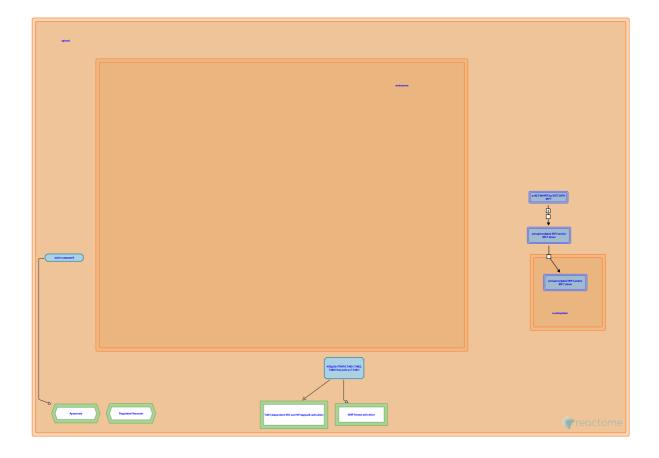


Toll Like Receptor 3 (TLR3) Cascade



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

Fabregat, A., Sidiropoulos, K., Viteri, G., Forner, O., Marin-Garcia, P., Arnau, V. et al. (2017). Reactome pathway analysis: a high-performance in-memory approach. *BMC bioinformatics*, 18, 142.

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Fabregat, A., Korninger, F., Viteri, G., Sidiropoulos, K., Marin-Garcia, P., Ping, P. et al. (2018). Reactome graph database: Efficient access to complex pathway data. *PLoS computational biology, 14*, e1005968.

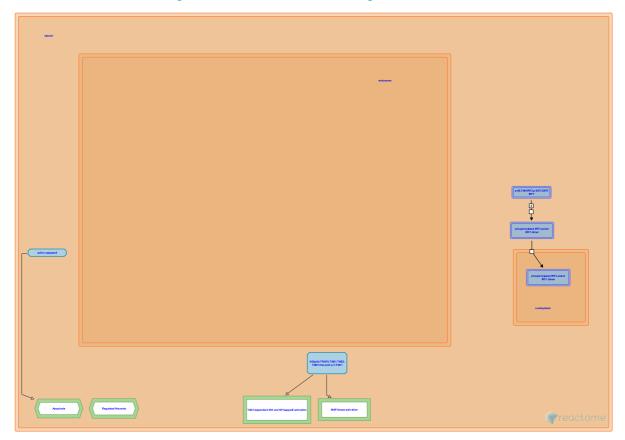
Reactome database release: 81

This document contains 4 pathways (see Table of Contents)

Toll Like Receptor 3 (TLR3) Cascade **ブ**

Stable identifier: R-XTR-168164

Inferred from: Toll Like Receptor 3 (TLR3) Cascade (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.

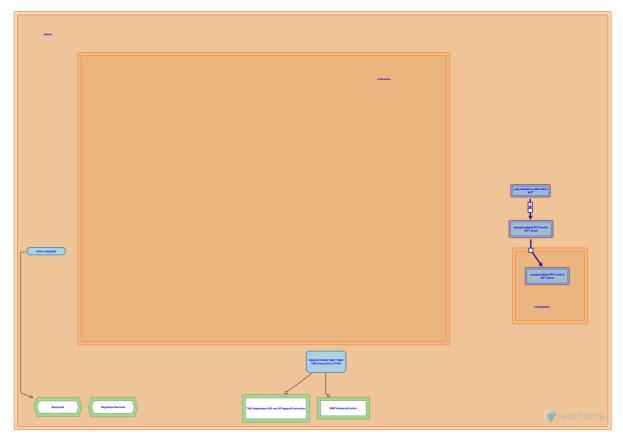
TICAM1-dependent activation of IRF3/IRF7 **对**

Location: Toll Like Receptor 3 (TLR3) Cascade

Stable identifier: R-XTR-9013973

Compartments: cytosol

Inferred from: TICAM1-dependent activation of IRF3/IRF7 (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.

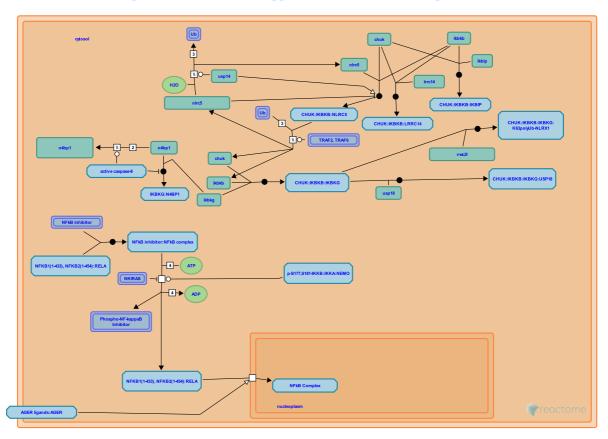
TAK1-dependent IKK and NF-kappa-B activation 7

Location: Toll Like Receptor 3 (TLR3) Cascade

Stable identifier: R-XTR-445989

Compartments: nucleoplasm, cytosol

Inferred from: TAK1-dependent IKK and NF-kappa-B activation (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.

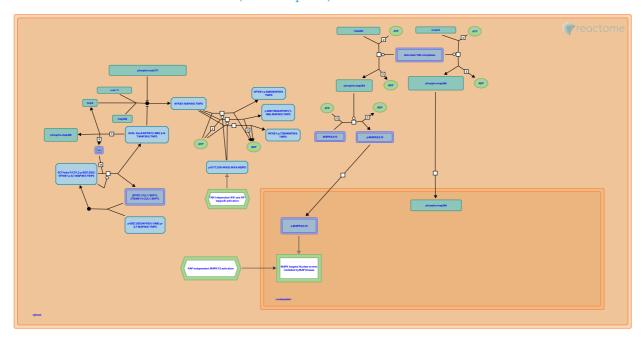
MAP kinase activation **↗**

Location: Toll Like Receptor 3 (TLR3) Cascade

Stable identifier: R-XTR-450294

Compartments: nucleoplasm, cytosol

Inferred from: MAP kinase activation (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.

Table of Contents

Introduction	1
Toll Like Receptor 3 (TLR3) Cascade	2
TICAM1-dependent activation of IRF3/IRF7	3
TAK1-dependent IKK and NF-kappa-B activation	4
MAP kinase activation	5
Table of Contents	6