

INTL1 binds bacterial glycans

Hains, DS., Jupe, S., Shamovsky, V.

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 <u>Creative Commons Attribution 4.0 International (CC BY 4.0)</u> <u>License</u>. For more information see our <u>license</u>.

10/05/2024

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. 7
- Sidiropoulos, K., Viteri, G., Sevilla, C., Jupe, S., Webber, M., Orlic-Milacic, M. et al. (2017). Reactome enhanced pathway visualization. *Bioinformatics*, 33, 3461-3467. A
- 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. *オ*

This document contains 1 reaction (see Table of Contents)

INTL1 binds bacterial glycans 7

Stable identifier: R-HSA-6804527

Type: binding

Compartments: extracellular region, peptidoglycan-based cell wall



Intelectin-1 (INTL1) is a 120-kDa secretory lectin that recognizes multiple glycan epitopes found exclusively on microbes: beta-linked D-galactofuranose (beta-Galf), D-phosphoglycerol-modified glycans, heptoses, D-glycero-D-talo-oct-2-ulosonic acid (KO) and 3-deoxy-D-manno-oct-2-ulosonic acid (KDO) (Tsuji S et al. 2001; Wesener DA et al. 2015). These glycan residues are widely distributed in bacteria, including S. pneumoniae, Proteus mirabilis, Proteus vulgaris, Yersinia pestis and K. pneumoniae (Wesener DA et al. 2015). The 1.6-A-resolution crystal structure of human INTL1 complexed with beta-Galf suggests that INTL1 binds its carbohydrate ligands bearing terminal 1,2-diols through calcium ion-dependent coordination (Wesener DA et al. 2015).

Secreted INTL1 functions as a disulfide-linked trimer (Tsuji S et al., 2001; Tsuji S et al., 2007; Wesener DA et al., 2015).

Literature references

- Forest, KT., Smith, DF., Kiessling, LL., Zarling, LC., Kraft, MB., McBride, R. et al. (2015). Recognition of microbial glycans by human intelectin-1. *Nat. Struct. Mol. Biol.*, 22, 603-10. 7
- Suzuki, Y., Seya, T., Matsumoto, M., Toyoshima, K., Matsuhisa, A., Tsuji, S. et al. (2001). Human intelectin is a novel soluble lectin that recognizes galactofuranose in carbohydrate chains of bacterial cell wall. J. Biol. Chem., 276, 23456-63. ↗

Editions

2015-10-05	Authored	Shamovsky, V.
2016-04-15	Reviewed	Jupe, S.
2016-08-02	Reviewed	Hains, DS.
2016-08-15	Edited	Shamovsky, V.