

BCL10 oligomerizes

Garapati, P V., Geijtenbeek, TB.

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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- Sidiropoulos, K., Viteri, G., Sevilla, C., Jupe, S., Webber, M., Orlic-Milacic, M. et al. (2017). Reactome enhanced pathway visualization. *Bioinformatics*, 33, 3461-3467. [↗](#)
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Reactome database release: 88

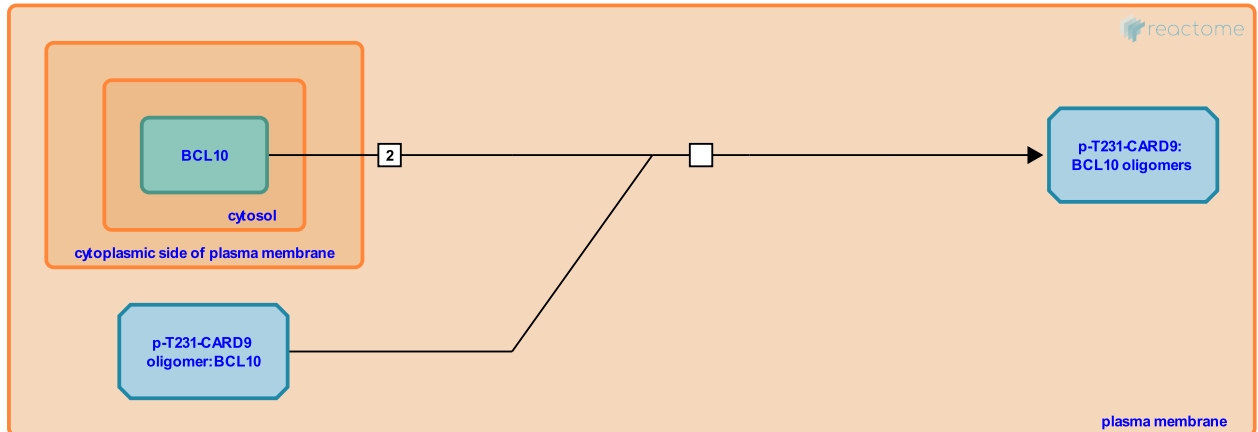
This document contains 1 reaction ([see Table of Contents](#))

BCL10 oligomerizes [↗](#)

Stable identifier: R-HSA-5607737

Type: transition

Compartments: plasma membrane, cytosol



CARD interactions between CARD9 and BCL10 induce BCL10 oligomerization (through its CARD domain), required for oligomerization and activation of MALT1.

Literature references

Nunez, G., Merino, J., Inohara, N., Hottiger, MO., Carrio, R., Chen, S. et al. (1999). CIPER, a novel NF kappaB-activating protein containing a caspase recruitment domain with homology to Herpesvirus-2 protein E10. *J. Biol. Chem.*, 274, 9955-61. [↗](#)

Editions

2014-07-14	Authored, Edited	Garapati, P V.
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