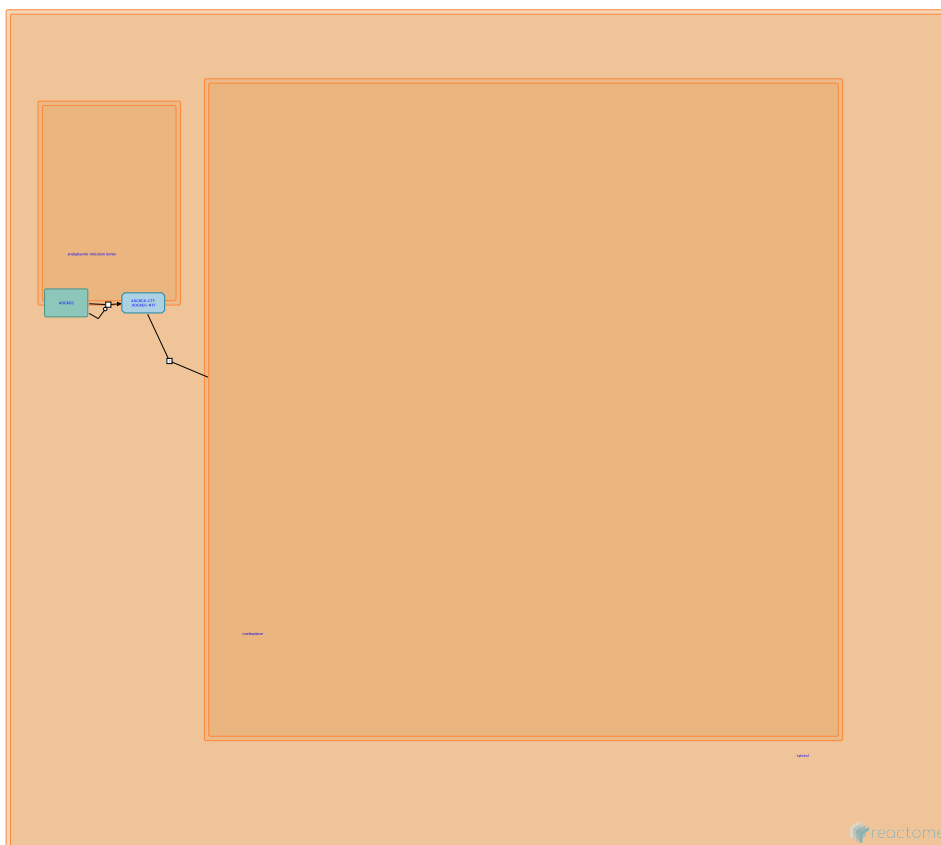


# EGR2 and SOX10-mediated initiation of Schwann cell myelination



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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.

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

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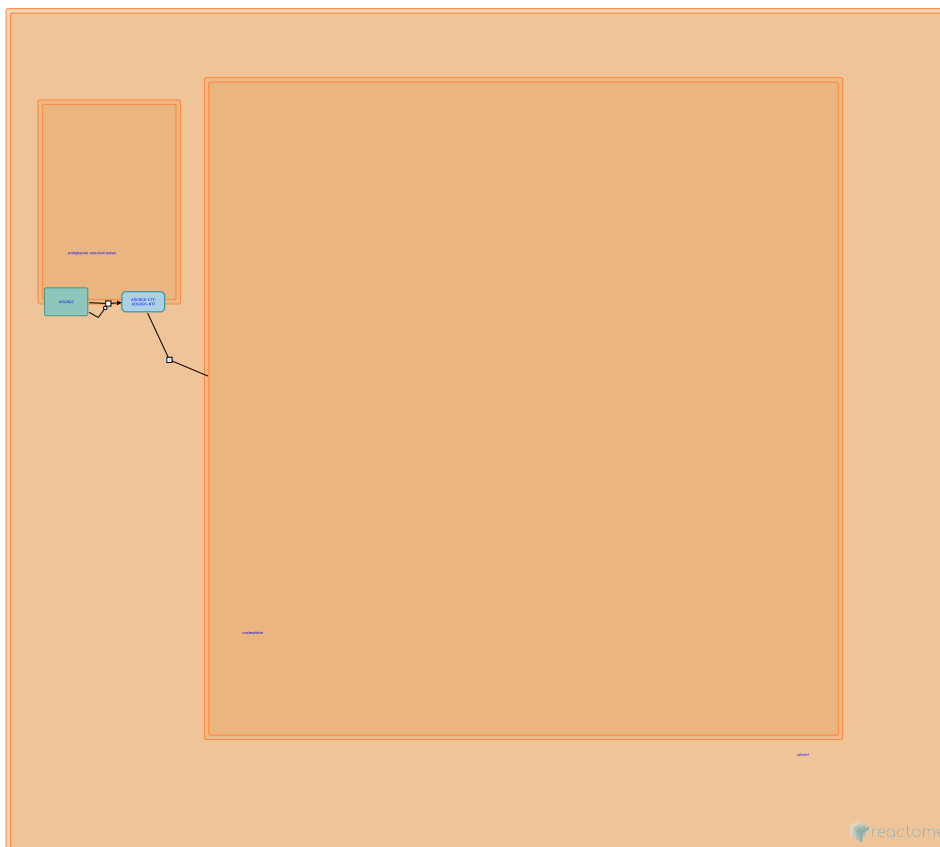
Reactome database release: 77

This document contains 1 pathway and 2 reactions ([see Table of Contents](#))

## EGR2 and SOX10-mediated initiation of Schwann cell myelination ↗

**Stable identifier:** R-SSC-9619665

**Inferred from:** [EGR2 and SOX10-mediated initiation of Schwann cell myelination \(Homo sapiens\)](#)



This event has been computationally inferred from an event that has been demonstrated in another species.

