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Using μ PIXE for quantitative mapping of metal concentration in *Arabidopsis thaliana* seeds

Magali Schnell Ramos^{1,2*}, Hicham Khodja³, Viviane Mary¹ and Sébastien Thomine¹

¹ Institut des Sciences du Végétal, UPR2355, Centre National de la Recherche Scientifique, Gif-sur-Yvette, France

² Chimica Agraria, Dipartimento di Scienze Agrarie e Ambientali, Università degli Studi di Udine, Udine, Italy

³ Laboratoire d'Etude des Eléments légers, SIS2M, UMR 3299, CEA-CNRS, CEA Saclay, Gif-sur-Yvette, France

Edited by:

Jean-François Briat, Centre National de la Recherche Scientifique, France

Reviewed by:

Stéphane Mari, Institut National pour la Recherche Agronomique, France

Marie-Pierre Isaure, Université de Pau et des Pays de l'Adour and CNRS, France

*Correspondence:

Magali Schnell Ramos, Chimica Agraria, Dipartimento di Scienze Agrarie e Ambientali, Università degli Studi di Udine, Via delle Scienze 208, 33100 Udine, Italy
e-mail: magali.schnell@uniud.it

Seeds are a crucial stage in plant life. They contain the nutrients necessary to initiate the development of a new organism. Seeds also represent an important source of nutrient for human beings. Iron (Fe) and zinc (Zn) deficiencies affect over a billion people worldwide. It is therefore important to understand how these essential metals are stored in seeds. In this work, Particle-Induced X-ray Emission with the use of a focused ion beam (μ PIXE) has been used to map and quantify essential metals in *Arabidopsis* seeds. In agreement with Synchrotron radiation X-ray fluorescence (SXRF) imaging and Perls/DAB staining, μ PIXE maps confirmed the specific pattern of Fe and Mn localization in the endodermal and subepidermal cell layers in dry seeds, respectively. Moreover, μ PIXE allows absolute quantification revealing that the Fe concentration in the endodermal cell layer reaches $\sim 800 \mu\text{g}\cdot\text{g}^{-1}$ dry weight. Nevertheless, this cell layer accounts only for about half of Fe stores in dry seeds. Comparison between *Arabidopsis* wild type (WT) and mutant seeds impaired in Fe vacuolar storage (*vit1-1*) or release (*nramp3nramp4*) confirmed the strongly altered Fe localization pattern in *vit1-1*, whereas no alteration could be detected in *nramp3nramp4* dry seeds. Imaging of imbibed seeds indicates a dynamic localization of metals as Fe and Zn concentrations increase in the subepidermal cell layer of cotyledons after imbibition. The complementarities between μ PIXE and other approaches as well as the importance of being able to quantify the patterns for the interpretation of mutant phenotypes are discussed.

Keywords: iron, seed, *Arabidopsis*, μ PIXE, elemental mapping, quantitative

INTRODUCTION

Thanks to their redox properties or Lewis acid strength under physiological conditions, iron (Fe), manganese (Mn), copper (Cu), or zinc (Zn) act as major cofactors in many enzymes such as proteases or antioxidant enzymes as well as in electron transfer chains of mitochondrial respiration or chloroplast photosynthesis (Marschner, 2012). While micronutrients are essential at adequate concentrations, excess leads to toxicity. Moreover, certain trace metals such as cadmium (Cd) or mercury (Hg) have no known function as nutrients and are potentially toxic at low concentrations (Clemens, 2006; Clemens et al., 2013). Therefore, mechanisms for tight regulation of metal homeostasis are vital (Kraemer and Clemens, 2006; Marschner, 2012).

Seeds are a crucial stage in the life of a plant. They contain the macro- and micronutrients necessary to initiate the development of a new organism. More specifically, essential micronutrients need to be safely stored and readily remobilized during early germination. Seeds also represent an important source of nutrient for human beings. Fe and Zn deficiencies affect over a billion people worldwide (Murgia et al., 2012). Consequently, studies on seed transition metal content are necessary. Despite its relatively high abundance, Fe availability remains limited for living organisms. In seeds, Fe is stored either as highly bio-available phytoferritins localized in plastids, or as poorly bio-available

Fe-phytate salts, localized in vacuolar inclusions called globoids (Harrison and Arosio, 1996; Briat and Lobreaux, 1997; Otegui et al., 2002; Lanquar et al., 2005). The balance between those two forms is different according to plant species and is greatly modified during germination. Besides, in seeds, phytate salts are also the main storage form of potassium (K), magnesium (Mg), calcium (Ca), and Zn (Mikus et al., 1992).

In *Arabidopsis*, VIT1, a vacuolar Fe transporter required for Fe storage in the vacuole during seed formation, was identified by Kim et al. (2006). Synchrotron radiation X-ray fluorescence (SXRF) tomographic imaging demonstrated that loss of VIT1 function disrupts the cell specific localization of Fe in dry seeds. Whereas Fe is stored in endodermal cells surrounding the provascular tissues in wild type (WT), it co-localizes with Mn in the subepidermal cell layer of *vit1-1* knockout mutant embryos (Kim et al., 2006; Roschztardt et al., 2009). Fe mislocalization results in drastically decreased viability of *vit1-1* seedlings under Fe deficiency. While VIT1 mediates Fe influx into the vacuole, NRAMP3 and NRAMP4 metal transporters have been shown to act redundantly to export Fe out of the vacuole (Lanquar et al., 2005). Energy-Dispersive X-ray (EDX) technique indicated that the *nramp3nramp4* double knockout mutant is defective in Fe retrieval from seed vacuolar globoids during germination. As a consequence, *nramp3nramp4* mutant seedlings display an early

developmental arrest when germinated on low Fe. Furthermore, the drastic morphological and biological changes that occur during germination must be accompanied by a relocation of nutrients to the sites where they are required for metabolism. Although several reports have addressed metal patterning in dry seed, the changes in metal localization and their kinetics upon seed germination have not been addressed at the tissue level. Three imaging techniques have been used to investigate Fe distribution in Arabidopsis WT and mutant seeds EDX (Lanquar et al., 2005), SXRF (Kim et al., 2006), and Perls/DAB staining (Roschztardt et al., 2009). However, those techniques provided either non-quantitative data (EDX, Perls/DAB) or approximate quantification (SXRF) of metal concentrations in the different seed tissues.

