

PIAS4 SUMOylates PPARA with SUMO1

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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: 77

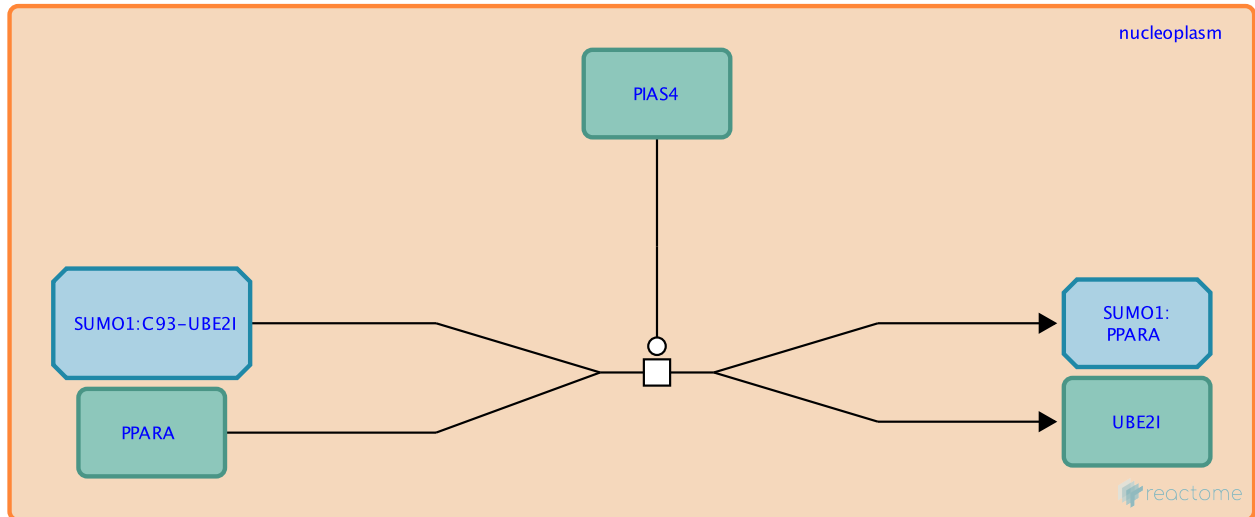
This document contains 1 reaction ([see Table of Contents](#))

PIAS4 SUMOylates PPARA with SUMO1 [↗](#)

Stable identifier: R-HSA-4341070

Type: transition

Compartments: nucleoplasm



PIAS4 SUMOylates PPARA at lysine-185 with SUMO1 (Pourcet et al. 2010). SUMOylation decreases the transactivation activity of PPARA. SUMOylation is decreased in the presence of ligand of PPARA.

Literature references

Pourcet, B., Pineda-Torra, I., Derudas, B., Staels, B., Glineur, C. (2010). SUMOylation of human peroxisome proliferator-activated receptor alpha inhibits its trans-activity through the recruitment of the nuclear corepressor NCoR. *J. Biol. Chem.*, 285, 5983-92. [↗](#)

Editions

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