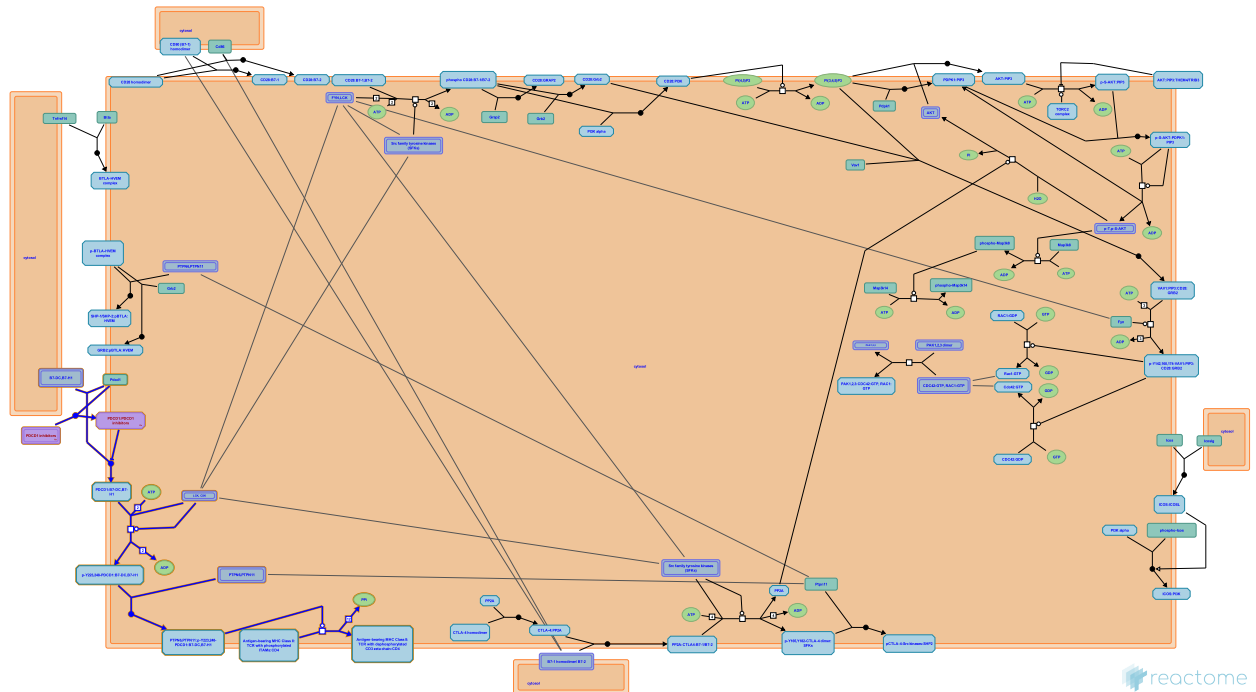


PD-1 signaling



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

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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: 75

