

FANCM binds FAAP24

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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

This document contains 1 reaction (see Table of Contents)

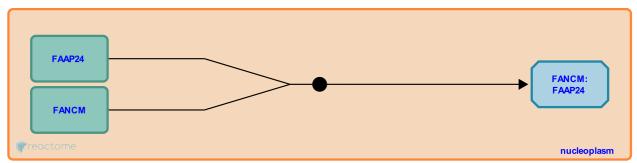
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FANCM binds FAAP24

Stable identifier: R-HSA-6785607

Type: binding

Compartments: nucleoplasm



FANCM binds FAAP24, forming a complex that recognizes DNA interstrand crosslinks, thus triggering the Fanconi anemia repair pathway (Ciccia et al. 2007, Kim et al. 2008).

Literature references

Gurtan, A., D'Andrea, AD., Kim, JM., Kee, Y. (2008). Cell cycle-dependent chromatin loading of the Fanconi anemia core complex by FANCM/FAAP24. *Blood, 111*, 5215-22.

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Editions

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