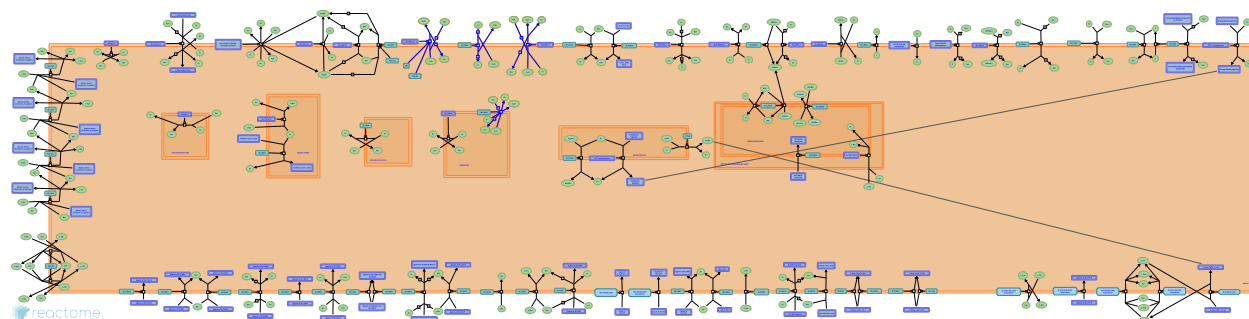


Sodium/Calcium exchangers



He, L., Jassal, B.

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

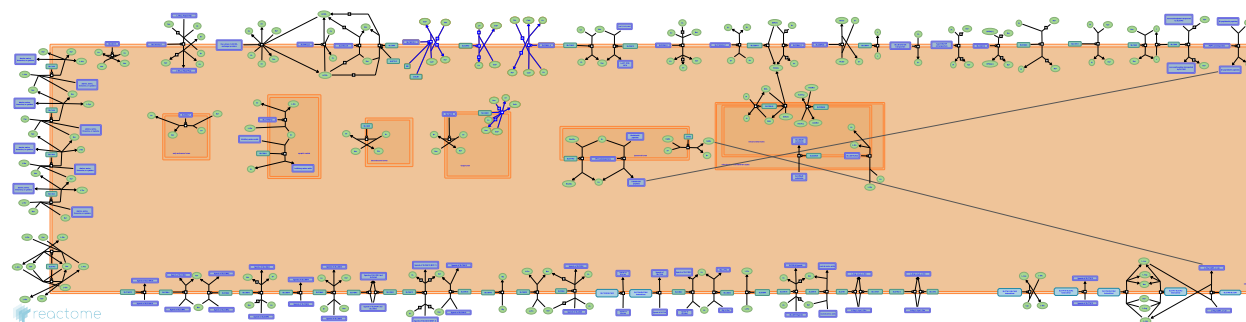
- Fabregat, A., Sidiropoulos, K., Viteri, G., Forner, O., Marin-Garcia, P., Arnau, V. et al. (2017). Reactome pathway analysis: a high-performance in-memory approach. *BMC bioinformatics*, 18, 142. [↗](#)
- Sidiropoulos, K., Viteri, G., Sevilla, C., Jupe, S., Webber, M., Orlic-Milacic, M. et al. (2017). 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: 70

This document contains 1 pathway and 4 reactions ([see Table of Contents](#))

Sodium/Calcium exchangers ↗

Stable identifier: R-HSA-425561



Calcium ions are used by cells as ubiquitous signalling molecules that control diverse physiological events. Three mammalian gene families control Ca^{2+} transport across plasma membranes and intracellular compartments (Lytton J, 2007). They are the $\text{Na}^{+}/\text{Ca}^{2+}$ exchanger family designated NCX (SLC8) (three members NCX1-3) (Quednau BD et al, 2004), the $\text{Na}^{+}/\text{Ca}^{2+}\text{-K}^{+}$ exchanger family designated NCKX (SLC24) (five members NCKX1-5) (Schnetkamp PP, 2004) and a Ca^{2+} /cation exchanger (NCKX6, NCLX) whose physiological function remains unclear.