The inference is based on the homology mapping from PANTHER. Briefly, reactions for which all involved PhysicalEntities (in input, output and catalyst) have a mapped orthologue/paralogue (for complexes at least 75% of components must have a mapping) are inferred to the other species. High level events are also inferred for these events to allow for easier navigation.

[More details and caveats of the event inference in Reactome.](/electronic_inference_compara.html) For details on PANTHER see also: <http://www.pantherdb.org/about.jsp>

## Autocleavage of ADGRG6 ↗

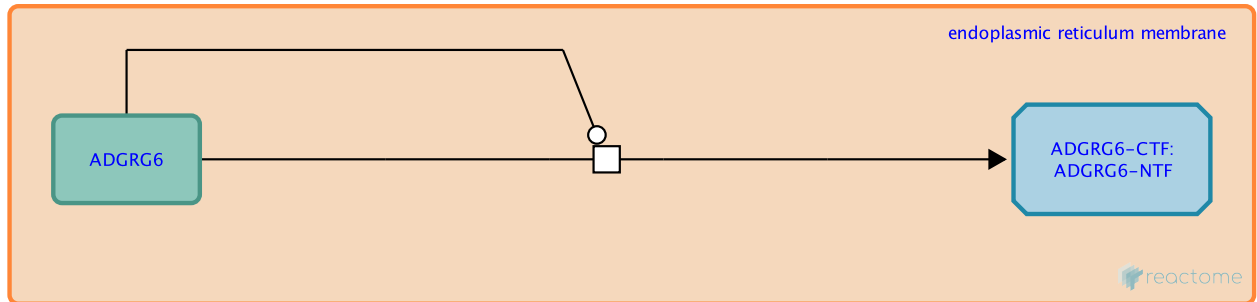
**Location:** [EGR2 and SOX10-mediated initiation of Schwann cell myelination](#)

**Stable identifier:** R-SSC-9614271

**Type:** transition

**Compartments:** endoplasmic reticulum membrane

**Inferred from:** [Autocleavage of ADGRG6 \(Homo sapiens\)](#)



This event has been computationally inferred from an event that has been demonstrated in another species.

The inference is based on the homology mapping from PANTHER. Briefly, reactions for which all involved PhysicalEntities (in input, output and catalyst) have a mapped orthologue/paralogue (for complexes at least 75% of components must have a mapping) are inferred to the other species. High level events are also inferred for these events to allow for easier navigation.

[More details and caveats of the event inference in Reactome.](/electronic_inference_compara.html) For details on PANTHER see also: <http://www.pantherdb.org/about.jsp>

**Followed by:** [ADGRG heterodimer translocates to plasma membrane](#)

## ADGRG heterodimer translocates to plasma membrane ↗

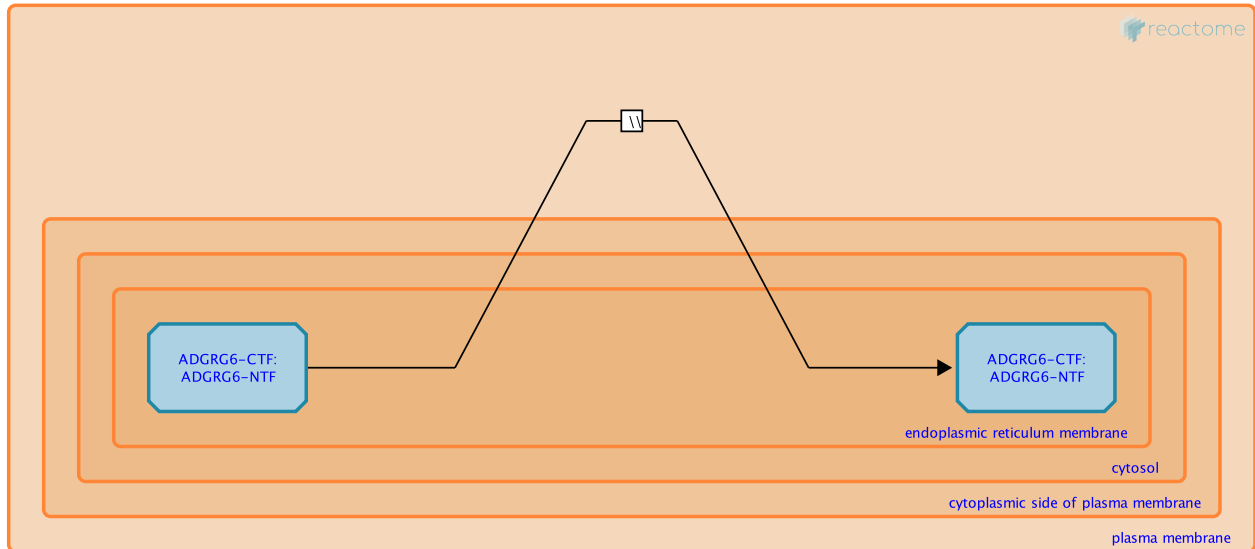
**Location:** EGR2 and SOX10-mediated initiation of Schwann cell myelination

**Stable identifier:** R-SSC-9614273

**Type:** omitted

**Compartments:** plasma membrane

**Inferred from:** ADGRG heterodimer translocates to plasma membrane (Homo sapiens)



This event has been computationally inferred from an event that has been demonstrated in another species.

The inference is based on the homology mapping from PANTHER. Briefly, reactions for which all involved PhysicalEntities (in input, output and catalyst) have a mapped orthologue/paralogue (for complexes at least 75% of components must have a mapping) are inferred to the other species. High level events are also inferred for these events to allow for easier navigation.

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**Preceded by:** [Autocleavage of ADGRG6](#)

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