

# TRAF3:NIK binds TRAF2:cIAP1/2

Garapati, P V., Rajput, A., Virgen-Slane, R., Ware, CF.

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

The contents of this document may be freely copied and distributed in any media, provided the authors, plus the institutions, are credited, as stated under the terms of [Creative Commons Attribution 4.0 International \(CC BY 4.0\) License](https://creativecommons.org/licenses/by/4.0/). For more information see our [license](https://reactome.org/licenses/).

04/11/2024

## 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

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. [↗](#)

Sidiropoulos, K., Viteri, G., Sevilla, C., Jupe, S., Webber, M., Orlic-Milacic, M. et al. (2017). Reactome enhanced pathway visualization. *Bioinformatics*, 33, 3461-3467. [↗](#)

Fabregat, A., Jupe, S., Matthews, L., Sidiropoulos, K., Gillespie, M., Garapati, P. et al. (2018). The Reactome Pathway Knowledgebase. *Nucleic Acids Res*, 46, D649-D655. [↗](#)

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: 90

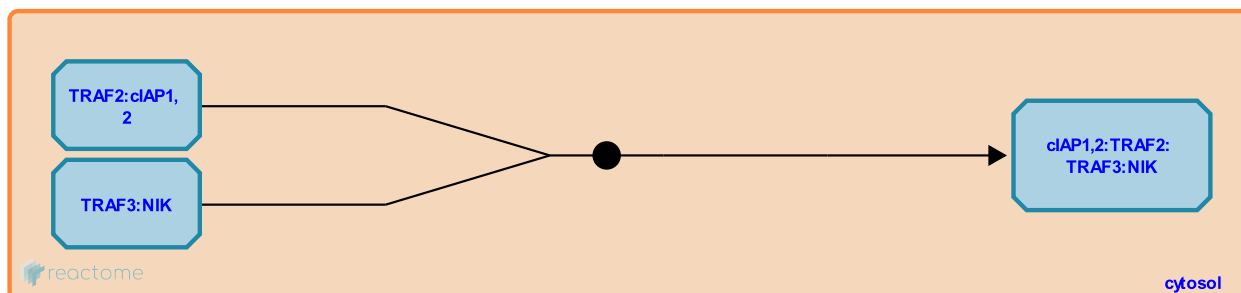
This document contains 1 reaction ([see Table of Contents](#))

## TRAF3:NIK binds TRAF2:cIAP1/2 ↗

**Stable identifier:** R-HSA-5668543

**Type:** binding

**Compartments:** cytosol



Mitogen-activated protein kinase kinase kinase 14 (MAP3K14 also named as NIK) is a central signalling component of the non-canonical pathway which integrates signals from TNFR2 and activates I $\kappa$ B kinase-alpha (I $\kappa$ BA) for triggering p100 phosphorylation and processing (Sun 2011). A tight control of NIK stability is essential to achieve controlled activation of the noncanonical NF- $\kappa$ B signalling upon TNFR2 activation. In unstimulated cells the level of NIK protein is extremely low, which is due to constant degradation by a ubiquitination-dependent mechanism (Liao et al. 2004). Proteasomal degradation of NIK occurs on assembly of a regulatory complex through TRAF3 complexed with NIK and TRAF2 which exists in a preassembled complex with cellular Inhibitor of apoptosis 1 (cIAP1) and cIAP2 (cIAP1,2:TRAF2::TRAF3:NIK). The c-IAPs do not directly contact TNFR2, but rather associate with TRAF2 through their N-terminal BIR motif-comprising domain (Rothe et al. 1995, Shu et al. 1996). TRAF3 functions as a bridging factor between cIAP1/2-TRAF2 E3 complex and NIK enabling cIAP to mediate K48 linked ubiquitination of NIK (Zarnegar et al. 2008, Vallabhapurapu et al. 2008, Li et al. 2004).

### Literature references

- Mahoney, DJ., Wang, Y., Shiba, T., Mak, TW., Yeh, WC., Cheng, G. et al. (2008). Noncanonical NF-kappaB activation requires coordinated assembly of a regulatory complex of the adaptors cIAP1, cIAP2, TRAF2 and TRAF3 and the kinase NIK. *Nat. Immunol.*, 9, 1371-8. ↗
- Zhang, W., Tseng, PH., Matsuzawa, A., Vignali, DA., Bergsagel, PL., Keats, JJ. et al. (2008). Nonredundant and complementary functions of TRAF2 and TRAF3 in a ubiquitination cascade that activates NIK-dependent alternative NF-kappaB signaling. *Nat. Immunol.*, 9, 1364-70. ↗

### Editions

2015-01-26	Authored, Edited	Garapati, P V.
2015-05-12	Reviewed	Ware, CF., Rajput, A., Virgen-Slane, R.