Association of RAD52 with the RPA complex at resected DNA DSBs

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**Introduction**

Reactome is an 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**


Reactome database release: 72

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