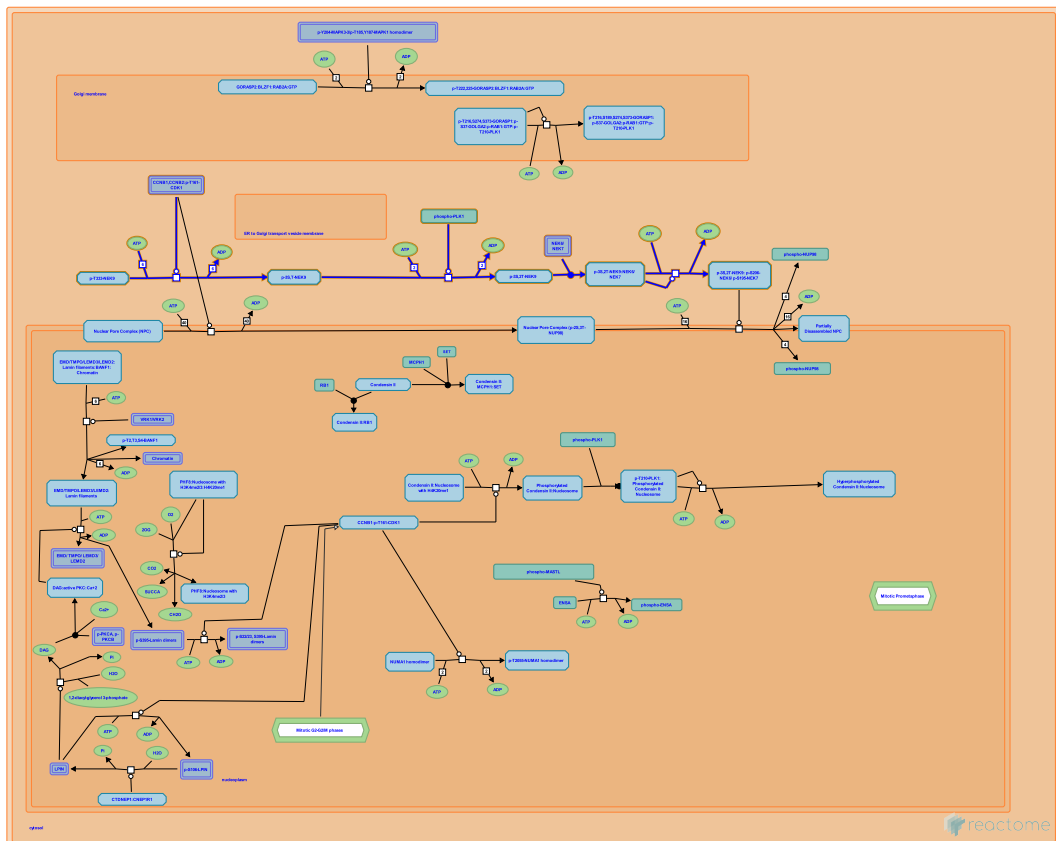


Activation of NIMA Kinases NEK9, NEK6, NEK7



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

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This is just an excerpt of a full-length report for this pathway. To access the complete report, please download it at the [Reactome Textbook](https://reactome.org/textbook).

