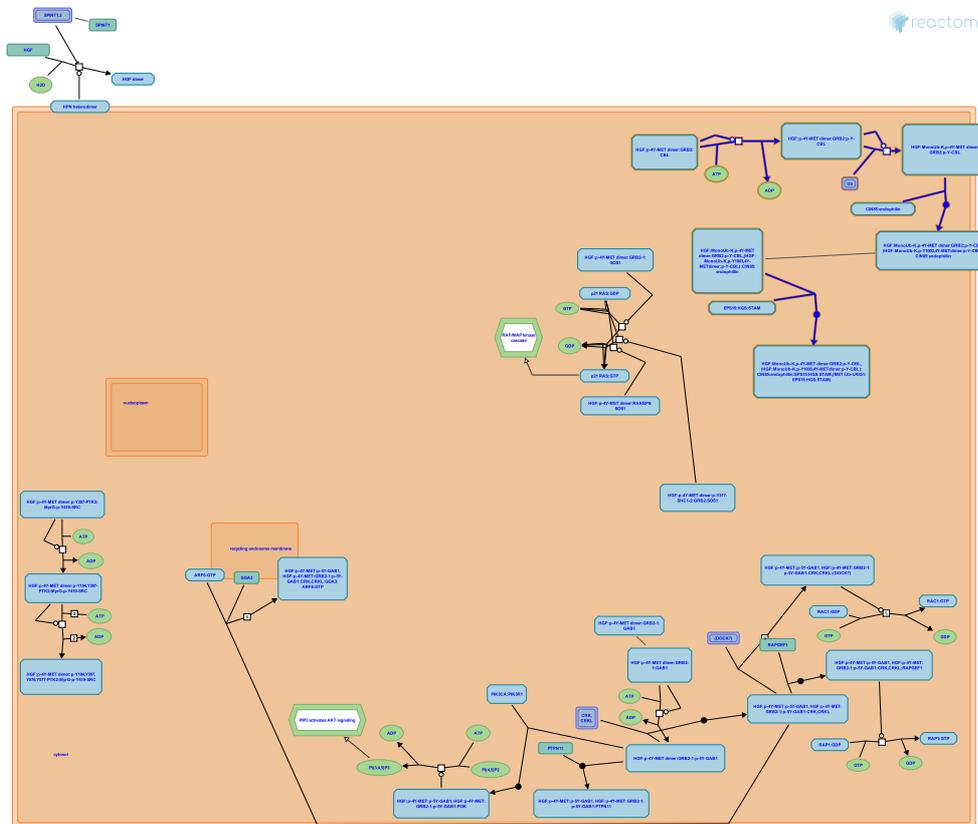


Negative regulation of MET activity



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.

