

CEBPB and phospho-STAT3 bind the promoter of the MYC gene

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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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Reactome database release: 88

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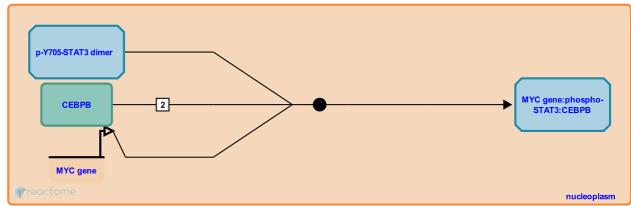
CEBPB and phospho-STAT3 bind the promoter of the MYC gene 7

Stable identifier: R-HSA-9618584

Type: binding

Compartments: nucleoplasm

Inferred from: Cebpb and phospho-Stat3 bind the promoter of the Myc gene (Mus musculus)



Activated (phosphorylated) STAT3 activates transcription of CEBPB and both phospho-STAT3 and CEBPB bind the promoter of the MYC gene (inferred from mouse homologs). The expression of MYC enhances proliferation of myeloid progenitors during emergency granulopoiesis in response to bacterial infection.

Editions

2018-09-10	Authored, Edited	May, B.
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