Particle-Induced X-ray Emission induced by a focused ion beam (μ PIXE) allows multi-elemental mapping in biological samples with high spatial resolution (1 μ m range) and high sensitivity (down to μ g·g⁻¹ range). Importantly, μ PIXE technique presents the unique advantage of providing quantitative results when used simultaneously with Rutherford Backscattering (RBS) and Scanning Transmission Ion Microscopy (STIM) analyses (Deves et al., 2005). The combined measurements of trace element amount by PIXE, charge monitoring and organic element determination by RBS and sample local mass determination by STIM are often referred as “fully” quantitative results in the literature in opposition to “semi”-quantitative results obtained by other imaging techniques.

In plants, μ PIXE has been used for the localization and quantification of essential macro- and micronutrients in specific tissues and organs such as elemental mapping of buckwheat seeds (Vogel-Mikus et al., 2009), Fe in barley roots (Schneider et al., 2002), Fe and Zn in Phaseolus seeds (Cvitanich et al., 2010, 2011), Cu in *Brassica carinata* leaf and root (Cestone et al., 2012). μ PIXE was also used to image and quantify non-essential elements. In the context of environmental contamination, μ PIXE was used to study cesium (Cs) in Arabidopsis leaf, stem, and trichome (Isaure et al., 2006), Cd and Ni in soybean seed (Malan et al., 2012), and uranium (U) in leaf and root of oilseed rape, sunflower, and wheat (Laurette et al., 2012). In the context of metal hyperaccumulation, it was used to analyze Cd in leaf and seed of *Thalpi praecox* (Vogel-Mikus et al., 2007, 2008) or nickel (Ni) in *Berkheya coddii* leaves (Budka et al., 2005).

Here, the μ PIXE approach was used to quantitatively analyze metal distribution in Arabidopsis seeds. A sample preparation protocol suitable for μ PIXE analysis of *Arabidopsis thaliana* dry but also imbibed seeds was established. Analysis of some nutritionally important elements by μ PIXE mapping confirmed the previously established pattern in WT dry seed metal distribution, exhibiting Mn accumulation at the abaxial side of cotyledons as well as Fe localization around the provascular tissues. Local Fe, Mn, and Zn concentrations were determined in these tissues in WT and both *nramp3nramp4* and *vit1-1* mutant seeds. Moreover, a comparison between elemental maps obtained with dry or imbibed seeds revealed early changes in metal localization. The μ PIXE results are put in perspective with other elemental analysis techniques raising questions regarding the input of element quantification in the interpretation of mutant phenotypes.

Finally, μ PIXE results obtained for WT dry and imbibed seeds are discussed with respect to the possible use of this technique to study dynamic element redistribution.

MATERIALS AND METHODS

PLANT MATERIAL

Mature (dry) and imbibed seeds of Arabidopsis (*Arabidopsis thaliana* accession Columbia-0) WT, *nramp3nramp4* (*nr3nr4*) and *vit1-1* mutants were used. Generation of both mutants has been described previously in Ravet et al. (2009b) and Kim et al. (2006), respectively. Dry seeds of all genotypes were harvested from plants grown on potting soil (Tonerde PAM argile, Brill France) in a greenhouse under a 16 h photoperiod with regular watering (2–3 times per week). Seeds were imbibed in deionized water on an orbital shaker (40 rotations·min⁻¹), under continuous light (60 μ mol photon·m⁻²·s⁻¹) at 21°C for 48 h.

AAS ANALYSES OF DRY SEEDS

Three to four replicates of circa 20 mg of dry seeds were digested in 2 ml of 70% nitric acid in a DigiBlock ED36 (LabTech, Italy) at 80°C for 1 h, 100°C for 1 h, and 120°C for 2 h. After dilution to 12 ml with ultrapure water, K, Ca, Mn, Fe, Cu, and Zn contents of the samples were determined by atomic absorption spectrometry using an AA240FS flame spectrometer (Agilent, USA).

SAMPLE PREPARATION FOR μ PIXE ANALYSIS

In order to avoid losses of elements and their redistribution in the analysed samples, cryotechniques were used. The seeds were fixed by high pressure cryofixation (Leica EM PACT2, Leica Microsystems, Germany) and stored in liquid nitrogen. Subsequent steps were performed in a cryostat (CM3050-S, Leica Microsystems, Germany) at -30°C: seeds were recovered from their cryofixation container in cold isopentane, embedded in an inert matrix (Tissue-Tek® OCT compound, Sakura Finetek, USA) and cut with a tungsten blade in sections of approximately 30 μ m. Sections were individually mounted on a cold aluminium sample holder covered with a pioloform film (1 g pioloform, 75 ml chloroform). Samples were carefully transferred to a freeze-dryer (Alpha 1-4, Christ Martin, Germany), freeze-dried at -10°C, under a vacuum of 0.37 mbar for 48 h and brought back to room temperature (RT) by steps of 5°C per hour. Finally, samples were covered by a second pioloform film and stored in an anhydrous environment at RT until μ PIXE analyses. Section integrity of seed sample was verified by putting a few freshly cut sections directly on a microscope slide (Menzel-Gläser SUPERFROST®, Thermo Fisher Scientific, USA) stored in the cryostat. After sectioning, the microscope slides mounted with seed sections were quickly brought to RT in a vacuum desiccator and checked under a light microscope.

