TFAP2 (AP-2) family regulates transcription of cell cycle factors

Bogachek, MV., Dawid, IB., Orlic-Milacic, M., Weigel, RJ., Zarelli, VE.

European Bioinformatics Institute, New York University Langone Medical Center, Ontario Institute for Cancer Research, Oregon Health and Science University.

The contents of this document may be freely copied and distributed in any media, provided the authors, plus the institutions, are credited, as stated under the terms of Creative Commons Attribution 4.0 International (CC BY 4.0) License. For more information see our license.

19/12/2019
**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.

The development of Reactome is supported by grants from the US National Institutes of Health (P41 HG003751), University of Toronto (CFREF Medicine by Design), European Union (EU STRP, EMI-CD), and the European Molecular Biology Laboratory (EBI Industry program).

**Literature references**


Reactome database release: 71

This document contains 1 pathway and 4 reactions (see Table of Contents)
TFAP2 (AP-2) family regulates transcription of cell cycle factors

Stable identifier: R-HSA-8866911


Literature references

Williams, CM., Scibetta, AG., Friedrich, JK., Canosa, M., Berlato, C., Moss, CH. et al. (2009). AP-2gamma promotes proliferation in breast tumour cells by direct repression of the CDKN1A gene. EMBO J., 28, 3591-601.


Editions

<table>
<thead>
<tr>
<th>Date</th>
<th>Event</th>
<th>Author(s)</th>
</tr>
</thead>
<tbody>
<tr>
<td>2016-03-14</td>
<td>Authored, Edited</td>
<td>Orlic-Milacic, M.</td>
</tr>
<tr>
<td>2016-05-04</td>
<td>Reviewed</td>
<td>Dawid, IB., Zarelli, VE.</td>
</tr>
<tr>
<td>2016-05-17</td>
<td>Reviewed</td>
<td>Weigel, RJ., Bogachek, MV.</td>
</tr>
</tbody>
</table>
TFAP2A homodimer binds CDKN1A (p21) gene

Location: TFAP2 (AP-2) family regulates transcription of cell cycle factors

Stable identifier: R-HSA-8865244

Type: binding

Compartments: nucleoplasm