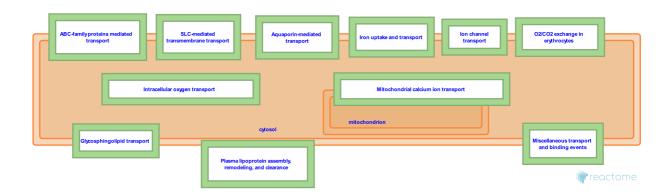


# Transport of small molecules



European Bioinformatics Institute, New York University Langone Medical Center, Ontario Institute for Cancer Research, Oregon Health and Science University.

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This is just an excerpt of a full-length report for this pathway. To access the complete report, please download it at the <a href="Reactome-Textbook">Reactome-Textbook</a>.

18/05/2024

#### 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.
- 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 data-base: Efficient access to complex pathway data. *PLoS computational biology, 14*, e1005968.

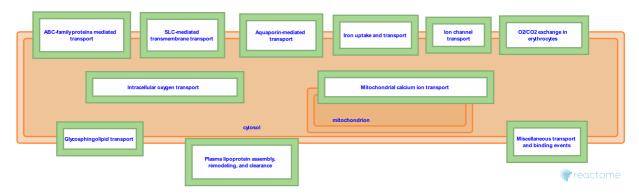
Reactome database release: 88

This document contains 12 pathways (see Table of Contents)

### **Transport of small molecules 对**

Stable identifier: R-XTR-382551

**Inferred from:** Transport of small molecules (Homo sapiens)



This event has been computationally inferred from an event that has been demonstrated in another species.

The inference is based on the homology mapping from PANTHER. Briefly, reactions for which all involved PhysicalEntities (in input, output and catalyst) have a mapped orthologue/paralogue (for complexes at least 75% of components must have a mapping) are inferred to the other species. High level events are also inferred for these events to allow for easier navigation.

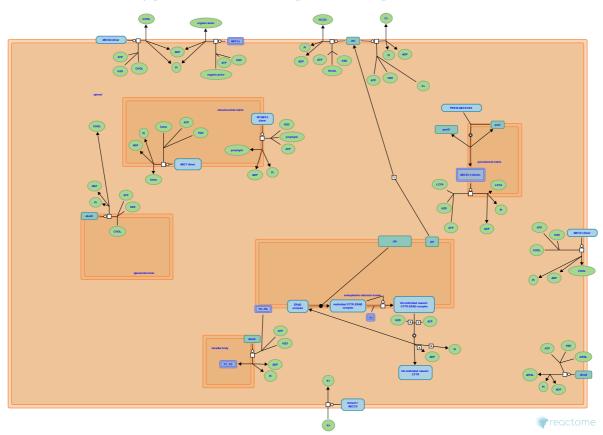
More details and caveats of the event inference in Reactome. For details on PANTHER see also: <a href="http://www.pantherdb.org/about.jsp">http://www.pantherdb.org/about.jsp</a>

### **ABC-family proteins mediated transport →**

**Location:** Transport of small molecules

Stable identifier: R-XTR-382556

Inferred from: ABC-family proteins mediated transport (Homo sapiens)



This event has been computationally inferred from an event that has been demonstrated in another species.

The inference is based on the homology mapping from PANTHER. Briefly, reactions for which all involved PhysicalEntities (in input, output and catalyst) have a mapped orthologue/paralogue (for complexes at least 75% of components must have a mapping) are inferred to the other species. High level events are also inferred for these events to allow for easier navigation.

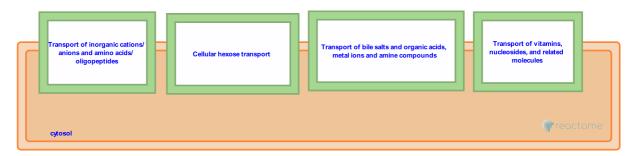
 $\underline{More\ details\ and\ cave ats\ of\ the\ event\ inference\ in\ Reactome.}\ For\ details\ on\ PANTHER\ see\ also: \\ \underline{http://www.pantherdb.org/about.jsp}$ 

### **SLC-mediated transmembrane transport →**

Location: Transport of small molecules

Stable identifier: R-XTR-425407

**Inferred from:** SLC-mediated transmembrane transport (Homo sapiens)



This event has been computationally inferred from an event that has been demonstrated in another species.

