Rapid accumulation of polyphosphate in extraradical hyphae of an arbuscular mycorrhizal fungus as revealed by histochemistry and a polyphosphate kinase/luciferase system

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Summary

The rate of polyphosphate accumulation in extraradical hyphae of an arbuscular mycorrhizal fungus was investigated by conventional histochemistry and a new enzymatic method using a bacterial enzyme, polyphosphate kinase. Marigold (Tagetes patula cv. Bonanza Orange) was inoculated with Archaeospora leptoticha and grown under P-deficient conditions. Extraradical hyphae were harvested at 0, 1, 3 and 24 h after 1 mM P-application. PolyP levels were assessed by both metachromasy of Toluidine blue O and polyphosphate kinase which converted polyP to ATP followed by the ATP-luciferase assay. Percentage of hyphae with metachromatic granules was increased from 25 to 44% from 0 to 1 h, and a maximum of 50% was reached by 3 h. Polyphosphate content was doubled from 1 to 3 h after P-application (4.8–10.0 µmol as Pi mg⁻¹ protein) at a rate of 46.4 ± 15.1 nmol min⁻¹ mg⁻¹. The rate of polyphosphate accumulation in the hyphae was surprisingly rapid as those of polyphosphate-hyper accumulating microorganisms. The enzymatic method employed in the present study allows highly specific and sensitive assessment of polyphosphate in the mycorrhizal system.

Key words: arbuscular mycorrhizal fungi, luciferase, metachromasy, polyphosphate, polyphosphate kinase, Toluidine Blue O.

Introduction

Arbuscular mycorrhizal (AM) fungi are ubiquitous and form symbiotic associations with most terrestrial plant species. It is well documented that the fungi promote growth of the host plant through enhanced uptake of phosphate. High-affinity type Pi-transporter genes have been isolated from AM fungi, and the gene products are responsible for uptake of Pi from the soil solution and into the fungal cytosol (Harrison & Van Buuren, 1995; Maldonado-Mendoza et al., 2001). However, there is a paucity of information on the mechanisms of Pi compartmentation, translocation along hyphae and transfer from fungi to the host.

Inorganic polyphosphate (polyP) is a linear polymer of phosphate linked by high-energy bonds; a wide range of microorganisms store P by accumulation of polyP (Kornberg et al., 1999). It has been suggested that AM fungi accumulate polyP for long distance translocation along hyphae (Callow et al., 1978; Cox et al., 1980; Solaiman et al., 1999). However, detailed information on the dynamics of polyP turnover in AM fungi is lacking, and further research is required to elucidate the role of polyP in the P-delivery system. To this end, establishment of a specific method to quantify polyP is essential. Several colorimetric methods for polyP measurement have been reported (Griffin et al., 1965; Allan & Miller, 1980; Lorenz et al., 1997). Recently, metachromasy of Toluidine
Materials and Methods

Polyphosphate kinase (PPK) was purified from the PPK-over-expressing *Escherichia coli* as described elsewhere (Ahn & Kornberg, 1990). Specific activity of PPK after purification was $3 \times 10^6$ U mg$^{-1}$ protein, in which 1 U of activity was designated to phosphorylate 1 pmol ADP min$^{-1}$ at 37°C using polyphosphate (polyP) as a phosphoryl donor (PPK reverse reaction). Purified PPK was diluted to 3000 U µl$^{-1}$ in 50% glycerol/50 mM Tris/HCl pH 7.5 and stored at −20°C.

Histochemical assessment by Toluidine blue O

The hyphae were removed from the ethanol, blotted on filter paper, stained in 0.05% TBO in 50 mM KCl/ HCl pH 1.0 for 15 min at room temperature, rinsed in 50 mM KCl/HCl pH 1.0 and mounted in the same buffer on a glass microscope slide. Percentage of hyphal length showing metachromasy was assessed using a grid line intersect method as described previously (Ezawa et al., 2001).

