

# EGR2 and SOX10 bind the MAG gene

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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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Reactome database release: 88

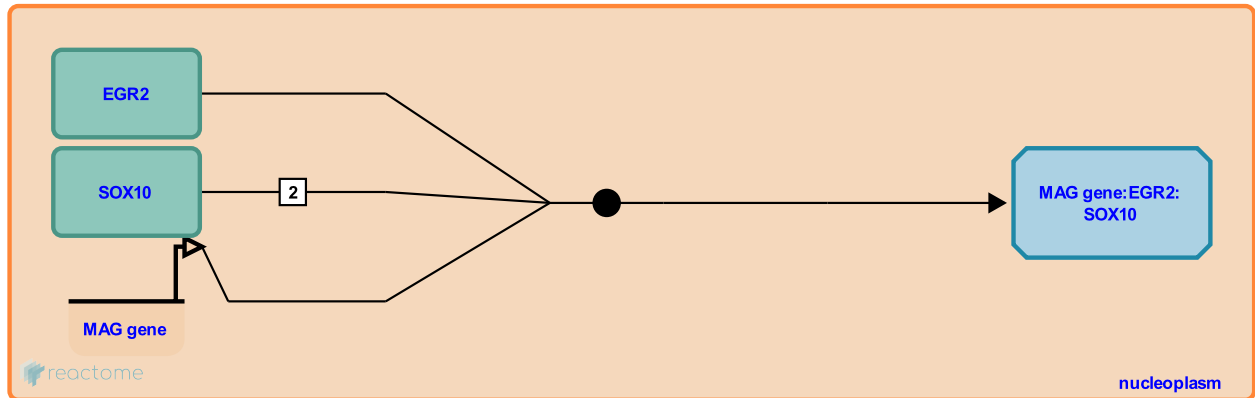
This document contains 1 reaction ([see Table of Contents](#))

## EGR2 and SOX10 bind the MAG gene ↗

**Stable identifier:** R-HSA-9616116

**Type:** binding

**Compartments:** nucleoplasm



EGR2 and SOX10 bind to elements in the MAG gene to promote expression during myelination in the peripheral and central nervous system (Jones et al, 2007; Jang et al, 2006; LeBlanc et al, 2007).

### Literature references

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### Editions

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