

Intracellular Transition Metal Homeostasis in Plants and Algae

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Keywords: epigenetics, heavy metal interactions, metalloproteins, metal uptake of organelles, transition metal sensing, transition metal signaling

The language style of the manuscript is British English.

Abstract

Essential transition metals such as copper (Cu), iron (Fe), manganese (Mn), molybdenum (Mo), and zinc (Zn) have prime importance in the metabolism of all living beings, including photosynthetic eukaryotes. Based on their redox-active nature these metals generally serve as cofactors of redox enzymes and thus contribute to a large diversity of enzymatic reactions, including the respiratory and the photosynthetic electron transport chains. Improper delivery and accumulation of these transition metals also challenge the redox balance of cells. Oxidized transition metals generate free radicals that damage lipids, macromolecules, and metabolites, and thus their production leads to disruption of metabolic and membrane function. Non-essential metals and metalloids, among others cadmium (Cd), lead (Pb), mercury (Hg), aluminium (Al), and arsenic (As) often interfere with the metabolism of essential metals, leading to, over-direct toxicity, and induce essential metal toxicity as well. In consequence, sensing, regulation, accumulation, and incorporation of the essential transition metals play a pivotal role in intracellular metal homeostasis. The collection “Intracellular Transition Metal Homeostasis in Plants and Algae” brings together pieces of current information on how plant cells sense and regulate the uptake of essential transition metals, and how non-essential transition metals and metalloids influence the epigenome. Articles in the collection highlight still open questions and indicate future research directions in the intracellular sensing, transport and interactions of transition metals.

1 Editorial overview

Plants established a highly structured regulatory network managing homeostasis of the essential transition metal which are widely employed as cofactors of redox enzymes. In the past century, dedicated transport systems have been identified for the uptake and translocation of essential transition metals (Eide *et al.* 1996; Curie *et al.* 2000). The current challenge in our understanding is how regulatory and intracellular transport processes contribute to the balanced metal homeostasis of plants.

Iron is the most abundant transition metal in all organisms and thus the most widely studied one in plants (Jeong 2023). Vélez-Bermúdez and Schmidt (2023) review the knowledge of the past decade on Fe sensing and regulation. Although Fe sensing of mammalian cells is long known, the analogous mechanism of plant cells has a long debate. In plants, hemerythrin domain-containing proteins were found to be the key Fe sensors: both monocot and dicot Fe sensors (HRZ; BTS, respectively) express E3 ubiquitin ligase activity by their RING domain. This domain indeed shares a functional analogy to the mammalian Fe sensor FBXL5. The authors also question whether multiple Fe sensor systems in plant cells could orchestrate responses. According to current information, this question cannot be answered without any doubts. Downstream elements of the currently accepted HRZ/BTS systems are transcription factors of the bHLH family. Spielmann *et al.* (2023) review this complex cascade of bHLH transcription factors including their target genes involved in Fe uptake, mobilization and storage. Spielman *et al.* also present the current knowledge of alternative splicing upon Fe deficiency and the effect of non-Fe-metals on the post-translational control of the two well-characterized Fe transporters, NRAMP1 and IRT1. Lacking knowledge relates to the effect of post-translational modifications on the activity of transporters, their removal from the plasma membrane and their cellular polarity for directional uptake.

Loading of Fe into plastids is highly important in photosynthetically active cells. Since the discovery of the plastid localized FRO7 (Jeong *et al.* 2008) a light-dependent, reduction-based plastidial Fe uptake (Solti *et al.* 2012) was considered. Fe uptake in etioplasts is essential for the transformation towards chloroplasts and accumulation of the photosynthetic electron transport chain members. Yet, Fe acquisition in the absence of light cannot be a light-dependent process. According to *in vivo* Fe uptake and ferric chelate reductase activity of chloroplast envelope membranes presented by Sági-Kazár *et al.* (2023), the Fe uptake in etioplasts operates as a reduction-based process.

Zn homeostasis is also tightly modulated to avoid oxidative stress and cytotoxic interactions. Moreover, Zn is required for multiple chloroplast functions (Hübner and Haase 2021). To achieve Zn

homeostasis, a similar diversity of Zn managing proteins is expected as for Fe management. Zhang *et al.* (2023) investigated two *Arabidopsis* nucleotide-dependent metallochaperones ZNG1 & 2, orthologues to human and fungal ZNG1s. Based on structural modelling, protein localization, RNA-seq, and proteomic data, they found that ZNG1 interacts with the methionine aminopeptidases, MAP1A, and serves as Zn transferase for AtMAP1A. The target(s) of ZNG2s were presumed to be conserved plastid-localized proteins but need to be identified.

