

Second phosphorylation of IRAK1 by IRAK4 bound to MyD88: activated TLR 7/8 or 9

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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

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Reactome database release: 81

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

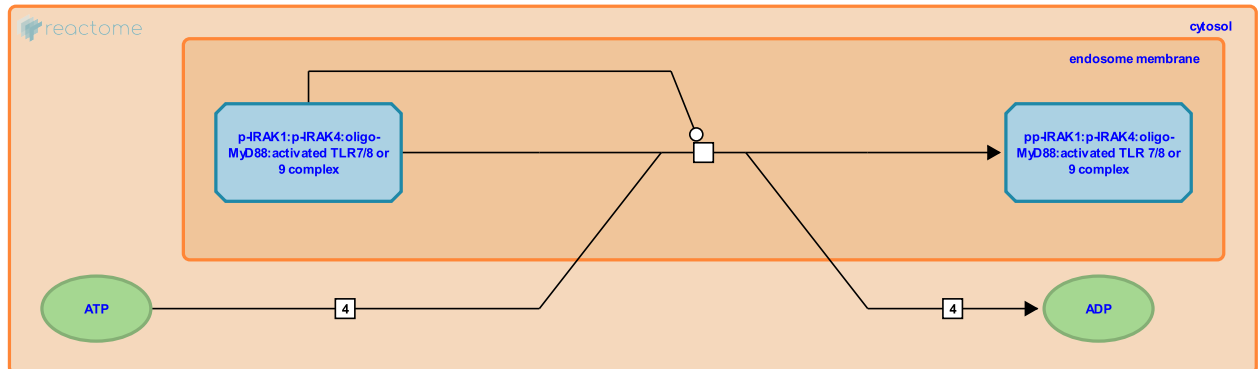
Second phosphorylation of IRAK1 by IRAK4 bound to MyD88: activated TLR 7/8 or 9



Stable identifier: R-HSA-975134

Type: transition

Compartments: endosome membrane, cytosol



Second, Thr387 in the activation loop is phosphorylated, leading to full enzymatic activity.

Literature references

Wesche, H., Li, S., Martin, MU., Knop, J., Neumann, D., Cao, P. et al. (2004). Sequential autophosphorylation steps in the interleukin-1 Receptor-associated Kinase-1 Regulate its Availability as an Adapter in Interleukin-1 Signaling. *J Biol Chem*, 279, 5227-36. [↗](#)

Editions

2010-08-25	Authored	Shamovsky, V.
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