The inference is based on the homology mapping from PANTHER. Briefly, reactions for which all involved PhysicalEntities (in input, output and catalyst) have a mapped orthologue/paralogue (for complexes at least 75% of components must have a mapping) are inferred to the other species. High level events are also inferred for these events to allow for easier navigation.

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## **Aquaporin-mediated transport**

**Location:** Transport of small molecules

**Stable identifier:** R-XTR-445717

Compartments: plasma membrane

Inferred from: Aquaporin-mediated transport (Homo sapiens)



This event has been computationally inferred from an event that has been demonstrated in another species.

The inference is based on the homology mapping from PANTHER. Briefly, reactions for which all involved PhysicalEntities (in input, output and catalyst) have a mapped orthologue/paralogue (for complexes at least 75% of components must have a mapping) are inferred to the other species. High level events are also inferred for these events to allow for easier navigation.

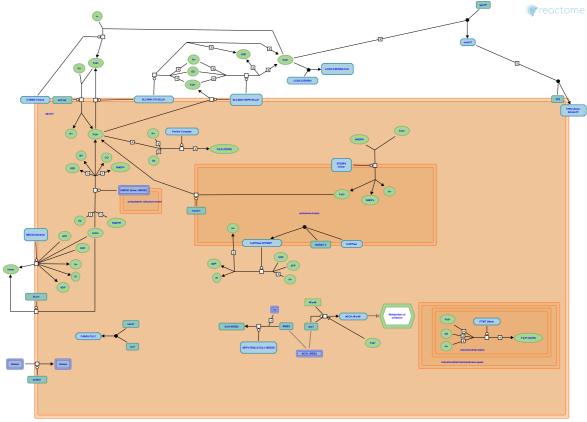
More details and caveats of the event inference in Reactome. For details on PANTHER see also: <a href="http://www.pantherdb.org/about.jsp">http://www.pantherdb.org/about.jsp</a>

### Iron uptake and transport **↗**

**Location:** Transport of small molecules

**Stable identifier:** R-XTR-917937

**Inferred from:** Iron uptake and transport (Homo sapiens)



This event has been computationally inferred from an event that has been demonstrated in another species.

The inference is based on the homology mapping from PANTHER. Briefly, reactions for which all involved PhysicalEntities (in input, output and catalyst) have a mapped orthologue/paralogue (for complexes at least 75% of components must have a mapping) are inferred to the other species. High level events are also inferred for these events to allow for easier navigation.

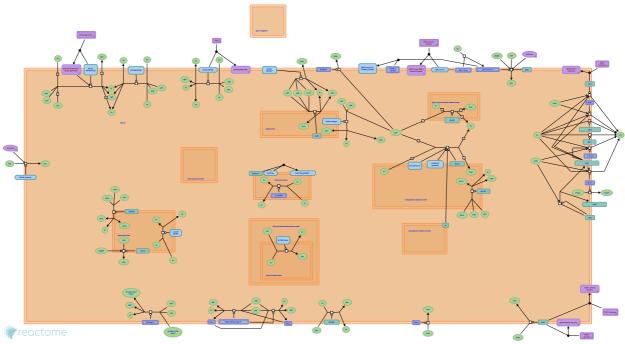
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### **Ion channel transport ₹**

**Location:** Transport of small molecules

**Stable identifier:** R-XTR-983712

**Inferred from:** Ion channel transport (Homo sapiens)



This event has been computationally inferred from an event that has been demonstrated in another species.

The inference is based on the homology mapping from PANTHER. Briefly, reactions for which all involved PhysicalEntities (in input, output and catalyst) have a mapped orthologue/paralogue (for complexes at least 75% of components must have a mapping) are inferred to the other species. High level events are also inferred for these events to allow for easier navigation.