Literature references

- Schnetkamp, PP. (2004). The SLC24 $\text{Na}^{+}/\text{Ca}^{2+}\text{-K}^{+}$ exchanger family: vision and beyond. *Pflugers Arch*, 447, 683-8. ↗
- Quednau, BD., Nicoll, DA., Philipson, KD. (2004). The sodium/calcium exchanger family-SLC8. *Pflugers Arch*, 447, 543-8. ↗
- Lytton, J. (2007). $\text{Na}^{+}/\text{Ca}^{2+}$ exchangers: three mammalian gene families control Ca^{2+} transport. *Biochem J*, 406, 365-82. ↗

Editions

2009-06-05	Authored, Edited	Jassal, B.
2009-08-24	Reviewed	He, L.

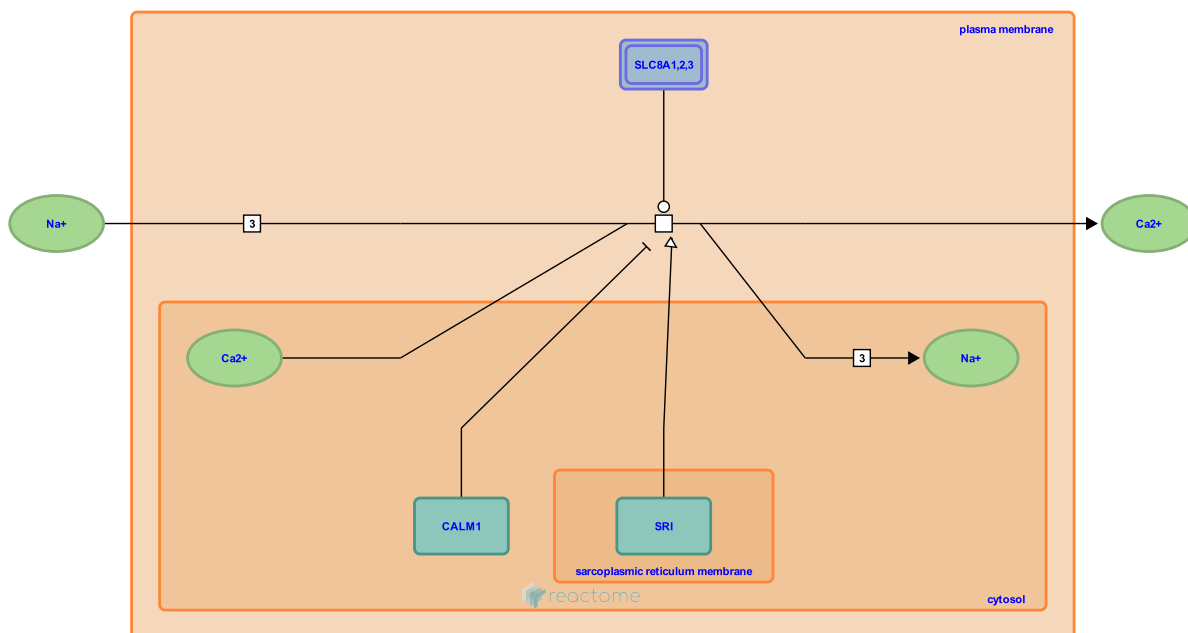
SLC8A1,2,3 exchange 3Na⁺ for Ca²⁺ ↗

Location: Sodium/Calcium exchangers

Stable identifier: R-HSA-425661

Type: transition

Compartments: plasma membrane, cytosol, extracellular region



The sodium/calcium exchangers 1, 2 and 3 (SLC8A1,2,3 aka NCX1,2,3) belong to one of three families that control Ca²⁺ flux across the plasma membrane or intracellular compartments. They extrude Ca²⁺ from the cell, using the electrochemical gradient of Na⁺ as it flows into the cell. One Ca²⁺ is exchanged for three Na⁺. During this electrogenic exchange, the membrane potential is altered. SLC8A1, 2, 3 play a minor role during phase 2, since they begin to restore ion concentrations. The high concentration of intracellular calcium starts contraction of those cells, which is sustained in the plateau phase. SLC8A1 has a ubiquitous expression profile (highest expression in heart, brain and kidney) and was originally cloned and characterized from human cardiac muscle (Komuro et al. 1992). Both SLC8A2 (Li et al. 1994) and SLC8A3 (Gabellini et al. 2002) are expressed in the brain.

In Rabbits, sorcin (SRI) activates SLC8A1, via the interaction of the respective Ca²⁺-binding domains (Zamparelli et al. 2010). Calmodulin (CALM1) binds to the cytoplasmic loop of NCX1 to negatively regulate exchange activity (Chou et al. 2015).

