

# pagA binds ANTXR1

D'Eustachio, P., Leppla, SH., Liu, S., Moayeri, M., Turk, BE.

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](#).

## 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. [↗](#)
- Sidiropoulos, K., Viteri, G., Sevilla, C., Jupe, S., Webber, M., Orlic-Milacic, M. et al. (2017). Reactome enhanced pathway visualization. *Bioinformatics*, 33, 3461-3467. [↗](#)
- 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: 77

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

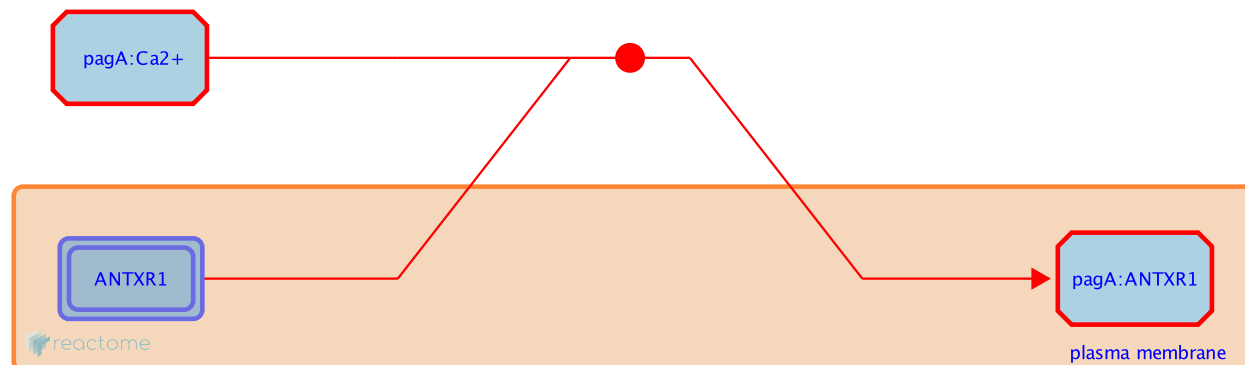
## pagA binds ANTXR1 [↗](#)

**Stable identifier:** R-HSA-5210921

**Type:** binding

**Compartments:** extracellular region, plasma membrane

**Diseases:** anthrax disease



Extracellular pagA (also known as PA83 - full length Protective Antigen - Petosa et al. 1997) produced by *Bacillus anthracis* binds to either of two isoforms of ANTXR1 (Anthrax Toxin Receptor 1, also known as TEM8 - Bradley et al. 2001; Liu and Leppla 2003) in the plasma membrane of a target human cell. The physiological ligand for ANTXR1 is not known nor are the physiological roles of the two ANTXR1 isoforms. Although ANTXR1 can act as a relatively low affinity pagA receptor in tissue culture model systems, it does not play a primary role in anthrax toxin induced effects in mouse models (Liu et al. 2009). While some studies suggest that ANTXR1 is associated with palmitoylated LRP6 (low density lipoprotein receptor related protein 6 - Abrami et al. 2008) in the plasma membrane and that the latter molecule can function as a co-receptor (Wei et al. 2006), the role of LRP6 in PA83 uptake remains uncertain (reviewed by van der Goot & Young 2009) and no function for LRP6 is annotated here.

### Literature references

- Wei, W., Lu, Q., Chaudry, GJ., Leppla, SH., Cohen, SN. (2006). The LDL receptor-related protein LRP6 mediates internalization and lethality of anthrax toxin. *Cell*, 124, 1141-54. [↗](#)
- Liu, S., Crown, D., Miller-Randolph, S., Moayeri, M., Wang, H., Hu, H. et al. (2009). Capillary morphogenesis protein-2 is the major receptor mediating lethality of anthrax toxin in vivo. *Proc. Natl. Acad. Sci. U.S.A.*, 106, 12424-9. [↗](#)
- Bradley, KA., Mogridge, J., Mourez, M., Collier, RJ., Young, JA. (2001). Identification of the cellular receptor for anthrax toxin. *Nature*, 414, 225-9. [↗](#)
- van der Goot, FG., Young, JA. (2009). Receptors of anthrax toxin and cell entry. *Mol. Aspects Med.*, 30, 406-12. [↗](#)
- Liu, S., Leppla, SH. (2003). Cell surface tumor endothelium marker 8 cytoplasmic tail-independent anthrax toxin binding, proteolytic processing, oligomer formation, and internalization. *J. Biol. Chem.*, 278, 5227-34. [↗](#)

### Editions

2013-12-13	Authored	D'Eustachio, P.
2014-05-19	Reviewed	Leppla, SH.
2014-05-23	Reviewed	Turk, BE.
2014-05-28	Reviewed	Moayeri, M., Liu, S.