

ACO1:4Fe-4S isomerises CIT to ISCIT

D'Eustachio, P., Jassal, 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](#).

04/05/2024

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

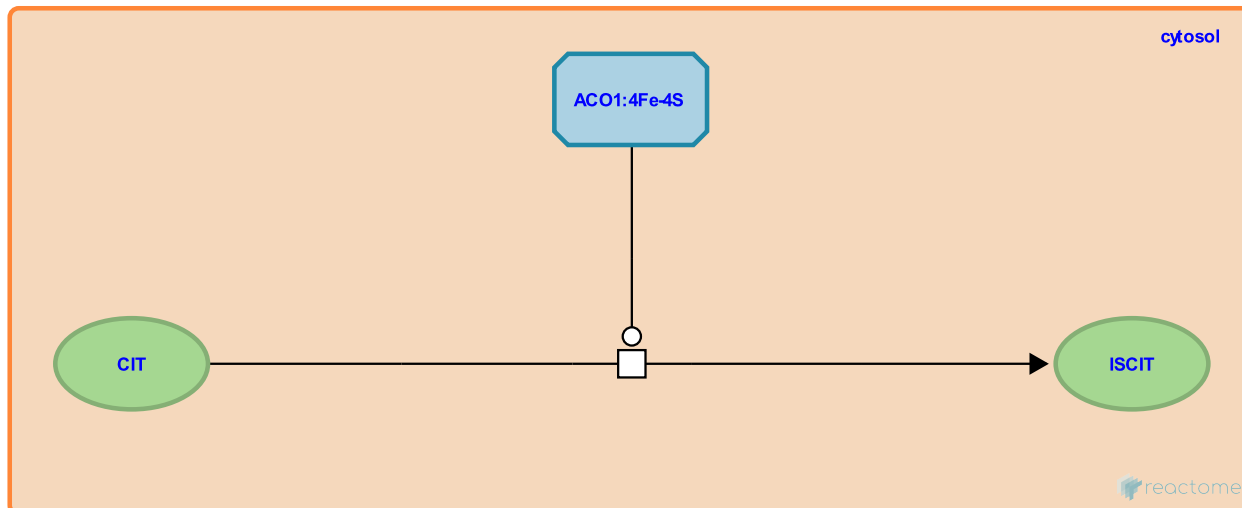
This document contains 1 reaction ([see Table of Contents](#))

ACO1:4Fe-4S isomerises CIT to ISCIT [↗](#)

Stable identifier: R-HSA-5690911

Type: transition

Compartments: cytosol



Cytoplasmic aconitate hydratase (ACO1, iron regulatory protein 1, IRP1) functions either as an RNA binding protein that regulates the uptake, sequestration, and utilisation of iron or an enzyme that isomerises citrate to isocitrate, depending on changes in cellular iron levels. Under iron-replete conditions, ACO1 binds the cofactor 4Fe-4S cluster and acts as an aconitase, isomerising citrate (CIT) to isocitrate (ISCIT) (Kaptain et al. 1991, Philpott et al. 1994, Dupuy et al. 2006).

Literature references

- Fontecilla-Camps, JC., Darnault, C., Carpentier, P., Volbeda, A., Dupuy, J., Moulis, JM. (2006). Crystal structure of human iron regulatory protein 1 as cytosolic aconitase. *Structure*, 14, 129-39. [↗](#)
- Haile, D., Downey, WE., Orloff, DG., Harford, JB., Philpott, C., Kaptain, S. et al. (1991). A regulated RNA binding protein also possesses aconitase activity. *Proc. Natl. Acad. Sci. U.S.A.*, 88, 10109-13. [↗](#)
- Philpott, CC., Klausner, RD., Rouault, TA. (1994). The bifunctional iron-responsive element binding protein/cytosolic aconitase: the role of active-site residues in ligand binding and regulation. *Proc. Natl. Acad. Sci. U.S.A.*, 91, 7321-5. [↗](#)
- Ortega, F., Moreno-Navarrete, JM., Fernández-Real, JM., Moreno, M., Xifra, G., Ricart, W. (2015). Cytosolic aconitase activity sustains adipogenic capacity of adipose tissue connecting iron metabolism and adipogenesis. *FASEB J*, 29, 1529-39. [↗](#)

Editions

2015-04-30	Authored, Edited	Jassal, B.
2017-01-06	Reviewed	D'Eustachio, P.
2024-02-08	Reviewed	D'Eustachio, P.