

# LDHAL6B reduces PYR to LACT

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](#).

26/10/2020

## 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: 74

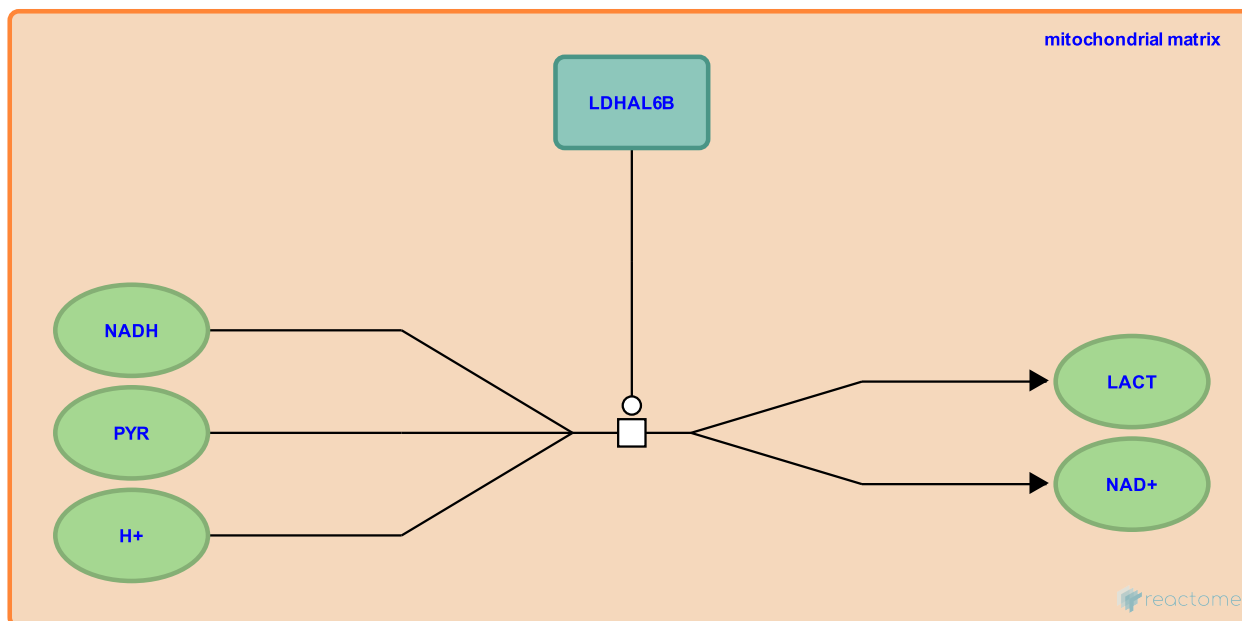
This document contains 1 reaction ([see Table of Contents](#))

## LDHAL6B reduces PYR to LACT ↗

**Stable identifier:** R-HSA-6807826

**Type:** transition

**Compartments:** mitochondrial matrix



LDHAL6B (L-lactate dehydrogenase A-like 6B) catalyzes the reaction of PYR (pyruvate) and NADH + H<sup>+</sup> to form LACT (lactate) and NAD<sup>+</sup>. The LDHAL6B protein is the inferred product of an open reading frame transcribed in the testis (Wang et al. 2005). Its mitochondrial localization is inferred from the presence of a mitochondrial localization sequence at its amino terminus (Holmes and Goldberg 2009). A physiological role for lactate formation from the abundant pyruvate and NADH expected in rapidly respiring mitochondria is not straightforward to imagine, however.

### Literature references

Holmes, RS., Goldberg, E. (2009). Computational analyses of mammalian lactate dehydrogenases: human, mouse, opossum and platypus LDHs. *Comput Biol Chem*, 33, 379-85. ↗

Wang, H., Zhou, Z., Lu, L., Xu, Z., Sha, J. (2005). Cloning and characterization of a novel intronless lactate dehydrogenase gene in human testis. *Int. J. Mol. Med.*, 15, 949-53. ↗

### Editions

2015-11-09	Authored, Edited, Revised	D'Eustachio, P.
2015-11-09	Reviewed	Jassal, B.