MULTI-ELEMENTAL LOCALIZATION BY μ PIXE

Microanalyses of Arabidopsis sections were performed at the Saclay nuclear microprobe, France (Khodja et al., 2001). The sample holder with seed sections and reference samples were attached to a motorized vacuum goniometer in the analyzing chamber. After closing the chamber and establishing a vacuum (typically 10⁻⁶ mbar), rough positioning of the samples was

achieved using an optical camera. The proton beam of 3.0 MeV and 200 pA current was focused down to a $1.5 \times 1.5 \mu\text{m}^2$ spot, used to rapidly scan large areas of the sample and finally adjusted to scan the seed section areas (usually $500 \times 500 \mu\text{m}^2$) during circa 4 h. Simultaneous PIXE, RBS, and STIM analyses were performed. A 40 mm^2 -Bruker XFlash SDD detector placed at 75° relative to the beam path and at 21 mm working distance of the scanned sample was used to collect X-rays emitted by non-organic elements. A $50 \mu\text{m}$ Mylar foil was placed in front of the detector to stop backscattered protons and attenuate X-ray signals from major elements. Backscattered particles were collected by a 180 mm^2 -surface barrier detector positioned at 170° and 35 mm from incident beam. Energy-loss maps based on STIM were obtained by collecting scattered particles at 30° using a surface barrier detector located behind the sample. In total, 11 seed sections were analyzed: 5 from WT, 5 from *nramp3nramp4* mutant and 1 from *vit1-1* mutant.

MULTI-ELEMENTAL QUANTIFICATION BY μPIXE

In order to generate μPIXE qualitative elemental images, PIXE spectra of the scanned Arabidopsis sections were processed using RISMIN software (Daudin et al., 2003). Distinct regions of particular interest (ROIs) such as seed morphological structures or areas enriched for one particular element were selected as ROIs on the elemental images and the corresponding X-ray spectra were extracted. Associated RBS and STIM spectra and images were used to assess thickness and matrix composition of the different ROIs using SIMNRA (Mayer, 1999) and a modified version of RISMIN software. Finally, all elemental quantifications were obtained by processing the extracted data using SIMNRA and GUPIX (Campbell et al., 2000) softwares. Concentrations of the elements present in the selected ROIs (whole seed section, Fe- and Mn-enriched areas) are presented, either as elemental concentrations of one representative sample (Table 2), or as average elemental concentrations of three to four representative samples (Figures 5A,B).

DATA ANALYSES

Average elemental concentrations presented in (Figures 5A,B) as well as elemental fractions presented in Table 3, were analysed by a Kruskal–Wallis test followed by a Tukey *post-hoc* test for multiple comparisons ($p < 0.05$). For both dry and imbibed seeds, fractions of Fe and Mn that accumulate in their respective enriched areas compared to the whole seed section element content were calculated as followed: [element concentration in enriched area-area of enriched area]/[element concentration in whole seed section-area of whole seed section].

RESULTS

SAMPLE PREPARATION FOR ANALYSES

Seed samples were prepared as recommended in Mesjasz-Przybyłowicz and Przybyłowicz (2002) and described in Materials and Methods. The chosen section thickness was $30 \mu\text{m}$ to obtain the best compromise between 3 requirements: (1) analysing a sufficient amount of matter by μPIXE to obtain accurate quantification, (2) the ability to record RBS and STIM complementary signals, and (3) to resolve Arabidopsis seed morphology ($30 \mu\text{m}$

corresponds to circa 2 cell layers). A cryogenic approach was used to prevent element leakage or redistribution within the sample sections (Mesjasz-Przybyłowicz and Przybyłowicz, 2002). Due to Arabidopsis seed small size, high-pressure freezing was used. In order to confirm that cryosectioning did not alter seed structure, some sections were put aside and directly observed under light microscopy after being rapidly brought back to RT. In agreement with previous reports, Arabidopsis dry seed morphological structure could be clearly defined for $30 \mu\text{m}$ thick longitudinal and transversal sections (Figure 1) (e.g., Vaughan and Whitehouse, 1971). After high-pressure freezing, samples could be either freeze-dried or freeze-substituted as no module for frozen-hydrated state specimen analysis was available (Tylko et al., 2007). Freeze-drying technique has been demonstrated to be neither suitable for observations at the cellular level since it may alter the morphology of the samples, nor for metal speciation since water extraction induces metal speciation artifacts (Sarret et al., 2009). However, freeze-drying is appropriate for elemental mapping, in particular for Fe, at the tissue level and thus, was used in this research (Schneider et al., 2002; Sarret et al., 2013). Furthermore, preliminary tests showed that the substitution of vitrified water by EPOXY resin lead to partial to complete leakage of some elements (data not shown).

ARABIDOPSIS SEED ELEMENTAL DISTRIBUTION USING μPIXE TECHNIQUE

Freeze-dried $30 \mu\text{m}$ thick sections of *Arabidopsis thaliana* dry and imbibed seeds were analysed by Particle-Induced X-ray Emission induced by a focused ion beam (μPIXE) and elemental images were generated (Figure 2). Some macroelements, such as K, and microelements such as Zn are apparently evenly distributed over the dry seed section (Figure A1). Mapping of Ca, which is evenly distributed in the embryo and accumulates to a higher level in

