

# L1-EGFR trans-heterodimerization

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27/09/2020

## 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: 73

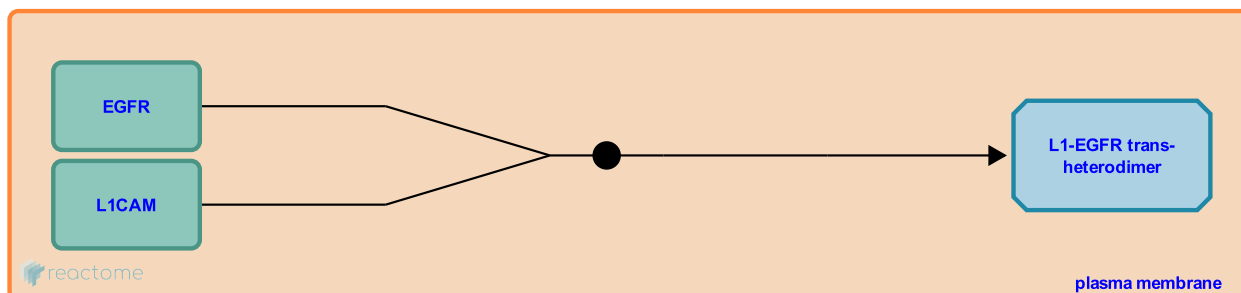
This document contains 1 reaction ([see Table of Contents](#))

## L1-EGFR trans-heterodimerization [↗](#)

**Stable identifier:** R-HSA-445069

**Type:** binding

**Compartments:** plasma membrane



L1CAM and EGFR engage in a weak heterophilic trans interaction and this induces EGFR tyrosine kinase activity and its activation. However, this trans interaction alone is not sufficient to induce EGFR auto-phosphorylation, which requires additional cis type interactions between the two proteins.

### Literature references

Islam, R., Kristiansen, L.V., Romani, S., Garcia-Alonso, L., Hortsch, M. (2004). Activation of EGF receptor kinase by L1-mediated homophilic cell interactions. *Mol Biol Cell*, 15, 2003-12. [↗](#)

### Editions

2008-07-30	Authored, Edited	Garapati, P V.
2010-02-16	Reviewed	Maness, PF.