds cadmium [↗](#)

Stable identifier: R-HSA-5662620

Type: binding

Compartments: cytosol



The metallothionein binds 7 atoms of cadmium(II) in 2 clusters, 4 atoms at the C-terminal alpha domain and 3 atoms at the N-terminal beta domain (Palumaa et al. 2002, Palumaa et al. 2005, Wu et al. 2014). MT3 binds cadmium with a lower affinity than MT2A does (Palumaa et al. 2005). MT3 can bind more than 7 cadmium atoms, however the MT3:7Cd²⁺ complex is most prevalent (Palumaa et al. 2002, Palumaa et al. 2005). Exposure of MT3:7Cd²⁺ to nitric oxide causes release of cadmium (Wang et al. 2008).

Literature references

- Sillard, R., Jörnvall, H., Kruusel, K., Palumaa, P., Kangur, L., Tammiste, I. (2005). Metal binding of metallothionein-3 versus metallothionein-2: lower affinity and higher plasticity. *Biochim. Biophys. Acta*, 1747, 205-11. [↗](#)
- Li, S., Cui, T., Wu, D., Li, H., Wu, F., Tang, D. et al. (2014). High level expression, efficient purification, and bioactivity of recombinant human metallothionein 3 (rhMT3) from methylotrophic yeast *Pichia pastoris*. *Protein Expr. Purif.*, 101, 121-6. [↗](#)
- Wang, H., Li, H., Huang, ZX., Cai, B., Sun, H. (2008). The effect of nitric oxide on metal release from metallothionein-3: gradual unfolding of the protein. *J. Biol. Inorg. Chem.*, 13, 411-9. [↗](#)
- Sillard, R., Njunkova, O., Jörnvall, H., Palumaa, P., Pokras, L., Eriste, E. (2002). Brain-specific metallothionein-3 has higher metal-binding capacity than ubiquitous metallothioneins and binds metals noncooperatively. *Biochemistry*, 41, 6158-63. [↗](#)

Editions

2015-01-10	Authored, Edited	May, B.
2015-09-19	Reviewed	Atrian, S.