Specific measurement by polyphosphate kinase

Five to 10 mg hyphae were disrupted in 300 µl ice-cold acetone with c. 100 mg of zirconia beads (0.5 mm in diam) by a Mini-Bead Beater (Biospec, OK, USA) at 5000 rpm for 3 × 10 s with 1 min intervals on dry ice. Acetone was evaporated by vacuum centrifugation after disruption. The pellet was suspended in 400 µl of the extraction buffer (8 mM HEPES/KOH pH 1.0 and 50 mM Tris/HCl pH 7.5) and centrifuged at 20 000 × g for 5 min at 4°C, and the supernatant was taken as the hyphal extract. Poly P extraction from hyphae was most efficient using 8 M urea, however, it interfered the PPK reaction. Therefore, following polyP extraction, urea was removed by adding 50–100 µl of the extract to a Bio-Spin 6 gel-filtration column (Bio-Rad Laboratories, Tokyo, Japan). PolyP concentration was quantified using the PPK-reverse reaction (Ahn & Kornberg, 1990) with slight modifications as follows: a PPK reaction mixture consisting of 2 µl of the gel-filtrated extract, 50 mM (NH$_4$)$_2$SO$_4$, 4 mM MgCl$_2$, 40 µM ADP and 600 U PPK, 40 mM HEPES/KOH pH 7.5 in a final volume of 20 µl was incubated at 37°C for 40 min. The reaction was terminated by heating at 95°C for 2 min. The mixture was diluted 1 : 100 in an ATP dilution buffer (4 mM EDTA in 100 mM Tris/HCl pH 7.5), and 20 µl aliquots were mixed with the same volume of the luciferase reaction mixture (ATP Bioluminescence Assay Kit CLS II, Roche Diagnostics, Tokyo, Japan). Luminescence was measured as total luminescent counts in 10 s with a Luminescence-PSN (ATTO, Tokyo, Japan). Background ATP levels were
determined by the PPK reaction control in which substrate (polyP) was omitted. Concentration of polyP is given in terms of Pi-residues per mg protein. Protein concentration was determined by the BCA Protein Assay Reagent Kit (PIERCE, IL, USA) using bovine serum albumin as a standard.

Qualitative analysis by gel-electrophoresis

Qualitative analysis of polyP was carried out by polyacrylamide gel-electrophoresis. Fifty to 100 µl of the hyphal extract prepared from samples harvested 0 and 3 h after P-application was desalted in a gel-filtration column, mixed with 1 µl of 10 mg ml\(^{-1}\) ribonuclease A and 0.5 –1.0 µl of 100 m\(\text{M}\) BeSO\(_4\) (1 m\(\text{M}\) in final) and incubated at 37°C for 1 h. BeSO\(_4\) was added to inhibit hydrolysis of polyP by polyphosphatase-type enzymes in the hyphal extract (Ezawa et al., 2001). The final solutions were loaded on 10% polyacrylamide gel buffered with Tris/borate/EDTA and run at 10 – 20 mA constant current. Loading volume was standardized between the samples on the basis of protein concentration of crude extracts. The gel was stained with 0.05% TBO/25% methanol/5% glycerol for 30 min and destained in 25% methanol/5% glycerol. Chain-length of polyP was estimated based on relative mobility (\(R_m\)) to bromophenol blue, a tracking dye, according to Clark & Wood (1987).

Results

Addition of P to P-deficient AM hyphae resulted in an exponential increase in TBO stained metachromatic granules over time. At the first harvest time (0 h after P-application) granules were observed in c. 25% of extraradical hyphae (Figs 1 and 2). One hour after P-application, the percentage of hyphae containing metachromatic granules increased to 44%. Heavy accumulation of metachromatic granules was observed in 49 – 50% of hyphae from 3 to 24 h after P-application.

Figure 3 shows polyP content in hyphae measured by the PPK/luciferase method. No increase in polyP was detected from 0 to 1 h after P-application (4.80 ± 0.60 and 4.40 ± 0.45 µmol mg\(^{-1}\) protein, respectively). The polyP level increased to 9.97 ± 1.75 µmol mg\(^{-1}\) at 3 h and was steady by 24 h after P-application. The net rate of polyP accumulation from 1 to 3 h was calculated as 46.4 ± 15.1 nmol min\(^{-1}\) mg\(^{-1}\) protein (\(P < 0.05\)). Qualitative analysis of polyP by electrophoresis clearly showed heavy accumulation of polyP 3 h after the P-application (Fig. 4). Although the range of polyP chain-length was broad, there was a peak of high-molecular weight polyP around \(R_m\) 0.12, which was estimated as more than 300 Pi-residues.

Discussion

This study provides the first information about the rate of polyP accumulation in the AM symbiosis. Using two different approaches, the maximum level of polyP was shown to occur in less than 3 h after P-application to hyphae growing in soil.
because the TBO staining revealed that only 50% of extraradical hyphae were responsible for polyP biosynthesis, but the rate was calculated on the basis of total protein content of the hyphal material irrespective of the presence of polyP. Therefore, the polyP accumulation rate of the AM fungus could be comparable to that of the polyP-hyper accumulating bacteria, *A. johnsonii* 210 A. It is considered that P-delivery in AM association is regulated through the following four steps: P-uptake, -compartmentation (polyP biosynthesis), -translocation along hyphae and -transfer to the host. The fact that the P-uptake and subsequent polyP biosynthesis were surprisingly rapid suggests that the rate-limiting step of the P-delivery system is the translocation or transfer to the host. This hypothesis requires further confirmation.