04/05/2024

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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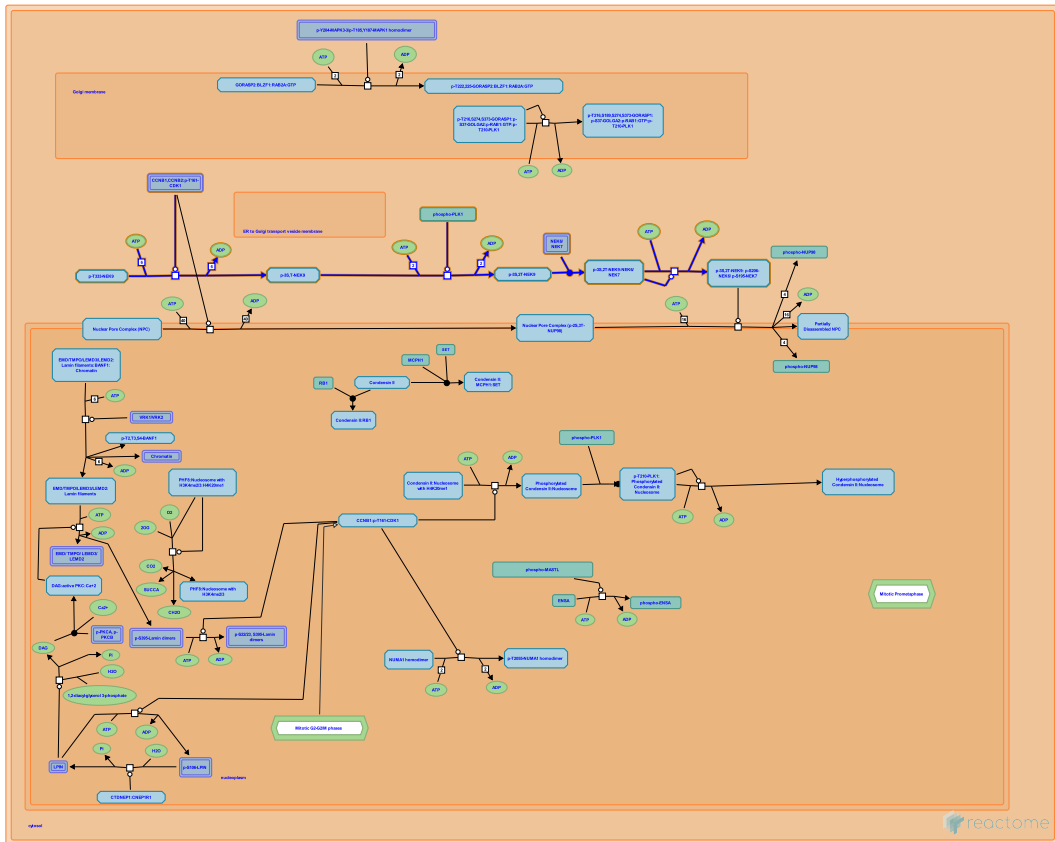
Reactome database release: 88

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

Activation of NIMA Kinases NEK9, NEK6, NEK7 ↗

Stable identifier: R-CFA-2980767

Inferred from: Activation of NIMA Kinases NEK9, NEK6, NEK7 (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.](http://www.pantherdb.org/about.jsp) For details on PANTHER see also: <http://www.pantherdb.org/about.jsp>

CDK1:CCNB phosphorylates NEK9 ↗

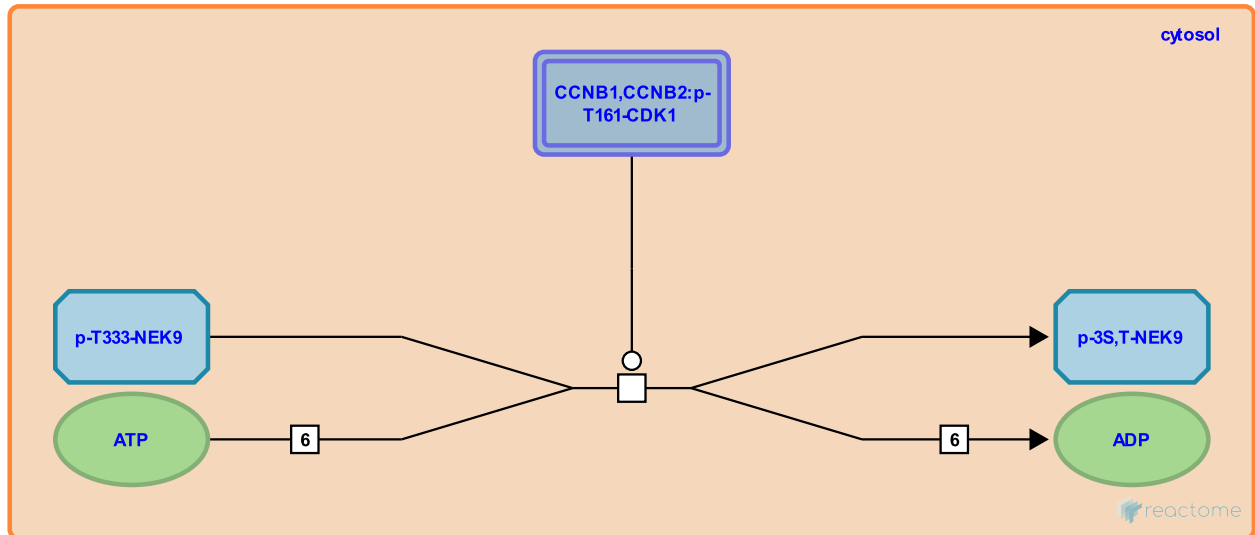
Location: [Activation of NIMA Kinases NEK9, NEK6, NEK7](#)