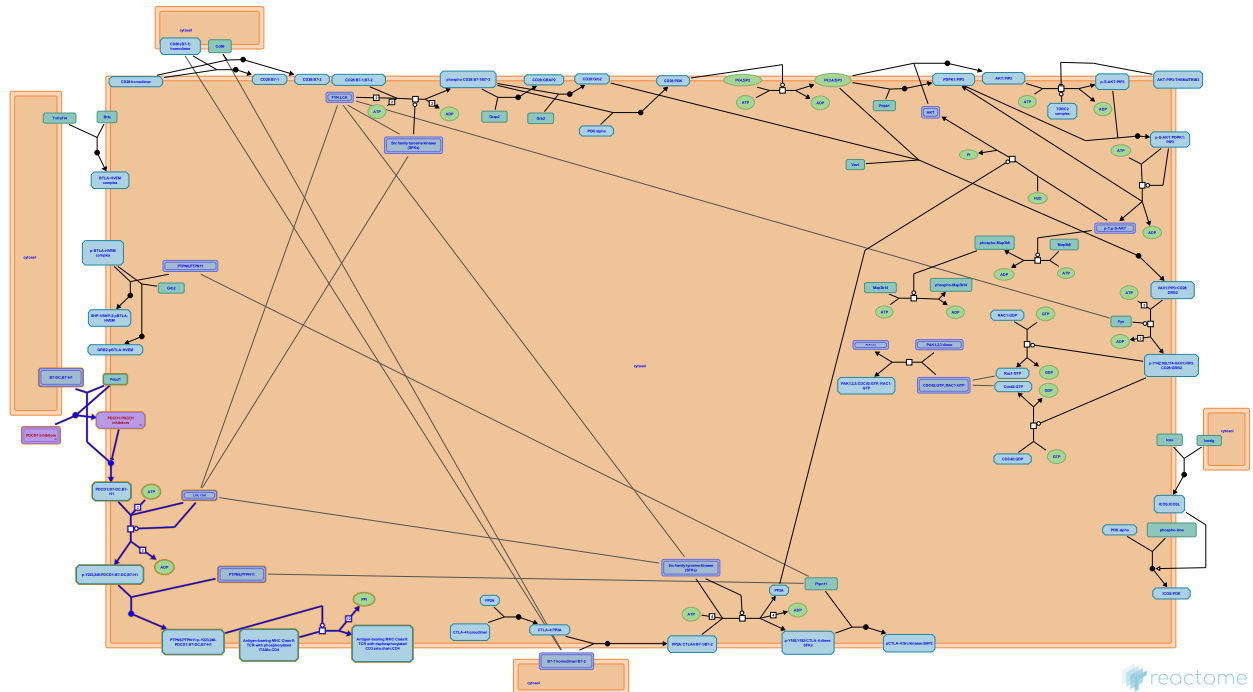
This document contains 1 pathway and 5 reactions ([see Table of Contents](#))

PD-1 signaling ↗

Stable identifier: R-MMU-389948

Compartments: plasma membrane

Inferred from: PD-1 signaling (Homo sapiens)



This event has been computationally inferred from an event that has been demonstrated in another species.

The inference is based on the homology mapping from PANTHER. Briefly, reactions for which all involved PhysicalEntities (in input, output and catalyst) have a mapped orthologue/paralogue (for complexes at least 75% of components must have a mapping) are inferred to the other species. High level events are also inferred for these events to allow for easier navigation.

[More details and caveats of the event inference in Reactome.](/electronic_inference_compara.html) For details on PANTHER see also: <http://www.pantherdb.org/about.jsp>

PD-1 binds B7DC and B7H1 ↗

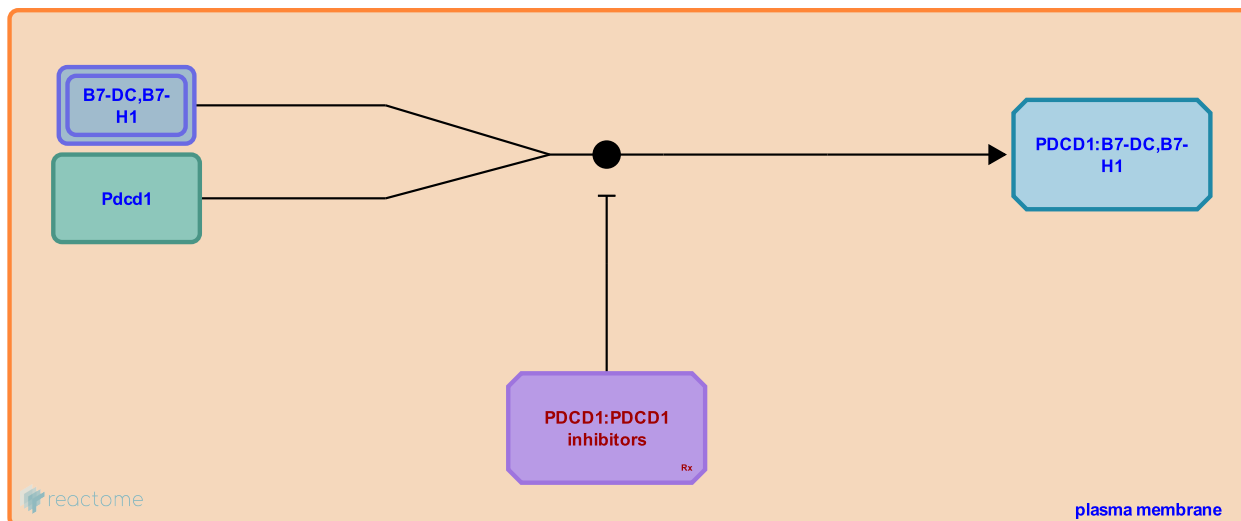
Location: [PD-1 signaling](#)

Stable identifier: R-MMU-388828

Type: binding

Compartments: plasma membrane

Inferred from: [PD-1 binds B7DC and B7H1 \(Homo sapiens\)](#)



This event has been computationally inferred from an event that has been demonstrated in another species.