It is predicted that external Pi concentration also affects the rate of polyP accumulation. In the present study we applied 1 mM Pi, which was extremely high compared with Pi concentration of normal soil solution. Therefore, the rate of polyP accumulation obtained in the present study can be regarded as a potential rate under optimal conditions. On the other hand, given the heterogeneous distribution of nutrients in soil both spatially and temporally, our results may reflect growth of P-starved hyphae exploiting a newly created or discovered patch of nutrient rich soil.

About 50% of extraradical hyphae showed metachromatic granules using TBO in the present study, consistent with observation of *Glomus coronatum* with *T. patula* (Ezawa et al., 2001). Boddington and Dodd (1999b) observed using DAPI that 50–70% of extraradical hyphae of *G. rosea* in symbiosis with *Desmodium ovalifolium* accumulated polyP irrespective of soil P fertility. The percentage of hyphae with polyP may indicate the percentage of hyphae that contribute to P-uptake and translocation in the hyphal network.

The chain-length distribution of polyP in AM fungi has not been determined precisely. An average chain-length of freely mobile polyP in *Gl. intraradices* was 15 as revealed by NMR (Rasmussen et al., 2000). A broad distribution of polyP chain-length was also observed in the extraradical hyphae of *Gl. etunicatum* in which high-molecular weight polyP was estimated as more than 190 residues (Ezawa et al., 1999). Yeast, a model eukaryotic microorganism, is known to accumulate three to several hundred Pi-residues of polyP (Kornberg et al., 1999). PolyP chain-length may depend on the balance between cell energy status (Schuddemat et al., 1989) and probably on P availability in environment and in turn affect solubility of polyP in situ (Ezawa et al., 2002).

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**Fig. 3** Time course assessment of polyphosphate in extraradical hyphae of *Archaeospora leptoticha* by the polyphosphate kinase/luciferase system. Polyphosphate was extracted and quantified as detailed in Materials and Methods, and polyphosphate content was expressed as Pi-residues per mg protein. Vertical bars show SE (*n* = 3).

**Fig. 4** Electrophoretic analysis of polyphosphate extracted from extraradical hyphae of *Archaeospora leptoticha* 0 and 3 h after P-application. Polyphosphate was extracted, purified as detailed in Materials and Methods and loaded on 10% polyacrylamide gel (buffered with TBE). Loading volume was standardized by protein concentration of the crude extracts (30 µg protein per lane). The position of xylene cyanol (XC) and bromophenol blue (BPB), tracking dyes of electrophoresis, are indicated. R<sub>m</sub>, relative mobility to BPB.
The PPK/luciferase method is highly specific and sensitive to polyP (Ault-Riché et al., 1998). This is the first case that has employed the method in AM associations. It is worthwhile that the TBO method, a conventional histochemical assessment of polyP, has been validated by the PPK method. Generally, results obtained by the TBO method were in good agreement with those obtained by the PPK method: the values at zero time were doubled at 3–24 h after P-application. The disagreement between the data at 1 h after P-application may be caused by the following two reasons. Firstly, we could not assess the density or size of metachromatic granules which might be less or smaller in hyphae at 1 h than those in hyphae at 3 and 24 h, resulting in an overestimation of the TBO-stained material at 1 h. Secondly, sensitivity of the two methods to short-chain polyP may differ. Reactivity of PPK to polyP shorter than 20 residues is much lower than that to longer polyP whereas TBO shows relatively constant reactivity to shorter polyP (T. Ezawa and R. Ohtomo, unpublished results).

It has been reported that short-chain polyP was accumulated at the initial phase of polyP biosynthesis in yeast (Schuddemat et al., 1989). It is likely that the AM fungus would have accumulated short-chain polyP with which TBO could react, but PPK could not react in the first hour after P-application. Little is known about mechanisms of polyP biosynthesis and compartmentation in euakaryotic microorganisms. Most of polyP is compartmentated in vacuoles in euakaryotic microorganisms, yeast (Kornberg et al., 1999) and Neurospora crassa (Yang et al., 1993). Rasmussen et al. (2000) suggested that polyP in AM fungi was also compartmentated in acidic compartments to short-chain polyP with which TBO could react, but PPK could not react in the first hour after P-application.

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