Stable identifier: R-CFA-2984220

Type: transition

Compartments: cytosol

Inferred from: [CDK1:CCNB phosphorylates NEK9 \(Homo sapiens\)](#)



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.](#) For details on PANTHER see also: <http://www.pantherdb.org/about.jsp>

Followed by: [PLK1 phosphorylates NEK9](#)

PLK1 phosphorylates NEK9 ↗

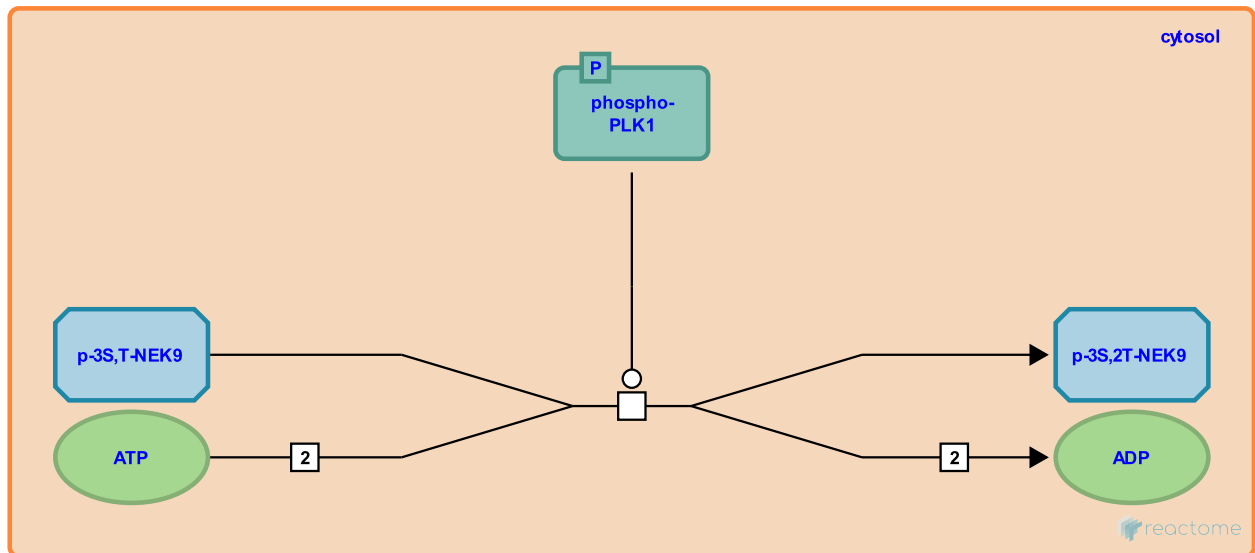
Location: [Activation of NIMA Kinases NEK9, NEK6, NEK7](#)

