

# Association of Acrosin Heavy and Light Chain

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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

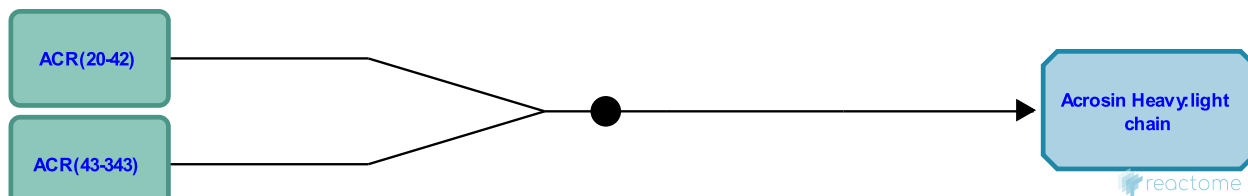
This document contains 1 reaction ([see Table of Contents](#))

## Association of Acrosin Heavy and Light Chain [↗](#)

**Stable identifier:** R-HSA-1297333

**Type:** binding

**Compartments:** extracellular region



The heavy and light chain proacrosin cleavage products bind to form an active acrosin complex.

### Literature references

Brady, RL., Tranter, R., Read, JA., Jones, R. (2000). Effector sites in the three-dimensional structure of mammalian sperm beta-acrosin. *Structure*, 8, 1179-88. [↗](#)

### Editions

2013-02-13	Authored	Gillespie, ME.
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