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

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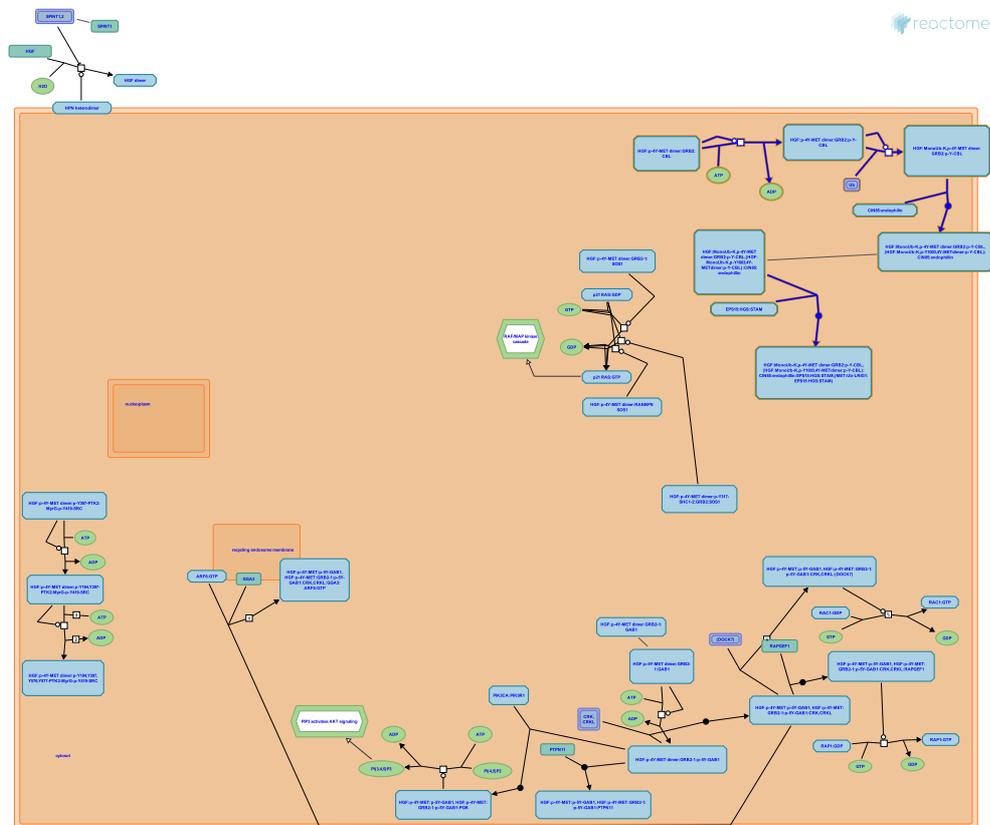
Reactome database release: 76

This document contains 1 pathway and 4 reactions ([see Table of Contents](#))

Negative regulation of MET activity ↗

Stable identifier: R-CFA-6807004

Inferred from: Negative regulation of MET activity (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>

MET phosphorylates CBL ↗

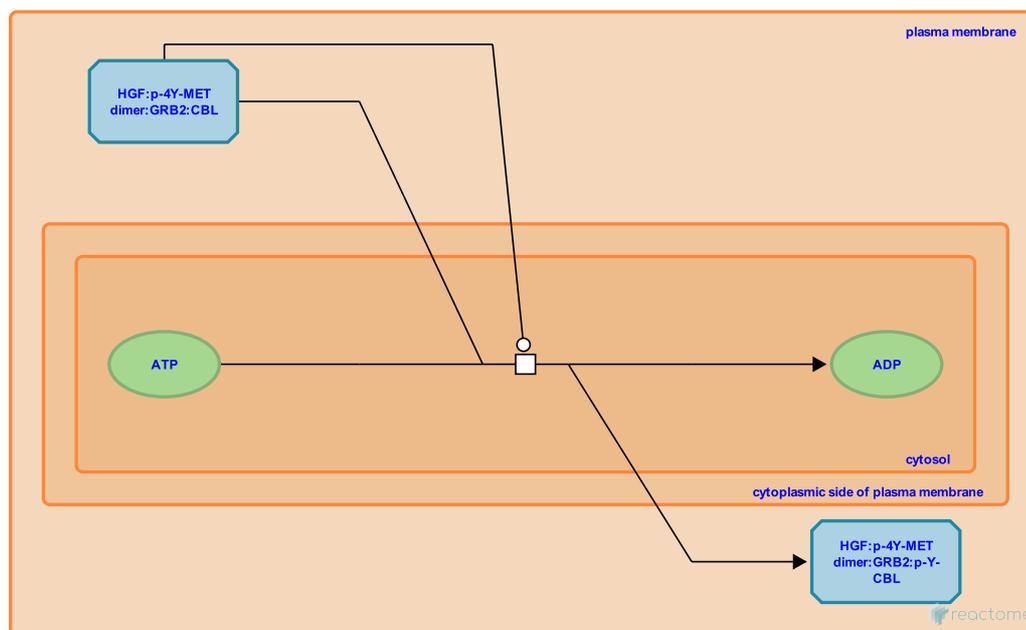
Location: [Negative regulation of MET activity](#)

Stable identifier: R-CFA-8875451

Type: transition

Compartments: cytosol, plasma membrane

Inferred from: [MET phosphorylates CBL \(Homo sapiens\)](#)