Stable identifier: R-CFA-2984226

Type: transition

Compartments: cytosol

Inferred from: [PLK1 phosphorylates NEK9 \(Homo sapiens\)](#)



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.](#) For details on PANTHER see also: <http://www.pantherdb.org/about.jsp>

Preceded by: [CDK1:CCNB phosphorylates NEK9](#)

Followed by: [NEK9 binds NEK6/NEK7 in the cytosol](#)

NEK9 binds NEK6/NEK7 in the cytosol ↗

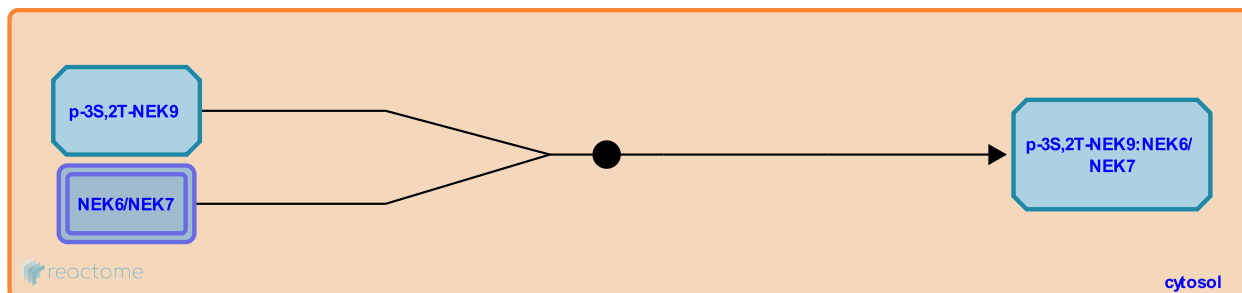
Location: Activation of NIMA Kinases NEK9, NEK6, NEK7

Stable identifier: R-CFA-2980720

Type: binding

Compartments: cytosol

Inferred from: NEK9 binds NEK6/NEK7 in the cytosol (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.](http://www.pantherdb.org/about.jsp) For details on PANTHER see also: <http://www.pantherdb.org/about.jsp>

Preceded by: PLK1 phosphorylates NEK9

Followed by: NEK9 phosphorylates NEK6/NEK7

NEK9 phosphorylates NEK6/NEK7 ↗

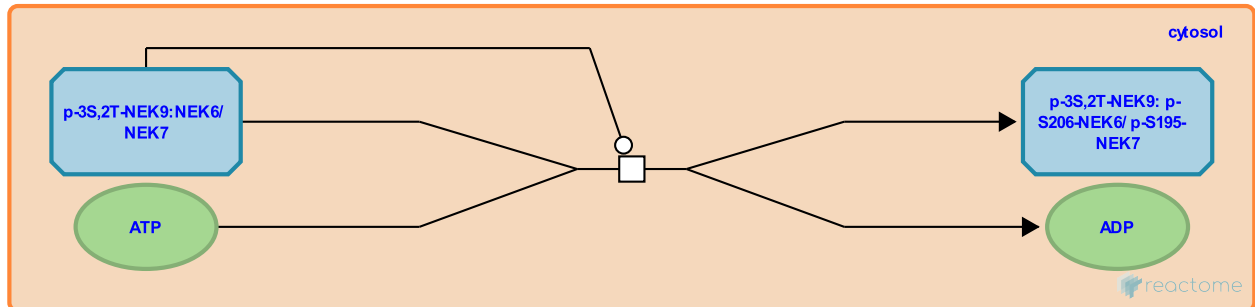
Location: [Activation of NIMA Kinases NEK9, NEK6, NEK7](#)

Stable identifier: R-CFA-2984258

Type: transition

Compartments: cytosol

Inferred from: [NEK9 phosphorylates NEK6/NEK7 \(Homo sapiens\)](#)



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.](#) For details on PANTHER see also: <http://www.pantherdb.org/about.jsp>

Preceded by: [NEK9 binds NEK6/NEK7 in the cytosol](#)

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