

Integrin alpha IIb beta3 T779 phosphorylation blocks SHC binding

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16/05/2024

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.

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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- 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)

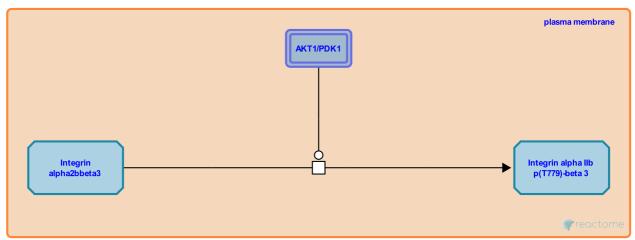
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Stable identifier: R-HSA-432110

Type: transition

Compartments: plasma membrane



The binding of SHC to integrin alpha IIb beta 3 is blocked by phosphorylation of beta 3 at Thr-779, or by substitution of this residue for Asp. PDK1 and Akt1/PKB-alpha both specifically target Thr-779 in in vitro assays.

Literature references

Kirk, RI., Lerea, KM., Sanderson, MR. (2000). Threonine phosphorylation of the beta 3 integrin cytoplasmic tail, at a site recognized by PDK1 and Akt/PKB in vitro, regulates Shc binding. *J Biol Chem*, 275, 30901-6. *¬*

Editions

2009-09-04	Authored	Akkerman, JW.
2010-09-01	Reviewed	Heemskerk, JW.
2010-09-01	Edited	Jupe, S.