

RUNX1 binds the PRKCB gene promoter

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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: 74

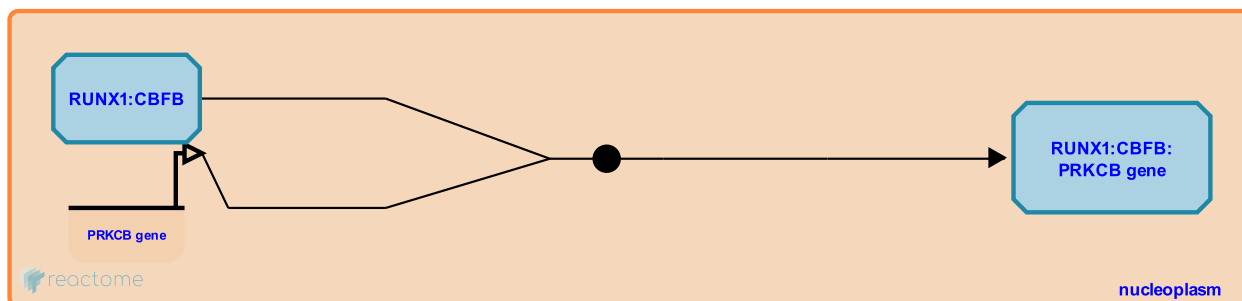
This document contains 1 reaction ([see Table of Contents](#))

RUNX1 binds the PRKCB gene promoter ↗

Stable identifier: R-HSA-8939054

Type: binding

Compartments: nucleoplasm



The RUNX1:CBFB complex binds the promoter of the PRKCB gene, encoding protein kinase C beta (Hug et al. 2004).

Literature references

Hug, BA., Ahmed, N., Robbins, JA., Lazar, MA. (2004). A chromatin immunoprecipitation screen reveals protein kinase Cbeta as a direct RUNX1 target gene. *J. Biol. Chem.*, 279, 825-30. ↗

Editions

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