Literature references

- Komuro, I., Wenninger, KE., Philipson, KD., Izumo, S. (1992). Molecular cloning and characterization of the human cardiac Na⁺/Ca²⁺ exchanger cDNA. *Proc Natl Acad Sci U S A*, 89, 4769-73. ↗
- Li, Z., Matsuoka, S., Hryshko, LV., Nicoll, DA., Bersohn, MM., Burke, EP. et al. (1994). Cloning of the NCX2 isoform of the plasma membrane Na⁺-Ca²⁺ exchanger. *J Biol Chem*, 269, 17434-9. ↗
- Gabellini, N., Bortoluzzi, S., Danieli, GA., Carafoli, E. (2002). The human SLC8A3 gene and the tissue-specific Na⁺/Ca²⁺ exchanger 3 isoforms. *Gene*, 298, 1-7. ↗
- Zamparelli, C., Macquaide, N., Colotti, G., Verzili, D., Seidler, T., Smith, GL. et al. (2010). Activation of the cardiac Na⁺-Ca²⁺ exchanger by sorcin via the interaction of the respective Ca²⁺-binding domains. *J. Mol. Cell. Cardiol.*, 49, 132-41. ↗

Chou, AC., Ju, YT., Pan, CY. (2015). Calmodulin Interacts with the Sodium/Calcium Exchanger NCX1 to Regulate Activity. *PLoS ONE*, 10, e0138856. [↗](#)

Editions

2009-06-05	Authored, Edited	Jassal, B.
2009-08-24	Reviewed	He, L.

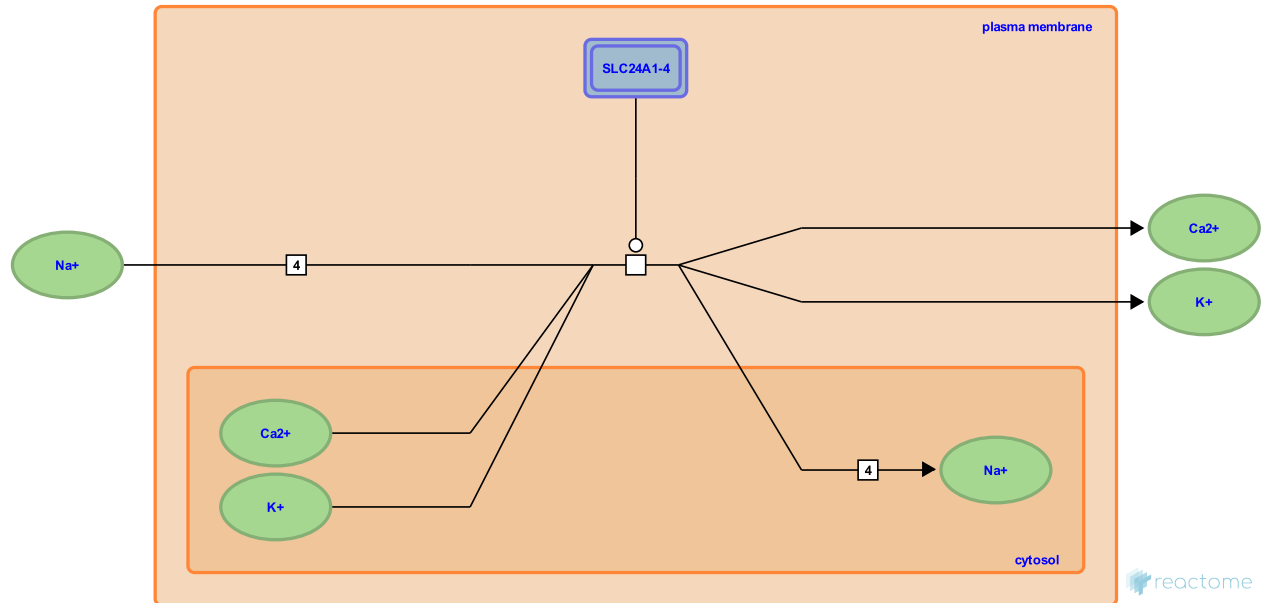
SLC24A1-4 exchange extracellular 4Na⁺ for cytosolic Ca²⁺, K⁺ ↗

Location: [Sodium/Calcium exchangers](#)

Stable identifier: R-HSA-425678

Type: transition

Compartments: plasma membrane



The five members of the NCKX (SLC24) family are all able to exchange one Ca²⁺ and one K⁺ for four Na⁺. NCKX1 (SLC24A1) encodes an exchanger protein which is the most extensively studied member (Tucker et al. 1998). It is highly expressed in the eye. Other members are expressed in the brain and skin as well as the eye (Prinsen et al. 2000, Kraev et al. 2001, Li et al. 2002, Lamason et al. 2005).