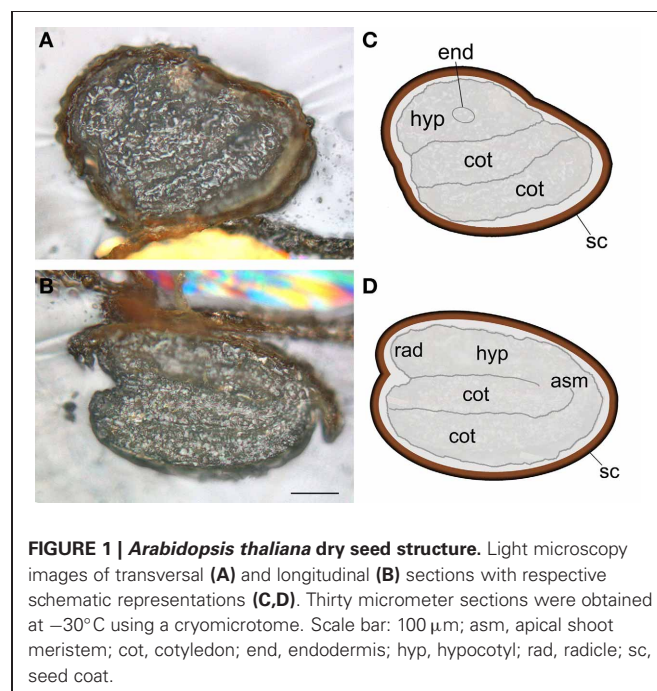


FIGURE 1 | Arabidopsis thaliana dry seed structure. Light microscopy images of transversal (A) and longitudinal (B) sections with respective schematic representations (C,D). Thirty micrometer sections were obtained at -30°C using a cryomicrotome. Scale bar: $100 \mu\text{m}$; asm, apical shoot meristem; cot, cotyledon; end, endodermis; hyp, hypocotyl; rad, radicle; sc, seed coat.

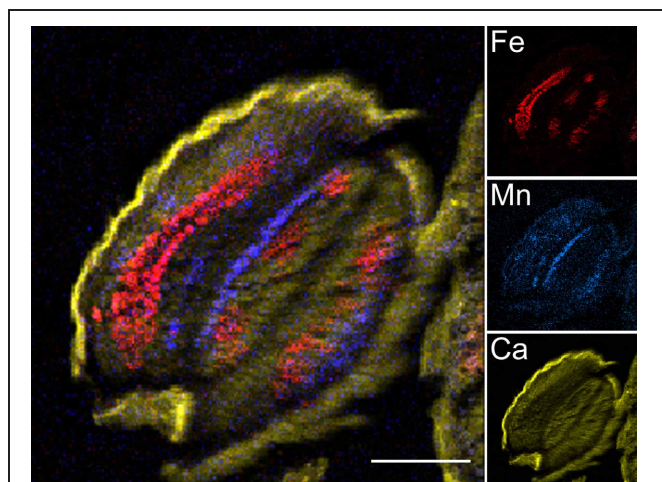


FIGURE 2 | μ PIXE elemental reconstituted image of *Arabidopsis thaliana* dry seed section. Total X-ray spectra of areas scanned with 3 MeV proton beam were collected. Specific distributions of Fe, Mn, and Ca in a 30 μ m thick longitudinal section of *Arabidopsis* dry seed are shown overlaid (large image) and separately (small images). False color images and reconstitution were obtained using ImageJ free software. Scale bar: 100 μ m; red, Fe; blue, Mn; yellow, Ca.

the seed coat, allows the visualization of general seed morphology, recalling the light microscopy image (compare **Figures 1B, 2, Ca** inset). Other elements, such as Fe and Mn, exhibited more specific localization in particular structures, organs or tissues. The elemental map of Fe highlighted the provascular tissue in the cotyledons, hypocotyl and radicle while the Mn elemental map showed concentration on the abaxial (lower) side of the cotyledons (**Figure 2**, Fe and Mn insets). Subsequently, elemental images were used to select regions specifically enriched in Fe or Mn within sections (regions of interest, ROIs; **Figure 3**) to determine the element concentrations within ROIs.

SEED ELEMENTAL CONTENT ANALYSES

Dry seed bulk sample analyses of *Arabidopsis* WT, *nramp3nramp4* and *vit1-1* were performed by atomic absorption spectroscopy (AAS) to determine seed concentrations of various macro- and microelements (**Table 1**). As previously reported, for all quantified elements, no notable differences were observed between seeds of the different genotypes (Lanquar et al., 2005; Kim et al., 2006; Young et al., 2006). Some seeds of the analysed batches were set aside and analysed by μ PIXE to determine elemental concentration in the ROI corresponding to the whole seed section (**Table 1**). In general, seed element average concentrations obtained by μ PIXE for each genotype were in the same range of concentrations but usually higher when compared to AAS analyses (**Table 1**). This difference can be partly explained by the huge difference in the sample sizes used for each technique. AAS elemental content analyses of circa 1800 *Arabidopsis* dry seeds (20 mg aliquots) are compared with μ PIXE analyses of few 30 μ m-thick seed sections (approximately 2 cell layers). Single seed elemental variability, seed cell heterogeneity as well as low replicate number inherent to the technique have a larger impact on the seed elemental content measured by μ PIXE.

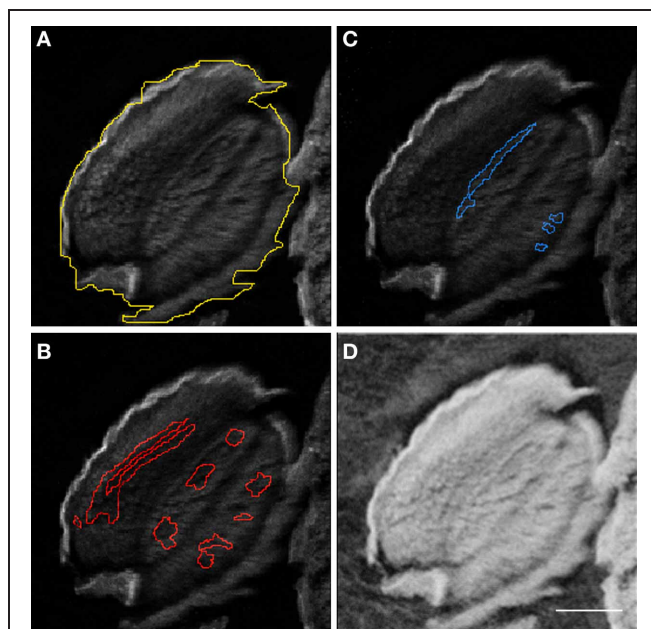


FIGURE 3 | Regions of interest for μ PIXE elemental quantification. Element distribution maps allowed selection of regions of interest (ROI). *Arabidopsis thaliana* whole seed (**A**), Fe- (**B**), and Mn- (**C**) ROIs are shown using the Ca distribution map as background. The associated STIM image (**D**) shows the local mass of the sample which is necessary for quantification. Scale bar: 100 μ m.

