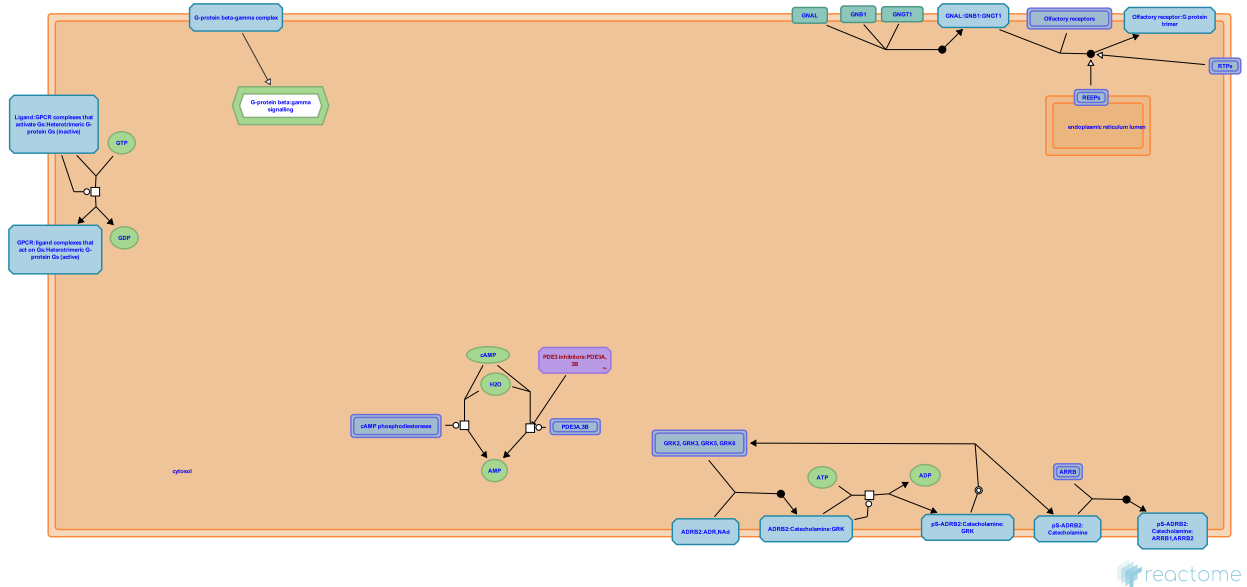


# G alpha (s) signalling events



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

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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

- Fabregat, A., Sidiropoulos, K., Viteri, G., Forner, O., Marin-Garcia, P., Arnau, V. et al. (2017). Reactome pathway analysis: a high-performance in-memory approach. *BMC bioinformatics*, 18, 142. [↗](#)
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- Fabregat, A., Jupe, S., Matthews, L., Sidiropoulos, K., Gillespie, M., Garapati, P. et al. (2018). The Reactome Pathway Knowledgebase. *Nucleic Acids Res*, 46, D649-D655. [↗](#)
- Fabregat, A., Korninger, F., Viteri, G., Sidiropoulos, K., Marin-Garcia, P., Ping, P. et al. (2018). Reactome graph database: Efficient access to complex pathway data. *PLoS computational biology*, 14, e1005968. [↗](#)

