

Recruitment of Profilin by Ena/VASP proteins

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https://reactome.org

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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- 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 data-base: Efficient access to complex pathway data. *PLoS computational biology, 14*, e1005968.

Reactome database release: 88

This document contains 1 reaction (see Table of Contents)

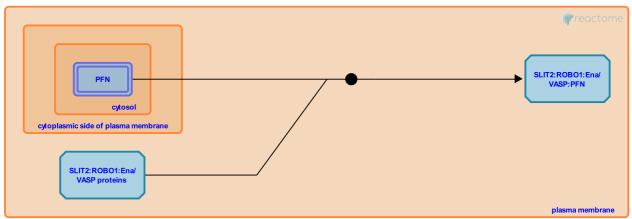
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Recruitment of Profilin by Ena/VASP proteins 对

Stable identifier: R-HSA-428534

Type: binding

Compartments: cytosol, plasma membrane



Ena/VASP proteins (ENAH, EVL1 and VASP) enhance actin filament elongation via the recruitment of profilin:actin complexes to the tips of spreading lamellipodia. Profilin (PFN1 or PFN2) binds to the central proline rich domain of an Ena/VASP protein (Bashaw et al. 2000).

Literature references

Pawson, T., Bashaw, GJ., Goodman, CS., Kidd, T., Murray, D. (2000). Repulsive axon guidance: Abelson and Enabled play opposing roles downstream of the roundabout receptor. *Cell*, 101, 703-15.

Editions

2008-09-05	Authored, Edited	Garapati, P V.
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2017-06-26	Edited	Orlic-Milacic, M.
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