Elemental images obtained for the three genotypes showed that Fe distribution is strongly perturbed in *vit1-1* mutant whereas it is not affected in *nramp3nramp4* dry seed at the same stage (**Figure 4**). Similar elemental distribution patterns were obtained using SXRF microtomography (Kim et al., 2006; Donner et al., 2012). ROIs corresponding to whole seed section, Fe- or Mn-enriched areas were selected on the images of each genotype and the corresponding PIXE, RBS, and STIM extracted spectra were analysed (**Table 2**). In WT and *nramp3nramp4* dry seed sections, element quantification within the ROIs confirmed the accumulation of Fe around the provascular tissues of the radicle and cotyledons (from 159 μ g·g⁻¹ DW in Fe-enriched area) and of Mn at the abaxial side of cotyledons (from 44 μ g·g⁻¹ DW up to 386 μ g·g⁻¹ DW). Surprisingly, slight accumulation of Zn also occurred in the Fe-enriched area (from 141 μ g·g⁻¹ DW up to 311 μ g·g⁻¹ DW). In *vit1-1* dry seed, Fe (up to 360 μ g·g⁻¹ DW) and Mn (up to 172 μ g·g⁻¹ DW) concentrated in a single Fe- and Mn-enriched area at the abaxial side of the cotyledons, while no Fe accumulation around provascular tissues was observed. Here also, Zn (up to 224 μ g·g⁻¹ DW) exhibited a slight accumulation within the combined enriched area.

The concentrations of Fe, Mn, and Zn in whole longitudinal and transversal seed sections as well as Fe- and Mn-enriched areas from dry and imbibed samples of WT and *nramp3nramp4* were measured. Statistical analyses of the results did not reveal any significant difference between WT and *nramp3nramp4* in any of the ROIs analysed. **Figure 5** shows the average concentrations

Table 1 | AAS analysis and μ PIXE quantification of various elements in three different *Arabidopsis thaliana* seed genotypes.

| EI. | AAS—dry seed | | | | μ PIXE—seed section | | | | | |
|-----|---|---|---|-------|---|-----|---|-----|---|-----|
| | WT | <i>nr3nr4</i> | <i>vit1-1</i> | LOD | WT | | <i>nr3nr4</i> | | <i>vit1-1</i> | |
| | $\mu\text{g} \cdot \text{g}^{-1} \text{DW}$ | $\mu\text{g} \cdot \text{g}^{-1} \text{DW}$ | $\mu\text{g} \cdot \text{g}^{-1} \text{DW}$ | | $\mu\text{g} \cdot \text{g}^{-1} \text{DW}$ | LOD | $\mu\text{g} \cdot \text{g}^{-1} \text{DW}$ | LOD | $\mu\text{g} \cdot \text{g}^{-1} \text{DW}$ | LOD |
| K | 11881 ± 188 | 11272 ± 463 | 11909 ± 663 | 0.008 | 17648 | 36 | 19989 | 30 | 27274 | 52 |
| Ca | 4200 ± 208 | 4590 ± 288 | 4445 ± 329 | 0.001 | 9165 | 85 | 11223 | 82 | 9757 | 77 |
| Mn | 32 ± 4 | 34 ± 3 | 33 ± 3 | 0.001 | 44 | 4 | 58 | 4 | 65 | 8 |
| Fe | 78 ± 5 | 73 ± 1 | 80 ± 13 | 0.006 | 159 | 3 | 189 | 3 | 157 | 8 |
| Cu | 10 ± 1 | 12 ± 1 | 11 ± 0 | 0.001 | 10 | 2 | 20 | 2 | 68 | 3 |
| Zn | 58 ± 5 | 59 ± 7 | 62 ± 5 | 0.001 | 141 | 2 | 197 | 2 | 169 | 6 |

AAS of bulk samples. Values are means \pm SE ($n = 3-4$ dry seed batches for each genotypes). X-ray and RBS spectra of the whole seed section were selected using RISMIND software and analysed using SIMNRA and GUPIX softwares. Concentrations of one representative measured sample are presented; LOD, limit of detection of the measurements ($\mu\text{g} \cdot \text{g}^{-1} \text{DW}$).