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### 02/C02 exchange in erythrocytes **₹**

Location: Transport of small molecules

Stable identifier: R-XTR-1480926

Compartments: plasma membrane, extracellular region, cytosol

**Inferred from:** O2/CO2 exchange in erythrocytes (Homo sapiens)

Erythrocytes take up carbon dioxide and release oxygen

Erythrocytes take up oxygen and release carbon dioxide



This event has been computationally inferred from an event that has been demonstrated in another species.

The inference is based on the homology mapping from PANTHER. Briefly, reactions for which all involved PhysicalEntities (in input, output and catalyst) have a mapped orthologue/paralogue (for complexes at least 75% of components must have a mapping) are inferred to the other species. High level events are also inferred for these events to allow for easier navigation.

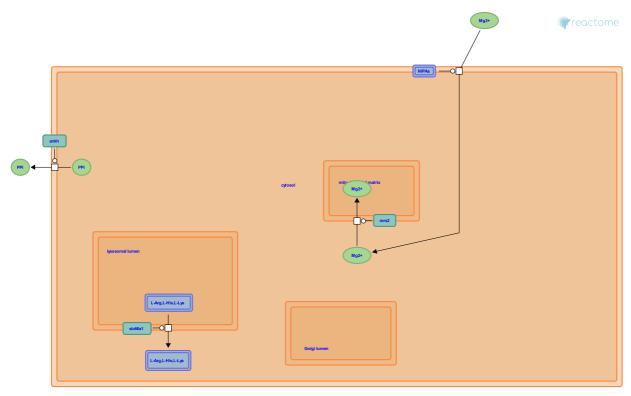
More details and caveats of the event inference in Reactome. For details on PANTHER see also: <a href="http://www.pantherdb.org/about.jsp">http://www.pantherdb.org/about.jsp</a>

### Miscellaneous transport and binding events **₹**

**Location:** Transport of small molecules

Stable identifier: R-XTR-5223345

**Inferred from:** Miscellaneous transport and binding events (Homo sapiens)



This event has been computationally inferred from an event that has been demonstrated in another species.

The inference is based on the homology mapping from PANTHER. Briefly, reactions for which all involved PhysicalEntities (in input, output and catalyst) have a mapped orthologue/paralogue (for complexes at least 75% of components must have a mapping) are inferred to the other species. High level events are also inferred for these events to allow for easier navigation.

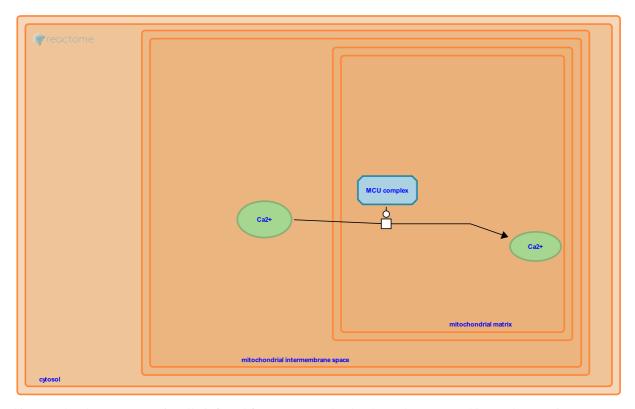
More details and caveats of the event inference in Reactome. For details on PANTHER see also: <a href="http://www.pantherdb.org/about.jsp">http://www.pantherdb.org/about.jsp</a>

### Mitochondrial calcium ion transport **→**

**Location:** Transport of small molecules

Stable identifier: R-XTR-8949215

**Inferred from:** Mitochondrial calcium ion transport (Homo sapiens)



This event has been computationally inferred from an event that has been demonstrated in another species.

The inference is based on the homology mapping from PANTHER. Briefly, reactions for which all involved PhysicalEntities (in input, output and catalyst) have a mapped orthologue/paralogue (for complexes at least 75% of components must have a mapping) are inferred to the other species. High level events are also inferred for these events to allow for easier navigation.