Literature references

- Prinsen, CF., Szerencsei, RT., Schnetkamp, PP. (2000). Molecular cloning and functional expression of the potassium-dependent sodium-calcium exchanger from human and chicken retinal cone photoreceptors. *J Neurosci*, 20, 1424-34. ↗
- Tucker, JE., Winkfein, RJ., Murthy, SK., Friedman, JS., Walter, MA., Demetrick, DJ. et al. (1998). Chromosomal localization and genomic organization of the human retinal rod Na-Ca+K exchanger. *Hum Genet*, 103, 411-4. ↗
- Kraev, A., Quednau, BD., Leach, S., Li, XF., Dong, H., Winkfein, R. et al. (2001). Molecular cloning of a third member of the potassium-dependent sodium-calcium exchanger gene family, NCKX3. *J Biol Chem*, 276, 23161-72. ↗
- Li, XF., Kraev, AS., Lytton, J. (2002). Molecular cloning of a fourth member of the potassium-dependent sodium-calcium exchanger gene family, NCKX4. *J Biol Chem*, 277, 48410-7. ↗
- Lamason, RL., Mohideen, MA., Mest, JR., Wong, AC., Norton, HL., Aros, MC. et al. (2005). SLC24A5, a putative cation exchanger, affects pigmentation in zebrafish and humans. *Science*, 310, 1782-6. ↗

Editions

2009-06-05	Authored, Edited	Jassal, B.
2009-08-24	Reviewed	He, L.

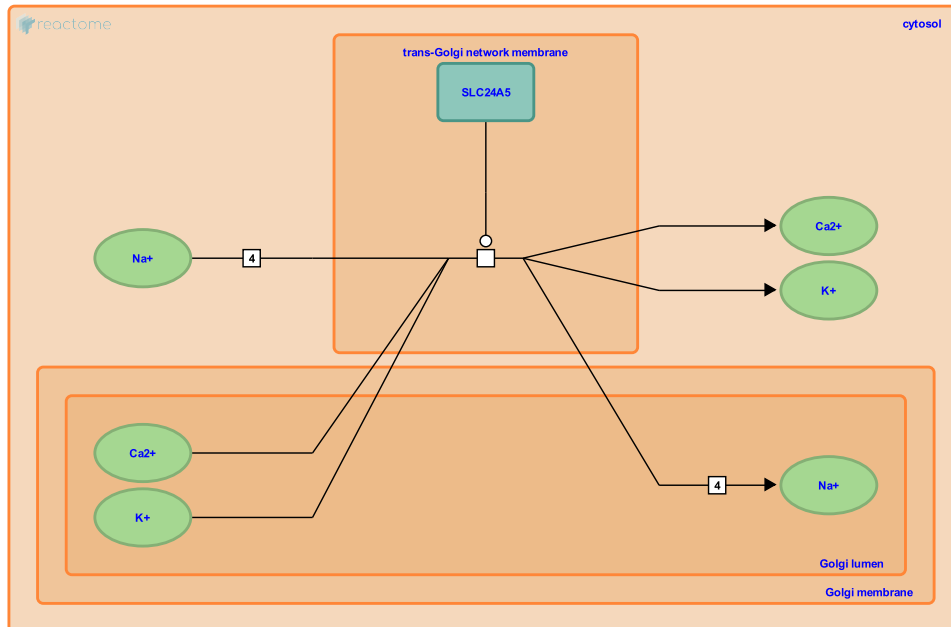
SLC24A5 exchanges cytosolic 4Na⁺ for Golgi luminal Ca²⁺, K⁺ ↗

Location: [Sodium/Calcium exchangers](#)

Stable identifier: R-HSA-5626316

Type: transition

Compartments: trans-Golgi network membrane, cytosol, Golgi lumen



The five members of the NCKX (SLC24) family are all able to exchange one Ca²⁺ and one K⁺ for four Na⁺. SLC24A5 encodes a trans-Golgi network exchanger protein NCKX5 which is expressed in melanocytes and regulates human epidermal melanogenesis (Ginger et al. 2008).