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: [CBL monoubiquitinates activated MET](#)

CBL monoubiquitinates activated MET ↗

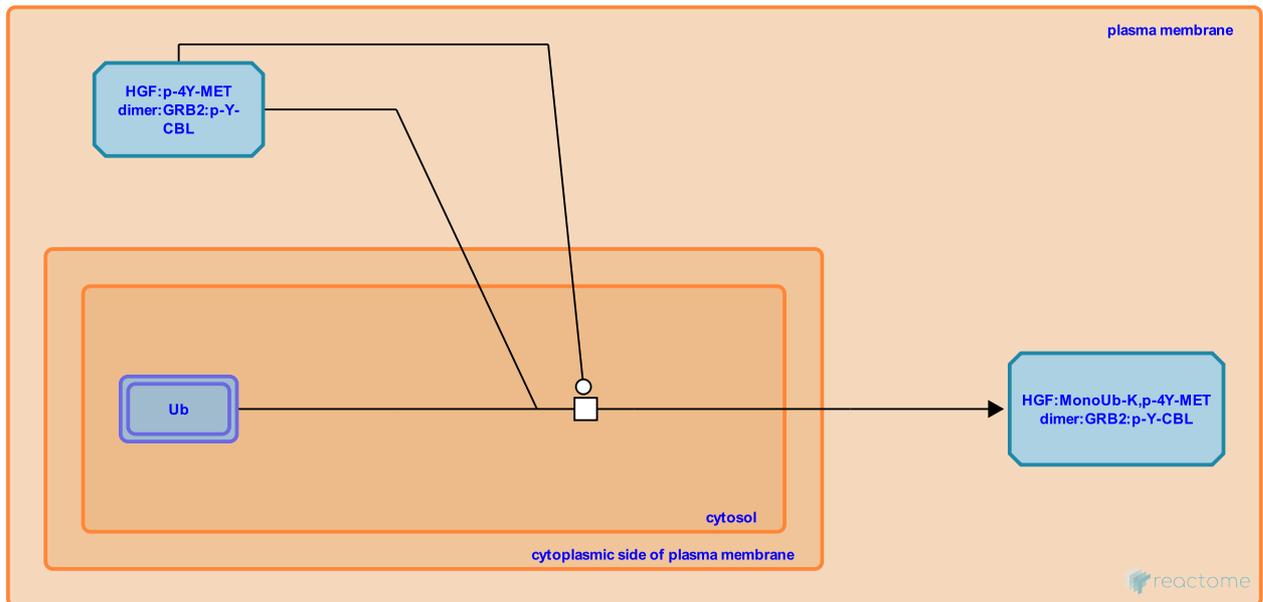
Location: [Negative regulation of MET activity](#)

Stable identifier: R-CFA-8875183

Type: transition

Compartments: cytosol, plasma membrane

Inferred from: [CBL monoubiquitinates activated MET \(Homo sapiens\)](#)



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: [MET phosphorylates CBL](#)

Followed by: [CBL recruits CIN85:endophilin to ubiquitinated MET](#)

