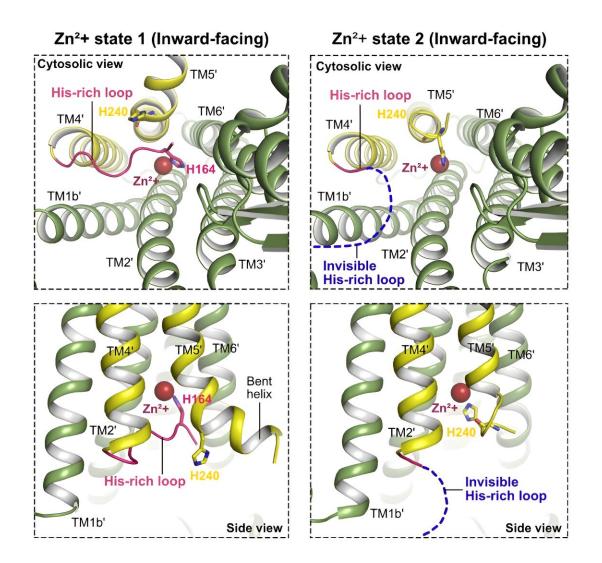


## Advanced technology reveals intricate details of zinc transportation in cells

August 31 2023



The histidine-rich loop segment is inserted into the cytosolic cavity and coordinated to  $Zn^{2+}$  at the transmembrane metal-binding site for efficient  $Zn^{2+}$  uptake. Credit: Han Ba Bui and Kenji Inaba



A group of researchers has unearthed the secrets behind a tiny but crucial protein that shuttles zinc ions  $(Zn^{2+})$  within our bodies. The discovery offers a deeper understanding of how our cells maintain optimal health.

 $Zn^{2+}$  may be small, but they play a mighty role in our cells. Zinc enables enzyme catalysis, <u>protein folding</u>, DNA binding, and regulating <u>gene</u> <u>expression</u>, with about 10% of the proteins in our body reliant on  $Zn^{2+}$  to function effectively.

The study, which was published in the journal *Nature Communications* on August 8, 2023, focused on the Golgi apparatus—a cellular compartment that processes, sorts and distributes cells to their final destination. Within the Golgi, three distinct zinc transporter (ZnT) complexes—ZnT4, ZnT5/6, and ZnT7—collaborate to usher Zn<sup>2+</sup> ions from the cellular interior (cytosol) into the Golgi. While these complexes have long been known to play pivotal roles, the precise mechanisms governing Zn<sup>2+</sup> transport within them have remained an enigma.

"We concentrated our study on the transport protein hZnT7," says Kenji Inaba, a corresponding author of the study and professor at Tohoku University's Institute of Multidisciplinary Research for Advanced Materials Sciences. "The study built upon our previous research that hZnT7 plays a vital role in Zn<sup>2+</sup> uptake into the cis-Golgi cisterna and regulates the localization, traffic and function of the chaperone protein ERp44."

To reveal more about hZnT7, Inaba and his colleagues employed an advanced technique called <u>cryo-electron microscopy</u> (cryo-EM). Two cryo-EM machines, one from Tohoku University and one from the University of Tokyo, could capture detailed images of hZnT7 in action.



By using a Fab fragment from a monoclonal antibody that specifically binds hZnT7 the researchers succeeded in determining the cryo-EM structures of hZnT7> at near-atomic resolutions, gaining critical insights into the mechanisms of  $Zn^{2+}$  transport.

Comparative analysis between hZnT7 and other zinc transporters, including human ZnT8 and bacterial YiiP, exposed distinct structural features of hZnT7. The existence of hZnT7 as a homodimer with varying  $Zn^{2+}$ -bound configurations holds particular significance.

Notably, hZnT7 boasts an elongated cytosolic histidine-rich loop (Hisloop) that interfaces with the transmembrane metal-binding site, a vital feature governing zinc transfer. During  $Zn^{2+}$  recruitment via the Hisloop, hZnT7 undergoes intricate conformational rearrangements, shedding light on an unparalleled mechanism of zinc transport.

It is widely known that hZnT7 is a key player in dietary zinc absorption and controlling body fat. When <u>zinc</u> levels drop in certain parts of the body, it can lead to issues like prostate cancer development in mice and disruptions in how our bodies process insulin.

Inaba adds their findings will result in greater understandings of the molecular processes with certain pathogens. "With it reported that abnormalities in Golgi-resident ZnT transporters result in fatal diseases such as diabetes, cancers, and immunodeficiency, it's essential to understand the pathogenic mechanisms of these diseases at the molecular and cellular level."

**More information:** Han Ba Bui et al, Cryo-EM structures of human zinc transporter ZnT7 reveal the mechanism of Zn2+ uptake into the Golgi apparatus, *Nature Communications* (2023). DOI: 10.1038/s41467-023-40521-5



## Provided by Tohoku University

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