

Application of nanoscale secondary ion mass spectrometry to plant cell research

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Imaging resource flow in soil-plant systems remains central to understanding plant development and interactions with the environment. Typically, subcellular resolution is required to fully elucidate the compartmentation, behavior, and mode of action of organic compounds and mineral elements within plants. For many situations this has been limited by the poor spatial resolution of imaging techniques and the inability to undertake studies *in situ*. Here we demonstrate the potential of Nanoscale Secondary Ion Mass Spectrometry (NanoSIMS), which is capable of the quantitative high-resolution spatial imaging of stable isotopes (e.g., ¹²C, ¹³C, ¹⁴N, ¹⁵N, ¹⁶O, ¹⁸O, ³¹P, ³⁴S) within intact plant-microbial-soil systems. We present examples showing how the approach can be used to investigate competition for ¹⁵N-labelled nitrogen compounds between plant roots and soil microorganisms living in the rhizosphere and the spatial imaging of ³¹P in roots. We conclude that NanoSIMS has great potential to elucidate the flow of isotopically-labelled compounds in complex media (e.g., soil) and opens up countless new opportunities for studying plant responses to abiotic stress (e.g., ¹⁸O₃, elevated ¹³CO₂), signal exchange, nutrient flow and plant-microbial interactions.

We have used the NanoSIMS technique to investigate the flow of nutrients between microbial and plant cells within the rhizosphere. Secondary Ion Mass Spectrometry

(SIMS) involves bombarding a sample with a high-energy ion beam, which sputters atoms, molecules and electrons from the sample surface. Ionized species (secondary ions) are extracted to a mass spectrometer, sorted according to their energy and their mass-to-charge ratio, and counted. NanoSIMS, a recent development in SIMS, combines high sensitivity with high spatial resolution (typically 100 nm) to allow elemental and isotopic imaging of secondary ions, such as ¹²C, ¹⁶O and ¹²C¹⁴N⁺, on a range of biological materials at the sub-cellular scale (Fig. 1A and B). An element map is obtained by scanning the primary ion beam over the sample surface and measuring the secondary ion intensities of any given ion species, at each pixel in the image. The intrinsically high mass resolution allows the separation of different ion species at the same nominal atomic mass (e.g., ¹²C¹⁵N⁺ from ¹³C¹⁴N⁺ at mass 27), while the multi-collection capability allows the simultaneous measurement of up to five ion species. This makes it possible to obtain images of different isotopes from the same area simultaneously, from which quantitative isotope ratios from individual components can then be extracted. As such, NanoSIMS offers a means of elucidating processes involved in the transport of ions and molecules into cells and their distribution within cells, at scales and sensitivities not attainable by other methods.¹⁻⁵

We previously demonstrated the use of NanoSIMS to image and map the location

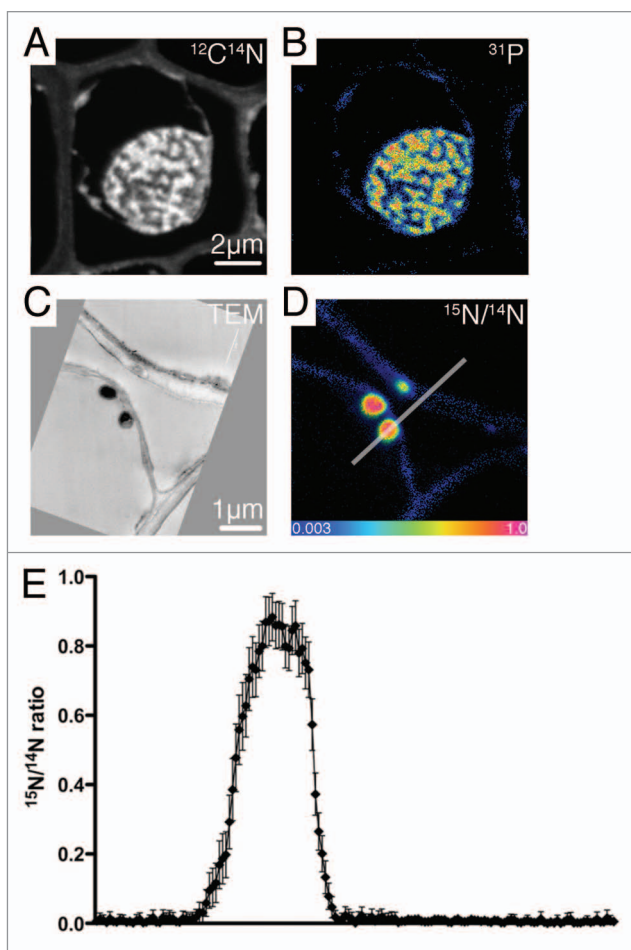


Figure 1. (A) $^{12}\text{C}^{14}\text{N}$ and (B) ^{31}P images of a wheat root cell nucleus from NanoSIMS illustrating the potential to map different elements at the sub-cellular scale; (C) TEM image of two bacteria attached to a cortical cell wall; (D) corresponding $^{15}\text{N}/^{14}\text{N}$ ratio image from NanoSIMS of the same bacteria. The differential uptake of ^{15}N is illustrated by the color scale; ranging from natural abundance (blue) to a $^{15}\text{N}/^{14}\text{N}$ ratio = 1.0 (i.e., 50 at% ^{15}N) (pink) for the plant cell and bacteria, respectively; (E) Linescan (3.5 μm) illustrating the variation in $^{15}\text{N}/^{14}\text{N}$ across an enriched bacterium and an un-enriched plant cell wall (line in D). Error bars are based on the Poisson counting statistics for each pixel.