Balancing intracellular transition metal homeostasis involves dynamic genomic and transcriptional regulation. Evidences are increasing that among others altered Fe homeostasis changes the epigenetic regulation of Fe responses (Shafiq *et al.* 2020; Sun *et al.* 2021). Along with DNA methylation, the modification of histones represents a second platform of epigenetic regulations. Under Fe excess, repressive lysine methylation occurs on H3 histones leading to the suppression of Fe uptake (Séré and Martin, 2020). Non-essential metals and metalloids are serious threats to intracellular essential transition metal homeostasis by impacting, among others, protein folding, redox balance, and the incorporation of cofactors into metalloproteins. Chmielowska-Bąk *et al.* (2023) review the findings of non-essential metals and metalloids on the epigenetic regulation. The importance of their summary is that comprehensive reviews of the plant's epigenetic responses to metals are still scarce. Although the information on epigenetic and epitranscriptomic alterations upon non-essential metal stresses is controversial, the higher methylation status and the activation of DNA methyltransferases, could result in a higher resilience of the nuclear DNA against damages, and could also affect the expression pattern of transporters, changes which would correlate with a higher tolerance against non-essential metals and metalloids.

2 Conclusions and Future Perspectives

Metal homeostasis is essential for development and environmental adaptation. In the intracellular transition metal homeostasis, characterization of Fe sensors, transporters, and transcription factor interactions are among the most important achievements. Indeed, sensing mechanisms for non-Fe metals and proposed additional Fe sensing systems are not yet completely resolved. Knowledge is slowly rising on the regulatory mechanisms at the epigenome and epitranscriptome level controlling the stability of the intracellular transition metal homeostasis. Future understanding of these mechanisms will reveal how stress memory, environmental adaptation, and intracellular interactions are induced by transition metals. Over molecular approaches, more data are needed from physical techniques that allow visualization of transition metals at tissue, cell, and intracellular levels. These

include X-ray absorption and emission spectroscopy for the analysis of biological metal complexes (among others μ XANES, EXAFS) and the distribution of transition metals (among others μ XRF, EDX, PIXE), respectively. For the detection of transition metals in metalloprotein improved methods are needed such as among others HPLC-ICP-MS based techniques. Subcellular, and quantitative proteomic approaches are also of a growing importance. Future advancements in proteomic methodologies will facilitate the comprehensive characterization of the plant metalloproteome. Integrating multi-omics datasets will deeper the insight into the mechanisms governing plant metal homeostasis and enables to develop strategies for enhancing metal nutrition, stress tolerance, and crop productivity.

3 Funding

Á.S. was supported by the János Bolyai Scholarship of the Hungarian Academy of Sciences, by the grant K-146865 of National Research, Development and Innovation Office, Hungary (NKFIH), and by the ÚNKP-23-5 New National Excellence Program of the Ministry for Culture and Innovation, Hungary from the source of the NKFIH Fund. M-T.H. was supported by the Austrian Science Fund (FWF) project Nr. 10.55776/DOC111 and the COST Action CA19116 “Plant Metals”.

4 Conflict of interest

All authors state that there is no conflict of interest in relation with the present work.

5 Data availability

This editorials do not include unpublished or original data.

6 References

- Curie, C., Alonso, J. M., Jean, M. L., Ecker, J. R., and Briat, J. F. (2000). Involvement of NRAMP1 from *Arabidopsis thaliana* in iron transport. *Biochem. J.* 347, 749-755.
- Eide, D., Broderius, M., Fett, J., and Gueriot, M. L. (1996). A novel iron-regulated metal transporter from plants identified by functional expression in yeast. *PNAS* 93, 5624-5628.
- Hübner, C., and Haase, H. (2021). Interactions of zinc-and redox-signaling pathways. *Redox Biol.* 41, 101916.
- Jeong, J., Cohu, C., Kerkeb, L., Pilon, M., Connolly, E. L., and Gueriot, M. L. (2008). Chloroplast Fe (III) chelate reductase activity is essential for seedling viability under iron limiting conditions. *PNAS* 105, 10619-10624.

Jeong, J. (Ed.). (2023). Plant iron homeostasis: Methods and protocols (Vol. 2665). New York, NY: Springer US.

Séré, D., and Martin, A. (2020). Epigenetic regulation: another layer in plant nutrition. *Plant Signal. Behav.* 15, 1-6.

Shafiq, S., Ali, A., Sajjad, Y., Zeb, Q., Shahzad, M., Khan, A. R., ... & Widemann, E. (2020). The interplay between toxic and essential metals for their uptake and translocation is likely governed by DNA methylation and histone deacetylation in maize. *Int. J. Mol. Sci.* 21, 6959.

Solti, Á., Kovács, K., Basa, B., Vértés, A., Sárvári, É., and Fodor, F. (2012). Uptake and incorporation of iron in sugar beet chloroplasts. *Plant Physiol. Biochem.* 52, 91-97.

Sun, S., Zhu, J., Guo, R., Whelan, J., and Shou, H. (2021). DNA methylation is involved in acclimation to iron deficiency in rice (*Oryza sativa*). *Plant J.* 107, 727-739.