

p-SYK and LYN phosphorylate BTK

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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](#))

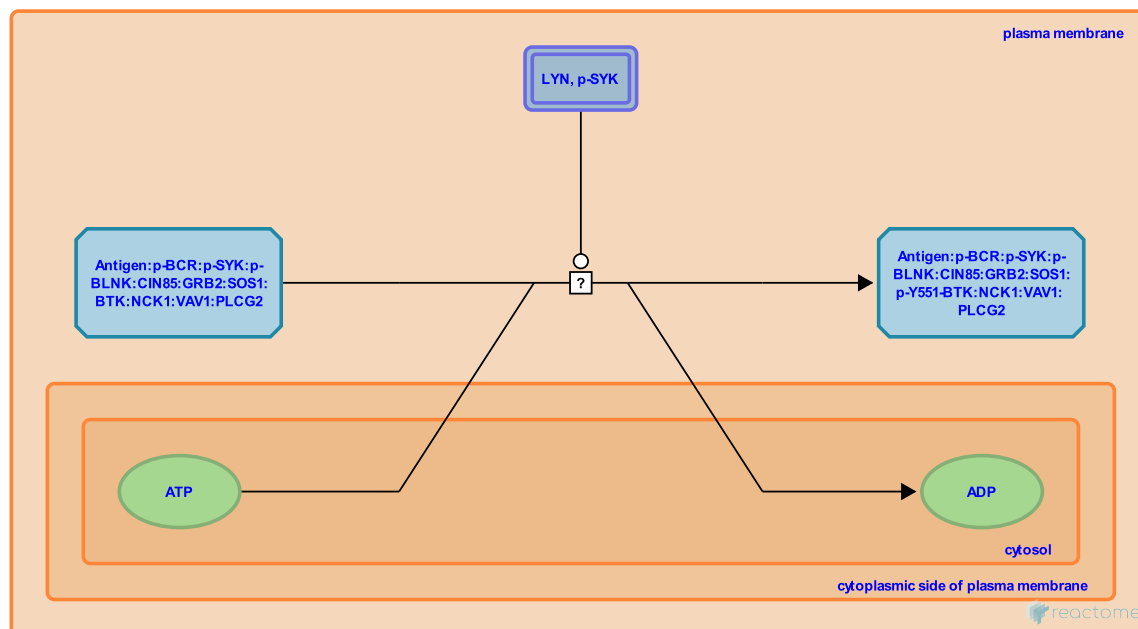
p-SYK and LYN phosphorylate BTK ↗

Stable identifier: R-HSA-9606163

Type: uncertain

Compartments: plasma membrane

Inferred from: LYN,SYK phosphorylate BTK (Gallus gallus), Lyn, p-Syk phosphorylate Btk (Mus musculus)



LYN and activated (phosphorylated) SYK phosphorylate BTK (Baba et al. 2001, Lin et al. 2009, also inferred from chicken homologs and mouse homologs) after BTK is recruited to the plasma membrane by phosphorylated BLNK (Baba et al. 2001) and phosphoinositol 3,4,5-trisphosphate (PIP3) (Salim et al. 1996). Phosphorylation of tyrosine-551 occurs within 30 seconds of B cell receptor activation and returns to low phosphorylation after 30 minutes (Nisitani et al. 1999).

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Editions

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