Reactome database release: 70

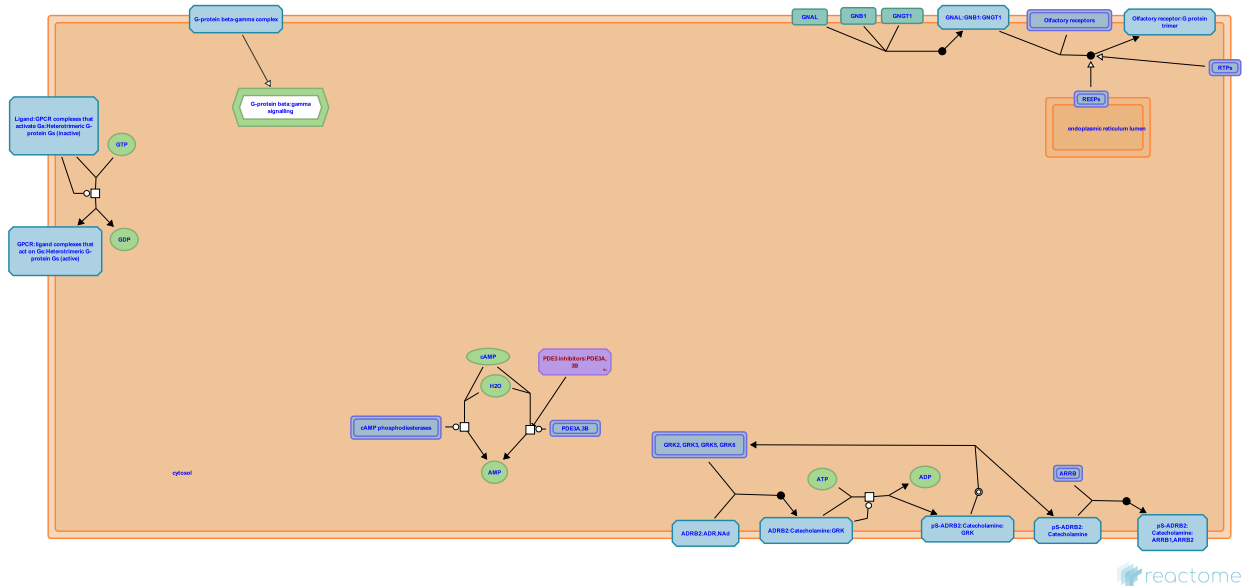
This document contains 2 pathways and 7 reactions ([see Table of Contents](#))

## G alpha (s) signalling events ↗

**Stable identifier:** R-RNO-418555

**Compartments:** plasma membrane

**Inferred from:** G alpha (s) signalling events (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>

## Liganded Gs-activating GPCR acts as a GEF for Gs ↗

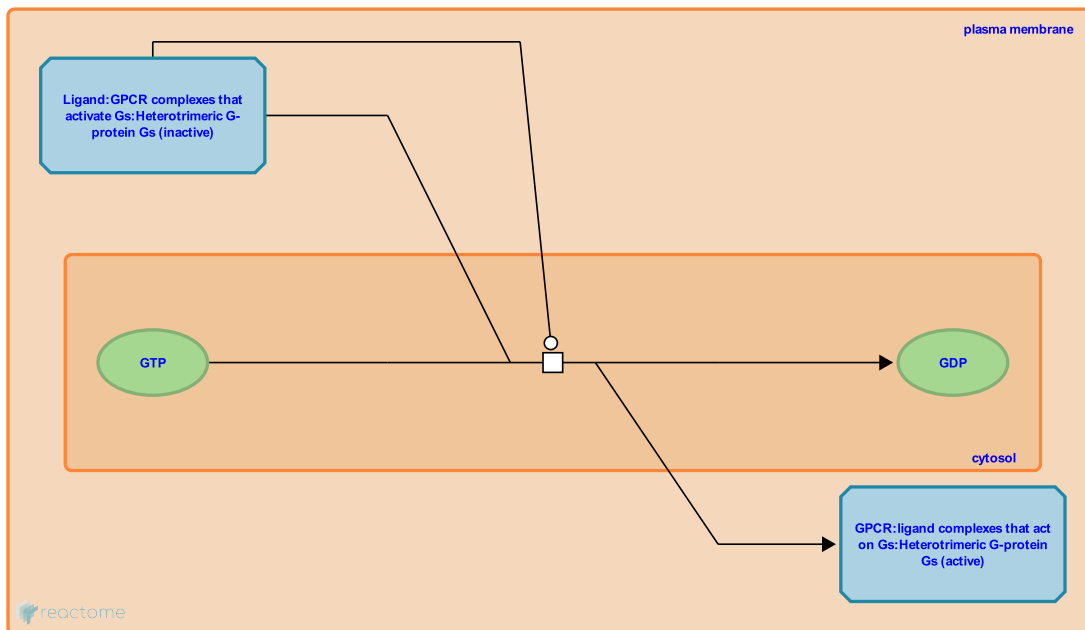
**Location:** G alpha (s) signalling events

**Stable identifier:** R-RNO-379044

**Type:** transition

**Compartments:** cytosol, plasma membrane

**Inferred from:** Liganded Gs-activating GPCR acts as a GEF for Gs (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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## cAMP degradation by Phosphodiesterases [↗](#)

