

HSPG2 binds FGF2(10-155), Fibronectin matrix, Transthyretin tetramer, PDGFA homodimer, PDGFB homodimer

Jupe, S., Ricard-Blum, S.

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

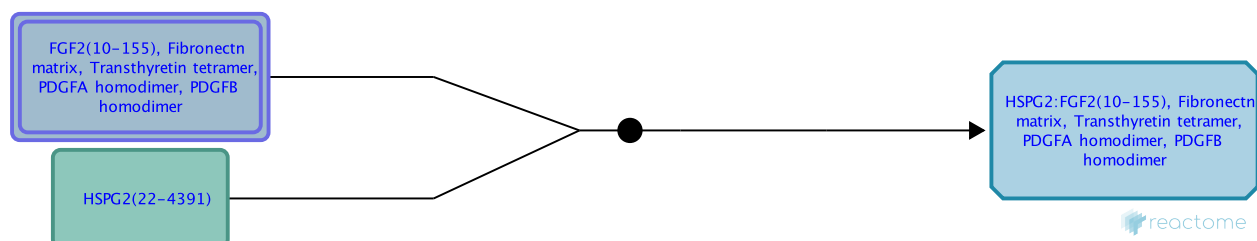
This document contains 1 reaction ([see Table of Contents](#))

HSPG2 binds FGF2(10-155), Fibronectn matrix, Transthyretin tetramer, PDGFA homodimer, PDGFB homodimer ↗

Stable identifier: R-HSA-2396337

Type: binding

Compartments: extracellular region



Perlecan (HSPG2) is a modular proteoglycan primarily located in the basement membranes of vascular tissues. It is involved in several developmental processes, both during embryogenesis and in human diseases such as cancer and diabetes (Iozzo et al. 1994). HSPG2 can self-aggregate into dimeric or multimeric forms (Yurchenco et al. 1987) and is involved in heterotypic interactions with numerous extracellular macromolecules (Whitelock et al. 2008, Perlecan entry in MatrixDB). HSPG2's GAG chains mediate interactions with fibroblast growth factor-2 (Vigny et al. 1988, Knox et al. 2002), and nidogens (Entactins, represented elsewhere). The core protein binds fibronectin (Isemura et al. 1987, Heremans et al. 1990, Vladavsky et al. 1991), transthyretin (Smeland et al. 1997) and platelet-derived growth factor A and B homodimers (Göhring et al. 1998).

Literature references

Knox, S., Merry, C., Stringer, S., Melrose, J., Whitelock, J. (2002). Not all perlecans are created equal: interactions with fibroblast growth factor (FGF) 2 and FGF receptors. *J. Biol. Chem.*, 277, 14657-65. ↗

Editions

2012-07-31	Authored	Jupe, S.
2013-04-26	Edited	Jupe, S.
2013-05-22	Reviewed	Ricard-Blum, S.