

AVP:AVPR2 Complex binds inactive heterotrimeric G(s) complex

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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

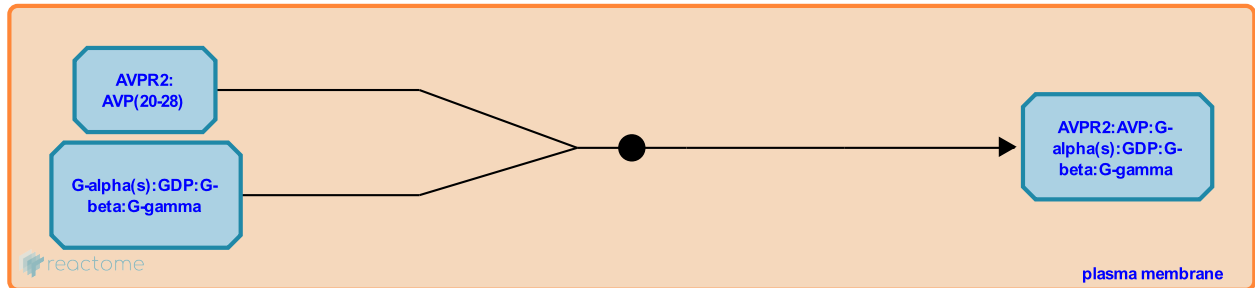
This document contains 1 reaction ([see Table of Contents](#))

AVP:AVPR2 Complex binds inactive heterotrimeric G(s) complex [↗](#)

Stable identifier: R-HSA-432188

Type: binding

Compartments: plasma membrane



The vasopressin receptor type 2 (AVPR2) interacts with G-protein alpha s via the third intracellular loop of AVPR2.

Literature references

Erlenbach, I., Wess, J. (1998). Molecular basis of V2 vasopressin receptor/Gs coupling selectivity. *J Biol Chem*, 273, 26549-58. [↗](#)

Liu, J., Wess, J. (1996). Different single receptor domains determine the distinct G protein coupling profiles of members of the vasopressin receptor family. *J Biol Chem*, 271, 8772-8. [↗](#)

Editions

2009-08-11	Authored, Edited	May, B.
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