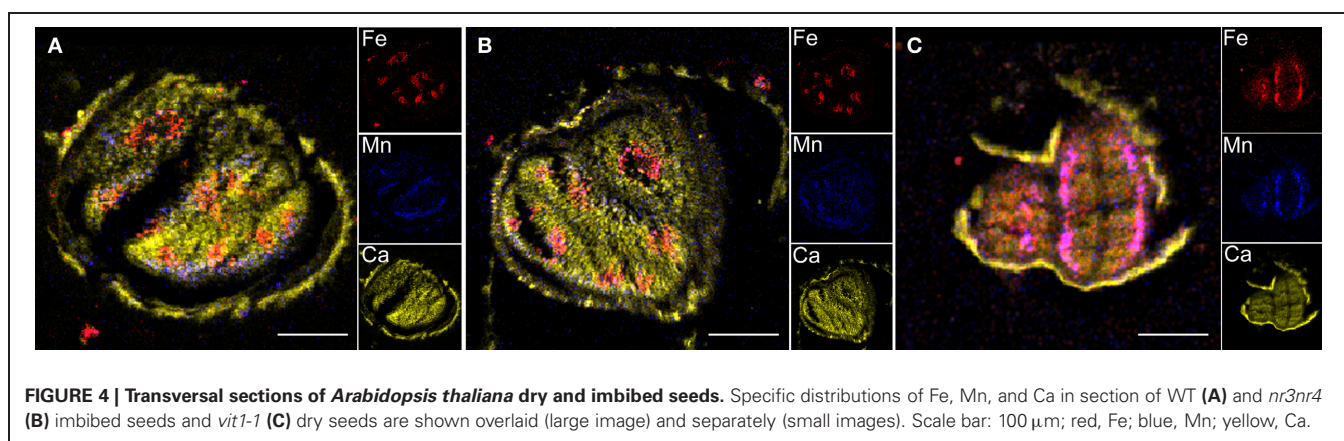


FIGURE 4 | Transversal sections of *Arabidopsis thaliana* dry and imbibed seeds. Specific distributions of Fe, Mn, and Ca in section of WT (A) and *nr3nr4* (B) imbibed seeds and *vit1-1* (C) dry seeds are shown overlaid (large image) and separately (small images). Scale bar: 100 μm ; red, Fe; blue, Mn; yellow, Ca.

of Fe, Mn, and Zn in the whole seed section, Fe- and Mn-enriched areas from dry and imbibed samples. In both sample types, Fe ($751 \pm 134 \mu\text{g} \cdot \text{g}^{-1} \text{DW}$ and $910 \pm 74 \mu\text{g} \cdot \text{g}^{-1} \text{DW}$) significantly accumulated in Fe-enriched provascular area as expected (Figures 5A,B). As already indicated in Table 2, Zn average concentrations were also significantly higher in Fe-enriched areas of both dry and imbibed seeds ($298 \pm 9 \mu\text{g} \cdot \text{g}^{-1} \text{DW}$ and $284 \pm 18 \mu\text{g} \cdot \text{g}^{-1} \text{DW}$). Mn accumulated at significantly higher concentration in the Mn-enriched area at the abaxial side of the cotyledons of both dry and imbibed seeds (Figures 2, 4, 5 and Table 2). Interestingly, after 48 h of imbibition, Fe ($577 \pm 61 \mu\text{g} \cdot \text{g}^{-1} \text{DW}$) and Zn ($268 \pm 14 \mu\text{g} \cdot \text{g}^{-1} \text{DW}$) were also significantly more concentrated within Mn-enriched area at the abaxial side of the cotyledons. These changes were not detectable visually on the elemental maps.

Finally, combining the relative areas of Fe- and Mn-enriched areas and the metal concentration within these areas, we calculated the fractions of Fe and Mn accumulated in their corresponding enriched areas, namely the provascular tissue region for Fe and the abaxial side of the cotyledons for Mn (Table 3). These calculations revealed that at most 69 and 54% of Fe and Mn, respectively, were localized within these areas. A significant fraction of seed Fe and Mn stores is thus localized outside the areas where these metals are the most concentrated.

DISCUSSION

Images of Arabidopsis dry seed sections obtained by μ PIXE confirmed Fe, Mn, and Zn distribution patterns observed previously in WT, *nramp3nramp4* and *vit1-1* mutant dry seeds. In addition, valuable information about local metal concentrations could be obtained from the analyses of the different types of spectra collected at the microprobe installation.

μ PIXE COMPLEMENTS OTHER TECHNIQUES FOR ELEMENTAL IMAGING OF SEEDS

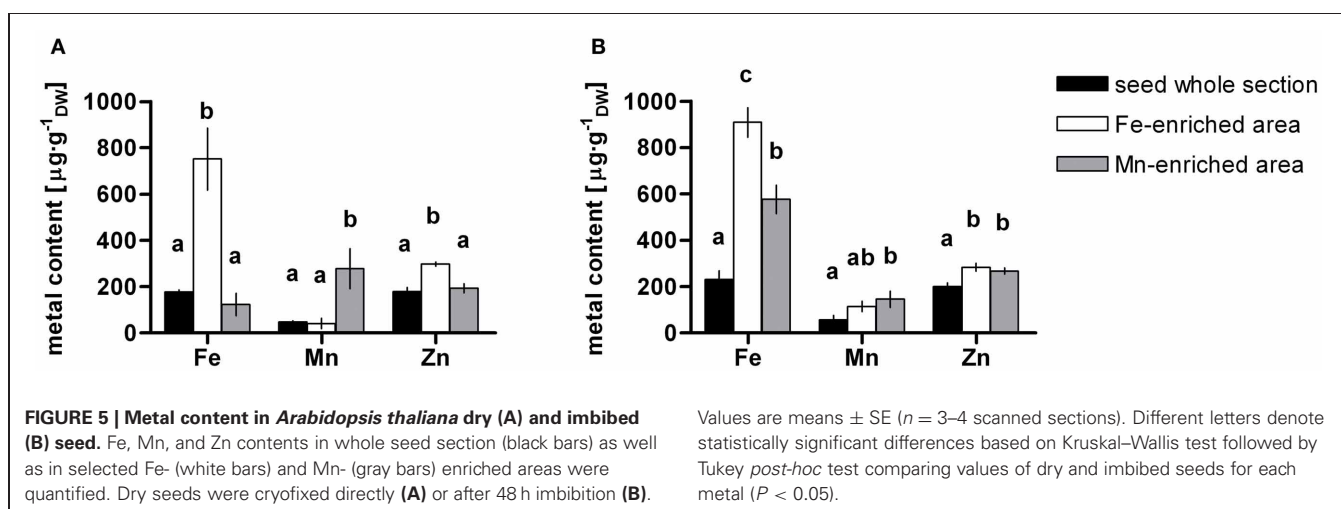
Fe distribution in Arabidopsis seeds has been already reported using different imaging techniques: EDX microanalysis on a transmission electron microscope (Lanquar et al., 2005), Fe histochemical localization using Perls staining with DAB enhancing (Roschztardt et al., 2009) and SXRF microtomography (Kim et al., 2006; Chu et al., 2010; Donner et al., 2012). The respective advantages and limitations of these different approaches have been recently discussed in several reviews (Deves et al., 2005; Lobinski et al., 2006; Ortega et al., 2009; Punshon et al., 2009, 2012; Donner et al., 2012; Sarret et al., 2013).