Literature references

Ginger, RS., Askew, SE., Ogborne, RM., Wilson, S., Ferdinando, D., Dadd, T. et al. (2008). SLC24A5 encodes a trans-Golgi network protein with potassium-dependent sodium-calcium exchange activity that regulates human epidermal melanogenesis. *J. Biol. Chem.*, 283, 5486-95. ↗

Editions

2009-08-24	Reviewed	He, L.
2014-10-09	Authored, Edited, Revised	Jassal, B.

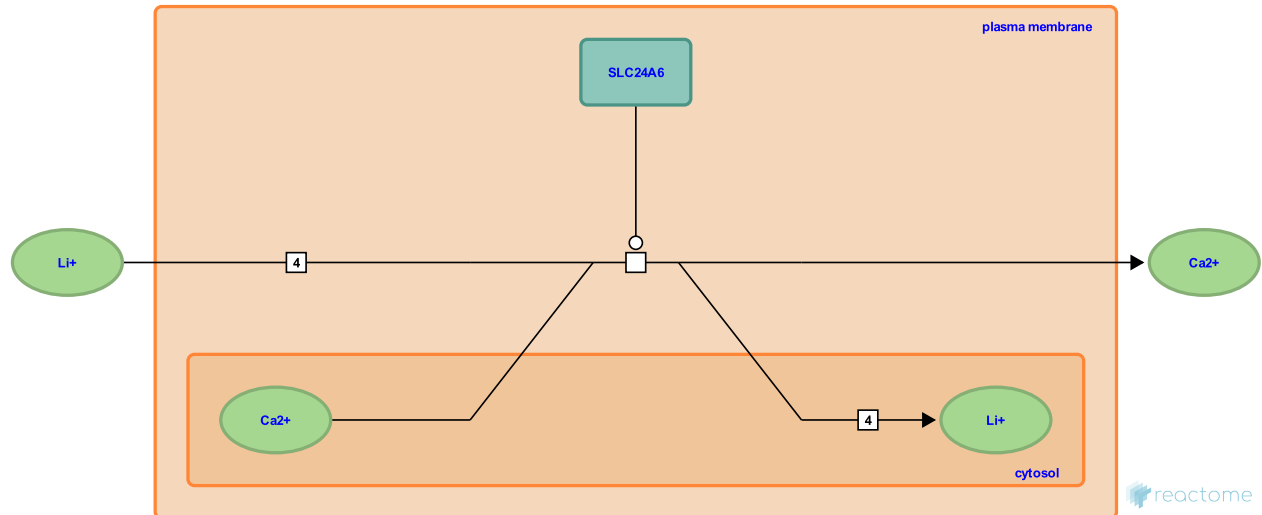
K⁺-independent Li⁺/Ca²⁺ exchanger transport ↗

Location: [Sodium/Calcium exchangers](#)

Stable identifier: R-HSA-425822

Type: transition

Compartments: plasma membrane



SLC24A6 (NCKX6, NCLX) (Palty R et al, 2004) encodes a protein which can transport Li⁺ or Na⁺ in exchange for Ca²⁺ in a K⁺-independent manner (Cai X and Lytton J, 2004). Lithium exchange with calcium is shown here.

Literature references

Palty, R., Ohana, E., Hershinkel, M., Volokita, M., Elgazar, V., Beharier, O. et al. (2004). Lithium-calcium exchange is mediated by a distinct potassium-independent sodium-calcium exchanger. *J Biol Chem*, 279, 25234-40. ↗

Cai, X., Lytton, J. (2004). Molecular cloning of a sixth member of the K⁺-dependent Na⁺/Ca²⁺ exchanger gene family, NCKX6. *J Biol Chem*, 279, 5867-76. ↗

Editions

2009-06-08	Authored, Edited	Jassal, B.
2009-08-24	Reviewed	He, L.

Table of Contents

Introduction	1
☒ Sodium/Calcium exchangers	2
↳ SLC8A1,2,3 exchange 3Na ⁺ for Ca ²⁺	3
↳ SLC24A1-4 exchange extracellular 4Na ⁺ for cytosolic Ca ²⁺ , K ⁺	5
↳ SLC24A5 exchanges cytosolic 4Na ⁺ for Golgi luminal Ca ²⁺ , K ⁺	6
↳ K ⁺ -independent Li ⁺ /Ca ²⁺ exchanger transport	7
Table of Contents	8