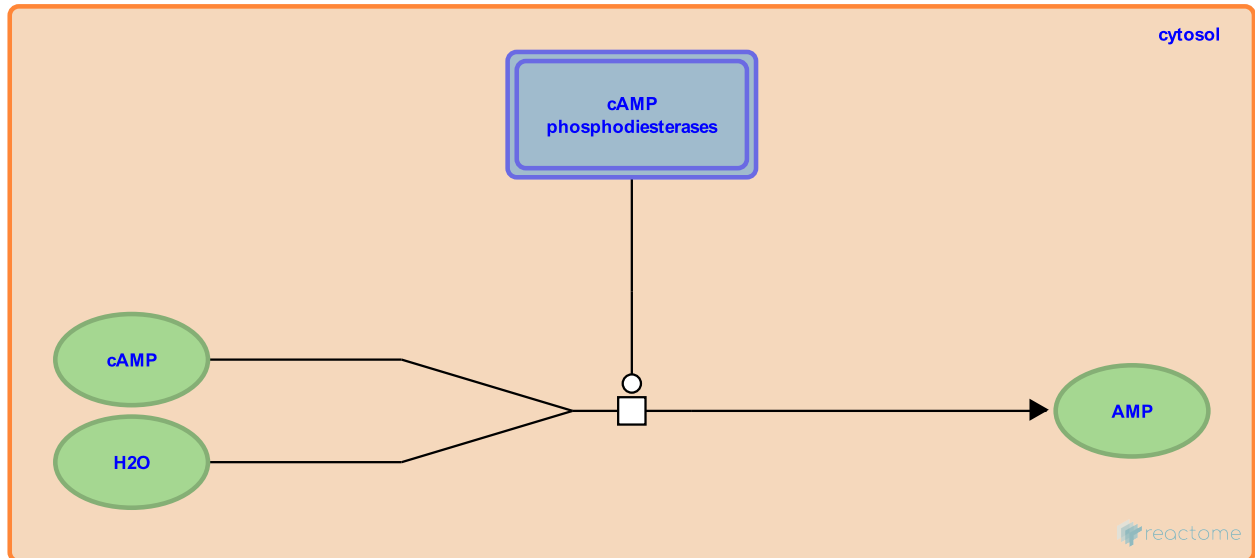
**Location:** [G alpha \(s\) signalling events](#)

**Stable identifier:** R-RNO-418553

**Type:** transition

**Compartments:** cytosol

**Inferred from:** [cAMP degradation by Phosphodiesterases \(Homo sapiens\)](#)



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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## PDE3A,3B hydrolyses cAMP to AMP ↗

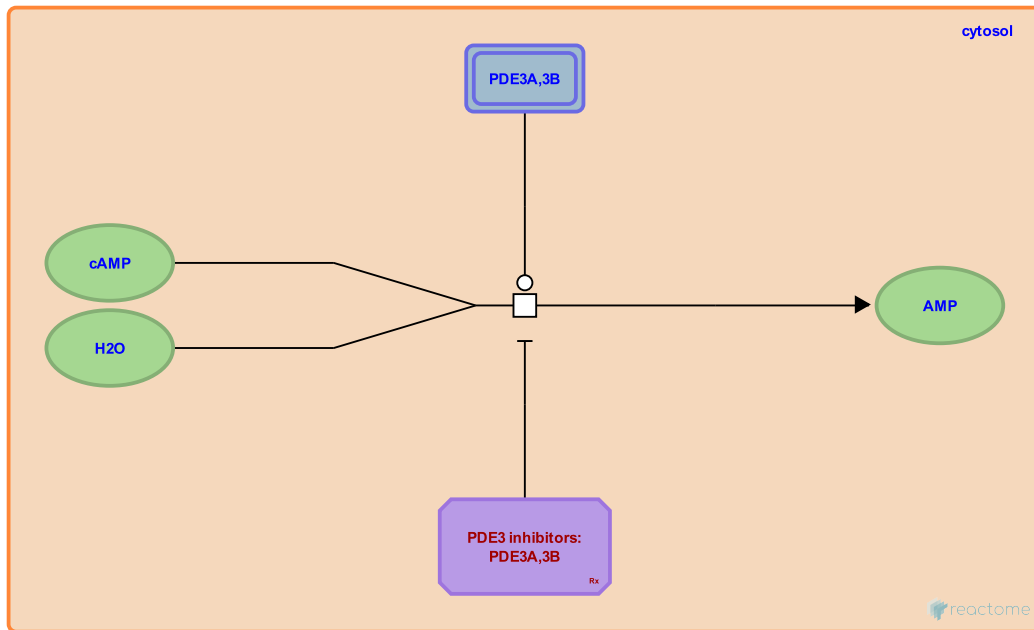
**Location:** [G alpha \(s\) signalling events](#)