of ^{15}N -labelled bacterial communities artificially introduced into soil microhabitats.^{6,7} We extended this approach to a natural ecosystem, by examining the differential partitioning of ^{15}N -labelled ammonium ($^{15}\text{NH}_4^+$) between plant roots and soil microbial communities at the nanometer scale (Fig. 1C and D).⁸ It was shown that introduced ^{15}N could be detected, and more importantly, mapped, in individual bacterial cells found in the soil matrix, within the rhizosphere, within root hairs, and intra-cellular within the root. The $^{15}\text{N}/^{14}\text{N}$ ratio data (determined as the ratio between the $^{12}\text{C}^{15}\text{N}$ and the $^{12}\text{C}^{14}\text{N}$ signals) could then be extracted from specific regions of interest—groups

of pixels bounding a particular feature, such as a bacterium or a root cell wall, or linescans (Fig. 1E). This unique approach allows the visualization of nutrient flows and metabolic pathways through complex, multi-component ecosystems. Here we consider further the application of the technique to study nutrient availability in plant cell research.

Uptake of Amino Acids by Plants and Microorganisms

Amino acids represent a large input of organic N to rhizosphere soil, constituting an important source of N to both plants and microorganisms, and they have been

implicated as a major factor regulating ecosystem productivity.⁹ Recently, we, and others, have confirmed that higher plants can capture amino acids from soil.^{10,11} This challenges the paradigm that N must first be microbially processed to inorganic N before becoming plant-available. This direct uptake of organic N by some plants provides an effective shortcut in the N cycle.^{11,12} Most experiments to date have used either ^{15}N - and/or ^{13}C -labelled amino acids to determine the relative degree of plant-microbe competition for dissolved organic N in the rhizosphere using bulk mass spectrometry methods.¹¹ This approach has been criticized, however, as it still remains difficult to distinguish between direct uptake and indirect uptake, where the amino acids are first mineralized to NH_4^+ before being taken up by the roots.⁹ We propose that by using dual-labelled ^{15}N and ^{13}C amino acids it will be possible to image the location of both isotopes within plant cells, associated microbial communities, and fungal associations at the sub-cellular scale. Where both isotopes are co-located within the same pixels of the image we can infer uptake as an organic molecule, while spatial separation of the isotopes in the image map would indicate a split of the molecule. Simultaneous labelling with ^{13}C and ^{15}N has proved effective on other biological systems analyzed by NanoSIMS,^{13–15} however, these experiments did not attach the two isotopes to the same starting molecule. Combining this with the use of halogenated DNA probes that are detectable by the NanoSIMS (e.g., Iodine),¹⁶ it should be possible to study such rhizosphere functions along with molecular identification of target organisms.

Spatial Mapping of Phosphorus (P) within the Soil Matrix

The spatial imaging of phosphorus (P) around roots also represents a real challenge within intact plant-soil systems. This is due to the poor solubility and slow diffusion of P in soil, which consequently produces steep and very narrow depletion zones around roots. Previous research in rhizosphere systems has tended to focus on the use of micro-autoradiography, using ^{32}P and ^{33}P radioisotopes.^{17,18} This

technique is typically limited to a spatial resolution of 0.25 to 1 mm, due to (1) the inability to get close contact between the soil particles and the screen and (2) the hemispherical spread of β -particles, which leads to image blurring. Although only one stable isotope of P exists (^{31}P), NanoSIMS has great potential to help spatially resolve P depletion and transfer in the plant-soil system. For example, in model systems, using sand grains coated with goethite [$\text{Fe}(\text{OH})_3$] to which ^{31}P is uniformly adsorbed, it would be possible to determine the exact spatial extent of the depletion zone in soil with high precision. In addition, it would enable examination of P depletion with root hairs of different ages, or whether the depletion zone extends past the root hair tips, which would imply mobilization by root exudates.¹⁹ NanoSIMS would also allow the spatial heterogeneity of P in the rhizosphere to be determined, as well as the spatial distribution of P within mycorrhizas associated with roots.²⁰

Existing work clearly proves that NanoSIMS has tremendous potential to allow the study of assimilatory processes at the sub-micron level in a wide range of biochemical and molecular applications. The ability to acquire quantitative data at high spatial resolution opens up countless new opportunities for ecosystems involving plant cells, microorganisms and animals.

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