The inference is based on the homology mapping from PANTHER. Briefly, reactions for which all involved PhysicalEntities (in input, output and catalyst) have a mapped orthologue/paralogue (for complexes at least 75% of components must have a mapping) are inferred to the other species. High level events are also inferred for these events to allow for easier navigation.

[More details and caveats of the event inference in Reactome.](/electronic_inference_compara.html) For details on PANTHER see also: <http://www.pantherdb.org/about.jsp>

Followed by: [Phosphorylation of PD-1](#)

PDCD1 binds PDCD1 inhibitors ↗

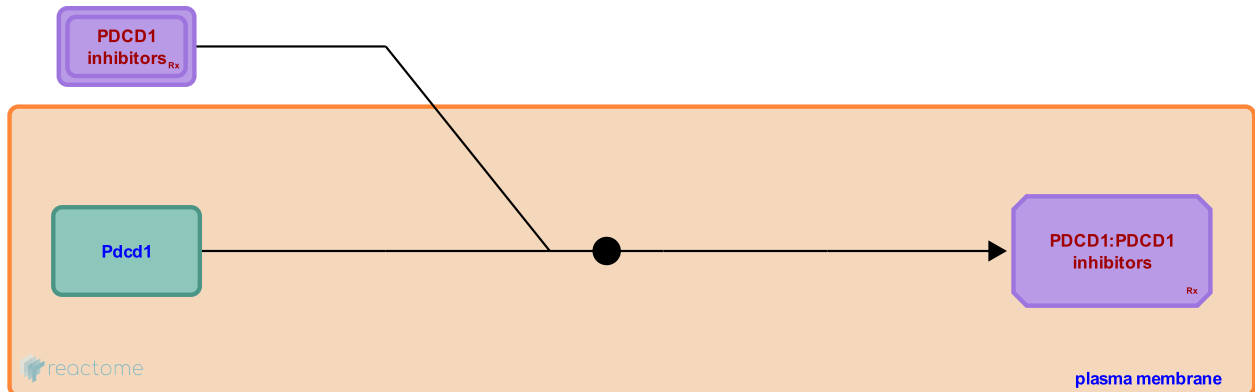
Location: [PD-1 signaling](#)

Stable identifier: R-MMU-9679421

Type: binding

Compartments: plasma membrane, extracellular region

Inferred from: [PDCD1 binds PDCD1 inhibitors \(Homo sapiens\)](#)



This event has been computationally inferred from an event that has been demonstrated in another species.

The inference is based on the homology mapping from PANTHER. Briefly, reactions for which all involved PhysicalEntities (in input, output and catalyst) have a mapped orthologue/paralogue (for complexes at least 75% of components must have a mapping) are inferred to the other species. High level events are also inferred for these events to allow for easier navigation.

[More details and caveats of the event inference in Reactome.](/electronic_inference_compara.html) For details on PANTHER see also: <http://www.pantherdb.org/about.jsp>

