

E2F7 binds E2F8

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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

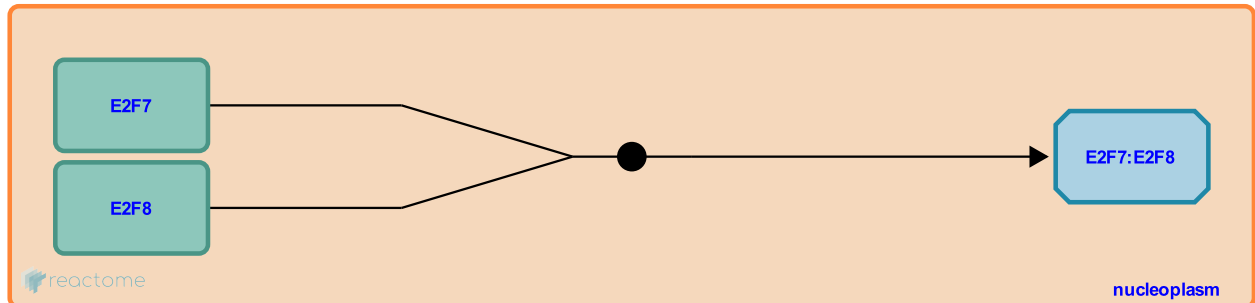
This document contains 1 reaction ([see Table of Contents](#))

E2F7 binds E2F8 [↗](#)

Stable identifier: R-HSA-8953013

Type: binding

Compartments: nucleoplasm



E2F7 forms heterodimers with E2F8 (Li et al. 2008, Zalmas et al. 2008).

Literature references

Coutts, AS., Reilly, C., Zalmas, LP., La Thangue, NB., Zhao, X., Fisher, R. et al. (2008). DNA-damage response control of E2F7 and E2F8. *EMBO Rep.*, 9, 252-9. [↗](#)

Siddiqui, H., Ran, C., Li, J., de Bruin, A., Chen, HZ., Li, E. et al. (2008). Synergistic function of E2F7 and E2F8 is essential for cell survival and embryonic development. *Dev. Cell*, 14, 62-75. [↗](#)

Editions

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