**Stable identifier:** R-RNO-9629675

**Type:** transition

**Compartments:** cytosol

**Inferred from:** [PDE3A,3B hydrolyses cAMP to AMP \(Homo sapiens\)](#)



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>

## ADRB2 in ADRB2:GRK complex is phosphorylated ↗

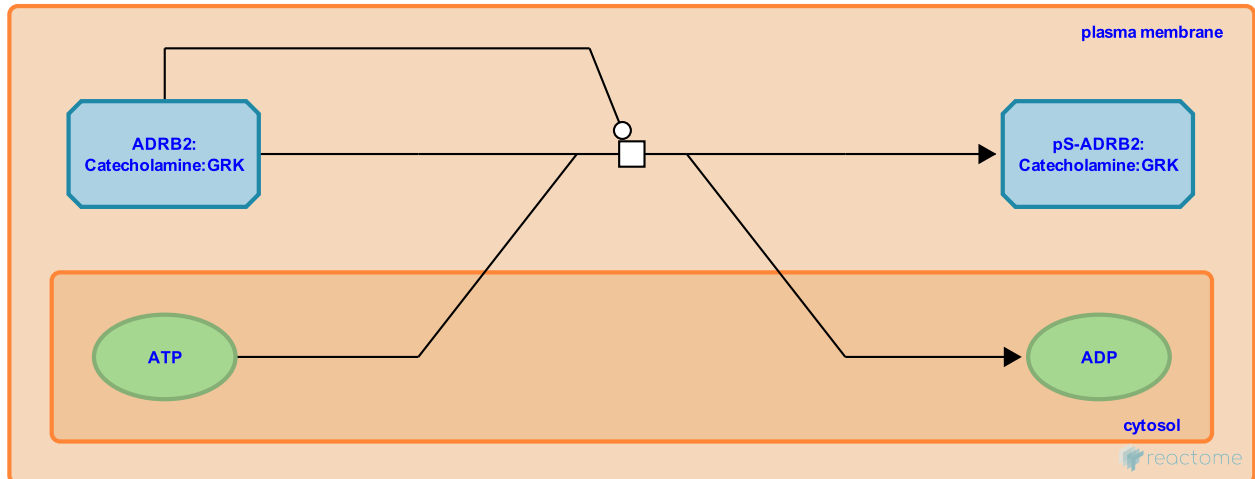
**Location:** [G alpha \(s\) signalling events](#)

**Stable identifier:** R-RNO-8851797

**Type:** transition

**Compartments:** plasma membrane, cytosol

**Inferred from:** [ADRB2 in ADRB2:GRK complex is phosphorylated \(Homo sapiens\)](#)



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>

**Preceded by:** [GRKs bind ADRB2:Catecholamine](#)

**Followed by:** [ADRB2:GRK complex dissociates to Phosphorylated ADRB2](#)

## ADRB2:Catecholamine binds ARRB1, ARRB2 ↗

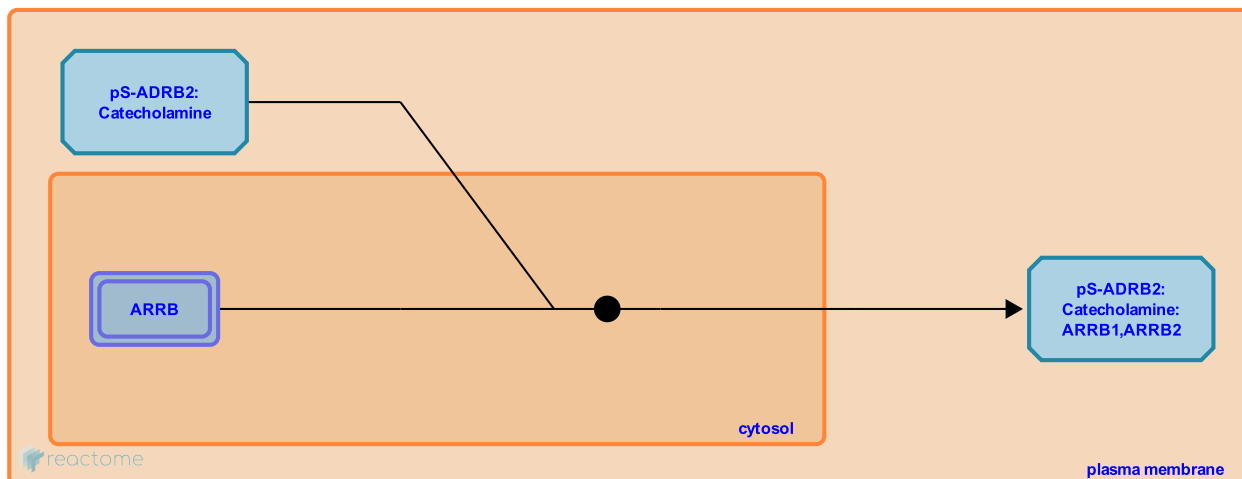
**Location:** [G alpha \(s\) signalling events](#)