Combined together, the information about Fe localization obtained by EDX, histochemical staining and SXRF techniques indicate that in Arabidopsis dry seed, Fe is concentrated in the endodermal cells of the embryo and, at subcellular level,

Table 2 | μ PIXE quantification of various elements in three different *Arabidopsis thaliana* seed genotypes with emphasis on Fe- and Mn-enriched areas.

| El. | Seed section | | Fe-enriched area | | Mn-enriched area | |
|---------------|---|-----|---|-----|---|------|
| | $\mu\text{g} \cdot \text{g}^{-1} \text{DW}$ | LOD | $\mu\text{g} \cdot \text{g}^{-1} \text{DW}$ | LOD | $\mu\text{g} \cdot \text{g}^{-1} \text{DW}$ | LOD |
| WT | | | | | | |
| K | 17648 | 36 | 21393 | 71 | 15851 | 121 |
| Ca | 9165 | 85 | 7509 | 140 | 9060 | 155 |
| Mn | 44 | 4 | 22 | 12 | 386 | 13 |
| Fe | 159 | 3 | 751 | 5 | 90 | 20 |
| Cu | 10 | 2 | 9 | 7 | n.d. | n.d. |
| Zn | 141 | 2 | 311 | 11 | 156 | 15 |
| nr3nr4 | | | | | | |
| K | 19989 | 30 | 19443 | 81 | 15582 | 57 |
| Ca | 11223 | 82 | 10672 | 128 | 10315 | 90 |
| Mn | 58 | 4 | 85 | 13 | 106 | 7 |
| Fe | 189 | 3 | 520 | 11 | 218 | 6 |
| Cu | 20 | 2 | 24 | 7 | 9 | 5 |
| Zn | 197 | 2 | 281 | 11 | 200 | 5 |
| El. | Seed section | | Fe- and Mn-enriched area | | | |
| | $\mu\text{g} \cdot \text{g}^{-1} \text{DW}$ | LOD | $\mu\text{g} \cdot \text{g}^{-1} \text{DW}$ | LOD | | |
| vit1-1 | | | | | | |
| K | 27274 | 25 | 22519 | 52 | | |
| Ca | 9757 | 63 | 8826 | 77 | | |
| Mn | 65 | 3 | 172 | 8 | | |
| Fe | 157 | 3 | 360 | 8 | | |
| Cu | 68 | 2 | 60 | 3 | | |
| Zn | 169 | 1 | 224 | 6 | | |

Concentrations of one representative measured sample of dry seed are presented; LOD, limit of detection of the measurements ($\mu\text{g} \cdot \text{g}^{-1} \text{DW}$); n.d., not detected.



is associated with vacuolar globoids (Lanquar et al., 2005; Kim et al., 2006; Roschztardt et al., 2009). Fe concentrations close to $1000 \mu\text{g} \cdot \text{g}^{-1} \text{DW}$ in provascular tissues were quantified by μ PIXE (Table 2 and Figure 5). As globoids represent no more than 10% of the cell volume but constitute the main site for Fe storage in

seed cells (Lanquar et al., 2005), endodermal vacuole globoids possibly contain close to 1% Fe. Besides, it has been hypothesized that Fe in the vacuole is mainly stored as Fe^{3+} but this still remains to be demonstrated using XAS (Pich et al., 2001; Otegui et al., 2002). Based on analyses of mutant phenotypes and metal

Table 3 | Fractions of Fe and Mn in *Arabidopsis thaliana* dry and imbibed seed.

| El. | Dry | | Imbibed | |
|-----|-----------------|----------------|-----------------|----------------|
| | % metal | % area | % metal | % area |
| Fe | 50.7 \pm 11.2 | 13.4 \pm 5.4 | 68.9 \pm 5.8 | 17.1 \pm 1.5 |
| Mn | 30.3 \pm 7.0 | 8.3 \pm 6.0 | 54.4 \pm 2.8* | 22.4 \pm 7.3 |

The fractions of Fe and Mn quantified in their corresponding enriched areas relative to the whole seed section element content were calculated for dry and imbibed seed. Proportion of each enriched area was also calculated. Values are means \pm SE ($n = 3-4$ scanned sections). Asterisk denotes statistically significant differences based on Mann-Whitney U test ($p < 0.05$).

localization, a mechanism of Fe loading into seed endodermal vacuoles mediated by VIT1 and its release into vasculature cells by NRAMP3 and NRAMP4 during germination has been suggested (Morrissey and Guerinot, 2009; Roschztardtz et al., 2009). Accordingly, recent data indicated that no more than 5% of total seed Fe is associated with the plastidial ferritin in Arabidopsis seed (Ravet et al., 2009a). Concentration of Fe in endodermal vacuoles in the vicinity of the provascular strand cells allows rapid and easy mobilization of Fe to the growing parts of the seedling during germination. Nevertheless, our determination of Fe fractions in dry seed sections by μ PIXE revealed that only about 50% of this metal is actually stored around the seed embryo provascular tissue (Table 3). This result indicates that Arabidopsis seeds likely contain two pools of Fe: a pool of concentrated Fe associated to endodermal vacuoles, previously identified by several imaging techniques (Kim et al., 2006; Roschztardtz et al., 2009), and a less concentrated pool spread in all seed tissues, which have so far been overseen. This finding does not contradict the previous assessments about the central role of vacuoles in Fe storage in Arabidopsis (Roschztardtz et al., 2009). However, it points to the existence of significant Fe stores outside of the endodermal cells that have to be taken into account in seed Fe storage models. Moreover, our results suggest that, besides VIT1, which is expressed near the vasculature, other vacuolar Fe uptake systems drive Fe accumulation into the vacuoles of other embryo cells.