CBL recruits CIN85:endophilin to ubiquitinated MET ↗

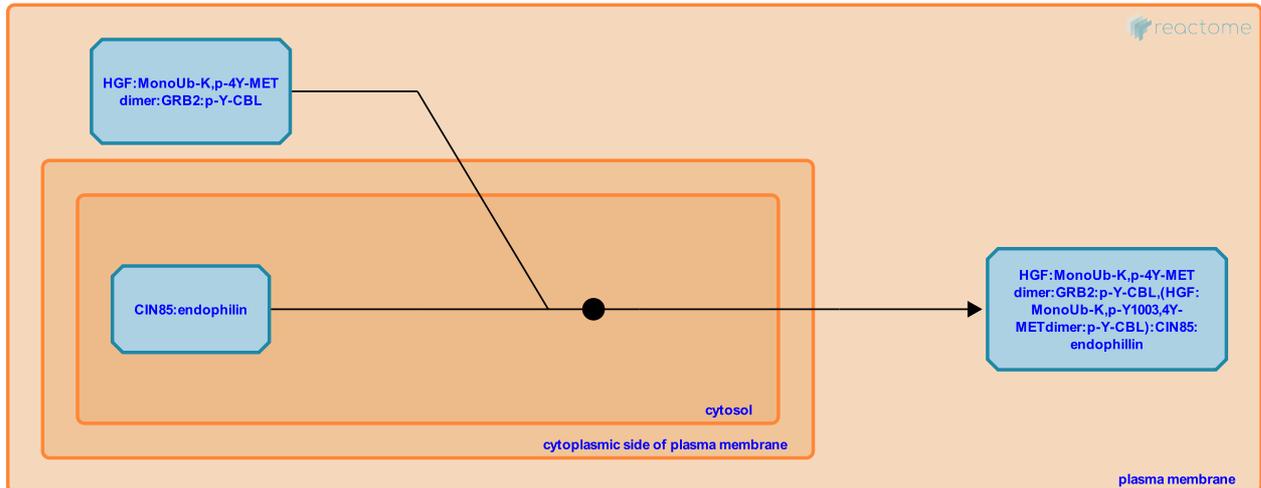
Location: [Negative regulation of MET activity](#)

Stable identifier: R-CFA-8875482

Type: binding

Compartments: cytosol, plasma membrane

Inferred from: [CBL recruits CIN85:endophilin to ubiquitinated MET \(Homo sapiens\)](#)



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: [CBL monoubiquitinates activated MET](#)

Followed by: [EPS15 and HGS bind ubiquitinated MET](#)

EPS15 and HGS bind ubiquitinated MET ↗

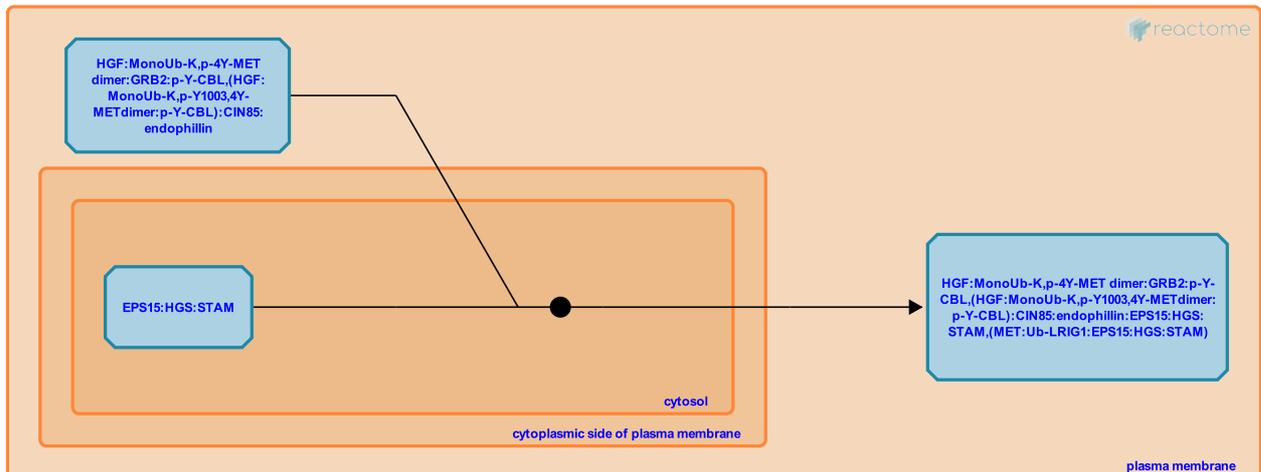
Location: [Negative regulation of MET activity](#)

Stable identifier: R-CFA-8875490

Type: binding

Compartments: cytosol, plasma membrane

Inferred from: [EPS15 and HGS bind ubiquitinated MET \(Homo sapiens\)](#)



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: [CBL recruits CIN85:endophilin to ubiquitinated MET](#)

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