Phosphorylation of PD-1 ↗

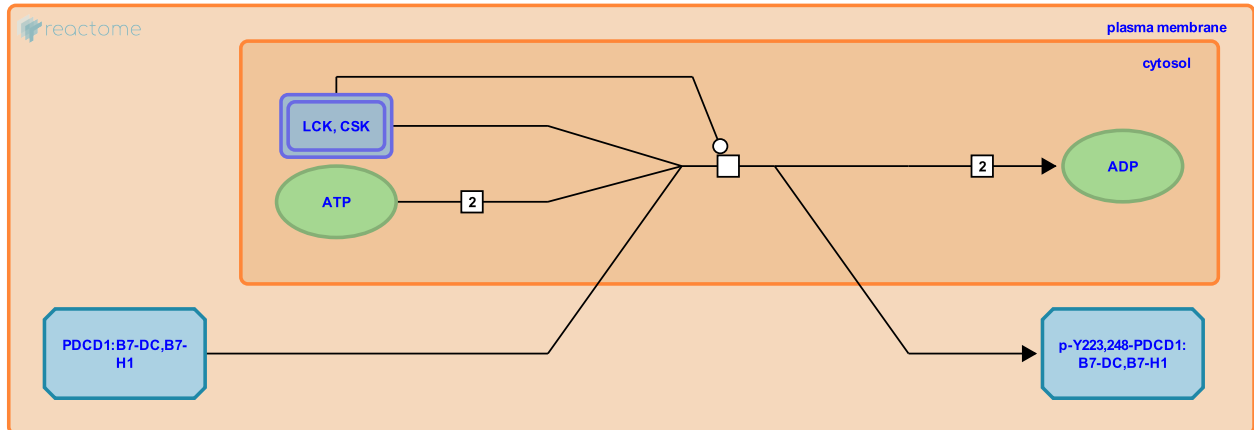
Location: [PD-1 signaling](#)

Stable identifier: R-MMU-389762

Type: transition

Compartments: cytosol, plasma membrane

Inferred from: [Phosphorylation of PD-1 \(Homo sapiens\)](#)



This event has been computationally inferred from an event that has been demonstrated in another species.

The inference is based on the homology mapping from PANTHER. Briefly, reactions for which all involved PhysicalEntities (in input, output and catalyst) have a mapped orthologue/paralogue (for complexes at least 75% of components must have a mapping) are inferred to the other species. High level events are also inferred for these events to allow for easier navigation.

[More details and caveats of the event inference in Reactome.](/electronic_inference_compara.html) For details on PANTHER see also: <http://www.pantherdb.org/about.jsp>

Preceded by: [PD-1 binds B7DC and B7H1](#)

Followed by: [Interaction of SHP-1 or SHP-2 with phospho PD-1](#)

Interaction of SHP-1 or SHP-2 with phospho PD-1 [↗](#)

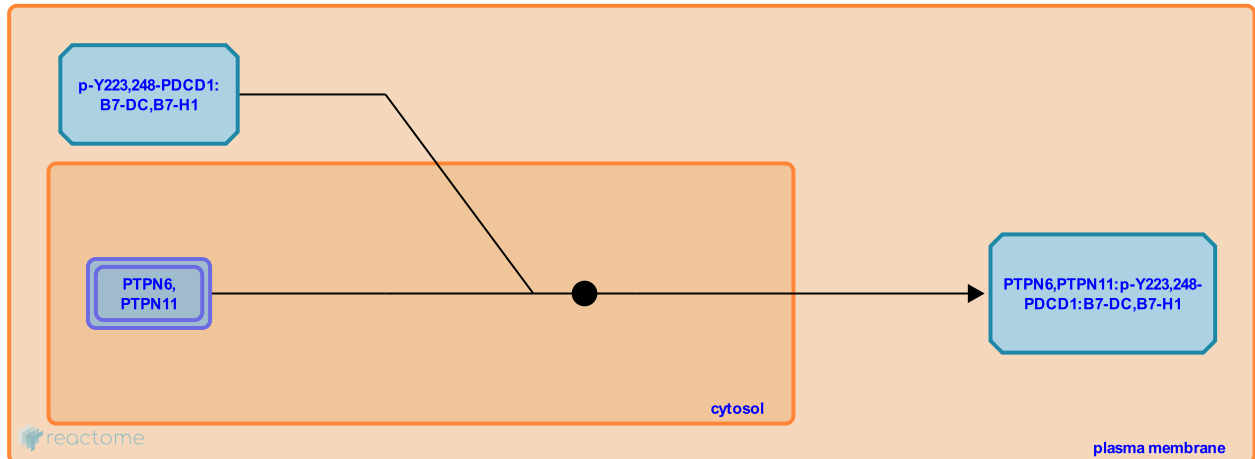
Location: [PD-1 signaling](#)

Stable identifier: R-MMU-389759

Type: binding

Compartments: cytosol, plasma membrane

Inferred from: [Interaction of SHP-1 or SHP-2 with phospho PD-1 \(Homo sapiens\)](#)



This event has been computationally inferred from an event that has been demonstrated in another species.