Imaging techniques also showed heterogeneous distribution of Mn and Zn in plant seeds (Kim et al., 2006; Vogel-Mikus et al., 2007; Takahashi et al., 2009; Cvitanich et al., 2011). Our analyses by μ PIXE showed that up to 30% of the Mn present in Arabidopsis dry seed is stored in subepidermal cells. Until now, the transporters which mediate Mn accumulation in those particular cells remain unknown. In the case of Zn, our quantification of Zn in seed sections by μ PIXE indicated that Zn is slightly more concentrated in the embryo provascular tissue. This is in agreement with SXRF microtomography of Arabidopsis seeds (Kim et al., 2006) and μ PIXE analysis of Phaseolus seeds (Cvitanich et al., 2011).

METAL REDISTRIBUTION IN ARABIDOPSIS SEED DURING GERMINATION MONITORED BY μ PIXE

During germination, seed undergoes drastic morphological and biological changes, necessarily inducing a redistribution of

macro- and micronutrients (Bewley, 1997; Bolte et al., 2011). Whereas mature dry seed is an excellent plant sample stage for imaging by SXRF or μ PIXE due to its natural dehydration state, monitoring of seed germination by X-ray imaging techniques remains challenging. Here, a μ PIXE analysis of Arabidopsis WT and *nramp3nramp4* mutant after seed imbibition was undertaken. Whereas EDX microanalyses revealed *nramp3nramp4* mutant defect in Fe remobilization from globoids in 2-day old seedling, μ PIXE failed to detect any qualitative or quantitative modification of the Fe pattern between the two genotypes at the same stage (Lanquar et al., 2005). As a matter of fact, elemental maps obtained on dry seeds or after imbibitions did not show any obvious difference. Two hypotheses may account for this apparent discrepancy: (1) even though the experiment was performed after 2 days in both case, seedling development may have been slower in the case of the μ PIXE experiment, due to differences in the conditions or seed age. Attempts to analyse sections of WT and mutant seeds at later stage were hampered by difficulties to obtain intact sections; (2) after 2 days, Fe may have been remobilized from the globoids, preventing its detection by EDX, but may have remained in the same tissues explaining why no change could be detected at the μ m resolution of the μ PIXE analysis.

Nevertheless, quantification of Fe, Mn, and Zn in seed sections by μ PIXE indicates significant distribution modifications after 48 h of imbibition. During imbibition, Fe and Zn concentrations increase in subepidermal cells of cotyledons, corresponding to the Mn-enriched area, while the fraction of Mn accumulating in those cells significantly increases from 30 to more than 50% of total Mn content. Although the origin of the metals could not be identified as no significant decrease of Fe, Mn, or Zn occurs in other areas, it is likely that during seed imbibition metals are already beginning to be redistributed toward the future photosynthetic cell layers. As the Fe concentration in the endodermal area is unchanged after imbibition, it is tempting to speculate that the Fe that accumulates in the sub epidermal area originates from the “diluted pool” spread in all seed tissues, which would therefore represent a more loosely bound Fe pool. The endodermal pool would be released at later stages of germination.

In conclusion, our study using μ PIXE complemented other approaches to analyse metal distribution in Arabidopsis seeds. This fully quantitative approach revealed unsuspected important Fe stores outside the endodermal cells. Analysis after seed imbibition provided a first glimpse at the rapid metal redistribution that occurs during germination.

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APPENDIX

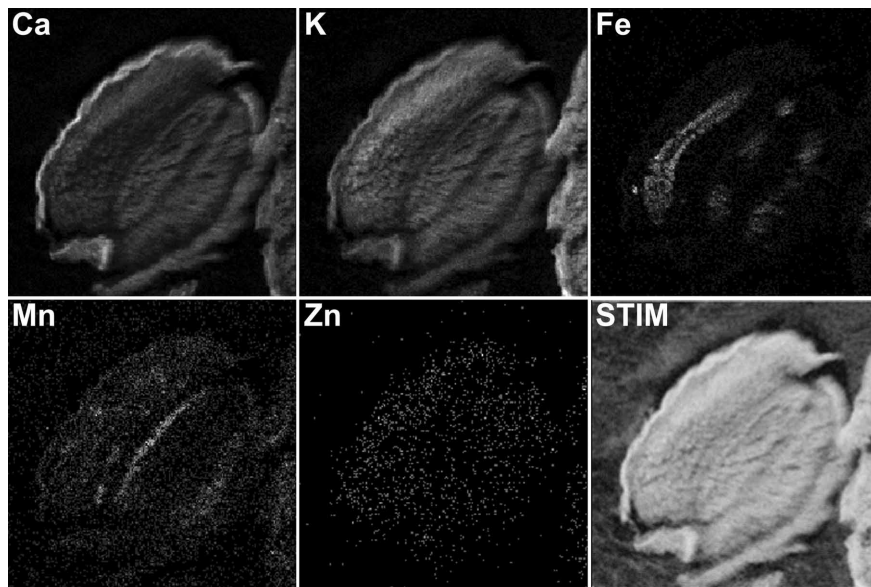


FIGURE A1 | μ PIXE elemental reconstituted image of *Arabidopsis thaliana* dry seed section. Total X-ray spectra of areas scanned with 3 MeV proton beam were collected. Specific distributions of Ca, K, Fe, Mn,

and Zn in a 30 μ m thick longitudinal section of *Arabidopsis* dry seed are separately. The associated STIM image shows the local mass of the sample. Scale bar: 100 μ m.