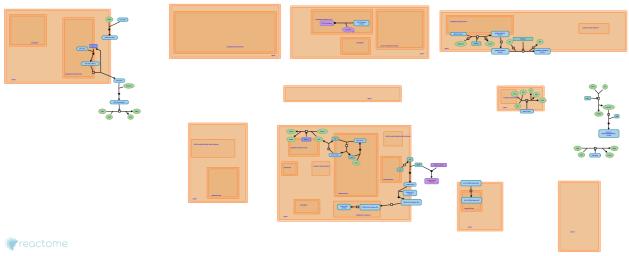
More details and caveats of the event inference in Reactome. For details on PANTHER see also: <a href="http://www.pantherdb.org/about.jsp">http://www.pantherdb.org/about.jsp</a>

### Plasma lipoprotein assembly, remodeling, and clearance **₹**

**Location:** Transport of small molecules

Stable identifier: R-XTR-174824

**Inferred from:** Plasma lipoprotein assembly, remodeling, and clearance (Homo sapiens)



This event has been computationally inferred from an event that has been demonstrated in another species.

The inference is based on the homology mapping from PANTHER. Briefly, reactions for which all involved PhysicalEntities (in input, output and catalyst) have a mapped orthologue/paralogue (for complexes at least 75% of components must have a mapping) are inferred to the other species. High level events are also inferred for these events to allow for easier navigation.

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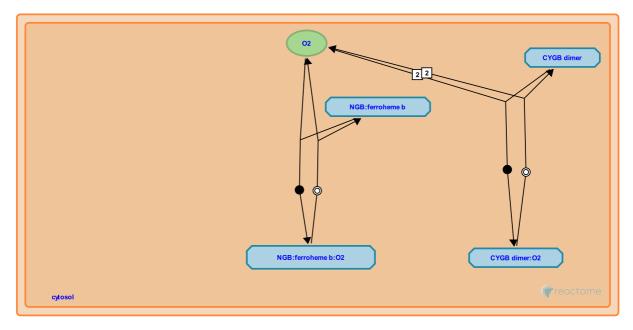
#### Intracellular oxygen transport 7

**Location:** Transport of small molecules

Stable identifier: R-XTR-8981607

**Compartments:** cytosol

Inferred from: Intracellular oxygen transport (Homo sapiens)



This event has been computationally inferred from an event that has been demonstrated in another species.

The inference is based on the homology mapping from PANTHER. Briefly, reactions for which all involved PhysicalEntities (in input, output and catalyst) have a mapped orthologue/paralogue (for complexes at least 75% of components must have a mapping) are inferred to the other species. High level events are also inferred for these events to allow for easier navigation.

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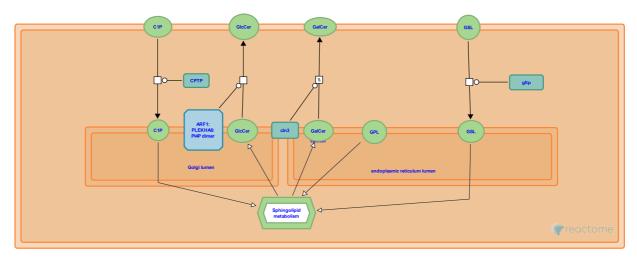
### **Glycosphingolipid transport**

**Location:** Transport of small molecules

Stable identifier: R-XTR-9845576

Compartments: endoplasmic reticulum membrane, plasma membrane, Golgi membrane, cytosol

**Inferred from:** Glycosphingolipid transport (Homo sapiens)



This event has been computationally inferred from an event that has been demonstrated in another species.

The inference is based on the homology mapping from PANTHER. Briefly, reactions for which all involved PhysicalEntities (in input, output and catalyst) have a mapped orthologue/paralogue (for complexes at least 75% of components must have a mapping) are inferred to the other species. High level events are also inferred for these events to allow for easier navigation.

More details and caveats of the event inference in Reactome. For details on PANTHER see also: <a href="http://www.pantherdb.org/about.jsp">http://www.pantherdb.org/about.jsp</a>

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