The inference is based on the homology mapping from PANTHER. Briefly, reactions for which all involved PhysicalEntities (in input, output and catalyst) have a mapped orthologue/paralogue (for complexes at least 75% of components must have a mapping) are inferred to the other species. High level events are also inferred for these events to allow for easier navigation.

[More details and caveats of the event inference in Reactome.](/electronic_inference_compara.html) For details on PANTHER see also: <http://www.pantherdb.org/about.jsp>

Preceded by: [Phosphorylation of PD-1](#)

Followed by: [Dephosphorylation of CD3-zeta by PD-1 bound phosphatases](#)

Dephosphorylation of CD3-zeta by PD-1 bound phosphatases ↗

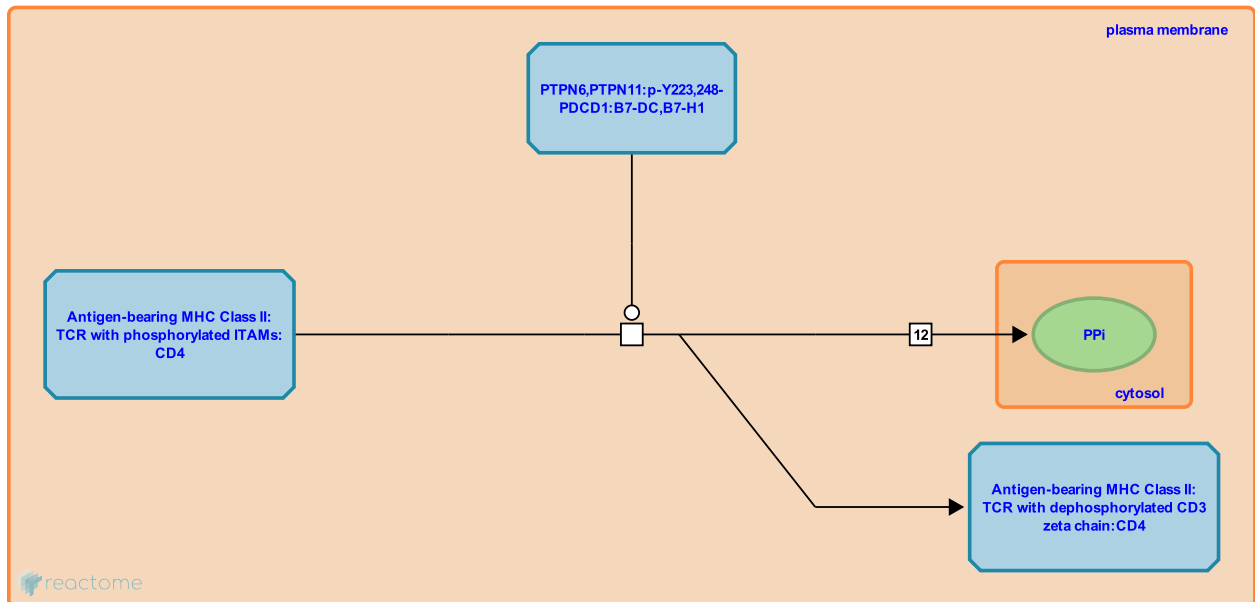
Location: PD-1 signaling

Stable identifier: R-MMU-389758

Type: transition

Compartments: plasma membrane, cytosol

Inferred from: [Dephosphorylation of CD3-zeta by PD-1 bound phosphatases \(Homo sapiens\)](#)



This event has been computationally inferred from an event that has been demonstrated in another species.

The inference is based on the homology mapping from PANTHER. Briefly, reactions for which all involved PhysicalEntities (in input, output and catalyst) have a mapped orthologue/paralogue (for complexes at least 75% of components must have a mapping) are inferred to the other species. High level events are also inferred for these events to allow for easier navigation.

[More details and caveats of the event inference in Reactome.](/electronic_inference_compara.html) For details on PANTHER see also: <http://www.pantherdb.org/about.jsp>

Preceded by: [Interaction of SHP-1 or SHP-2 with phospho PD-1](#)

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