**Stable identifier:** R-RNO-8852167

**Type:** binding

**Compartments:** cytosol, extracellular region, plasma membrane

**Inferred from:** [ADRB2:Catecholamine binds ARRB1, ARRB2 \(Homo sapiens\)](#)



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>

**Preceded by:** [ADRB2:GRK complex dissociates to Phosphorylated ADRB2](#)



## GRKs bind ADRB2:Catecholamine ↗

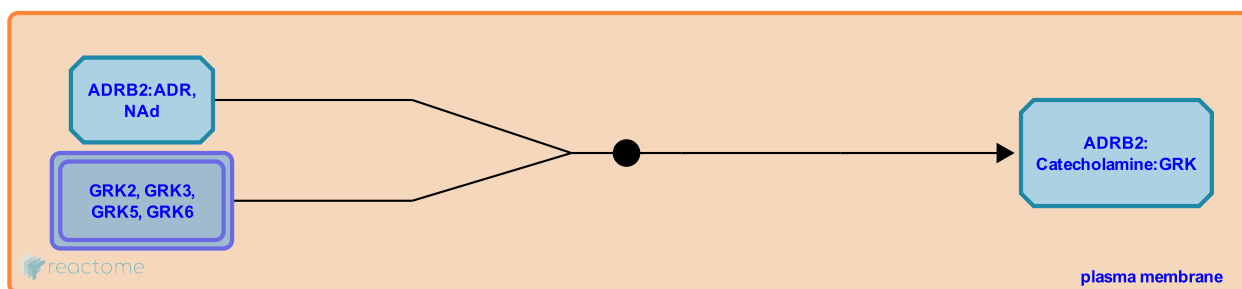
**Location:** G alpha (s) signalling events

**Stable identifier:** R-RNO-8982645

**Type:** binding

**Compartments:** plasma membrane

**Inferred from:** GRKs bind ADRB2:Catecholamine (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:** [ADRB2 in ADRB2:GRK complex is phosphorylated](#)

## ADRB2:GRK complex dissociates to Phosphorylated ADRB2 ↗

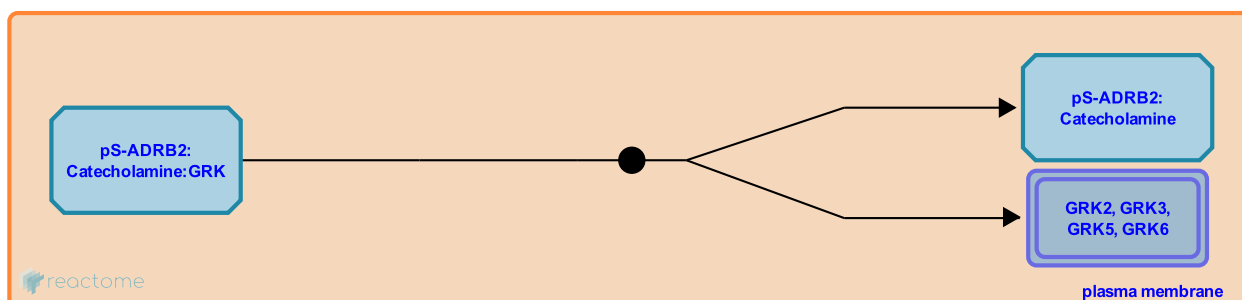
**Location:** [G alpha \(s\) signalling events](#)

**Stable identifier:** R-RNO-8982641

**Type:** dissociation

**Compartments:** plasma membrane, cytosol

**Inferred from:** [ADRB2:GRK complex dissociates to Phosphorylated ADRB2 \(Homo sapiens\)](#)



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>

**Preceded by:** [ADRB2 in ADRB2:GRK complex is phosphorylated](#)

**Followed by:** [ADRB2:Catecholamine binds ARRB1, ARRB2](#)



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